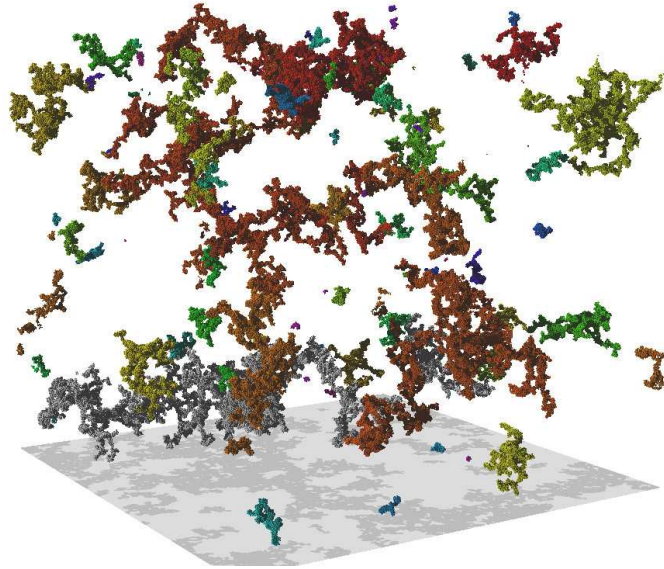


Aggregation and Gelation in Random Networks



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“Believe those who are seeking the truth; doubt those who find it.”

— André Gide (1869–1951)

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Contents

| | | |
|----------|--|-----------|
| 1 | Introduction | 1 |
| 2 | Highly Cross-Linked Brownian Particles | 3 |
| 2.1 | Introduction | 3 |
| 2.2 | Model: Randomly Cross-Linked Particles | 4 |
| 2.3 | Disorder Average | 6 |
| 2.4 | Replica Calculation of the Free Energy | 7 |
| 2.4.1 | Disorder-Averaged Free Energy and Introduction of Replicas | 7 |
| 2.4.2 | Introduction of the Replicated Density Fields | 9 |
| 2.4.3 | Free Energy of the Replicated Density Field | 11 |
| 2.4.4 | Introduction of a Field Theory and Decoupling | 12 |
| 2.4.5 | Universality and Connection to Other Models | 13 |
| 2.5 | Goldstone Fluctuations in General | 14 |
| 2.6 | Ansatz for the Order Parameter | 15 |
| 2.6.1 | Spontaneously Broken Translational Invariance | 17 |
| 2.6.2 | Goldstone Fluctuations and Ansatz for Ω | 18 |
| 2.6.3 | Physical Shear Deformations | 20 |
| 2.7 | Replica Free Energy with Shear Deformations | 21 |
| 2.8 | Returning to the Disorder-Averaged Free Energy | 23 |
| 2.9 | Results | 24 |
| 2.9.1 | Average Cross-Link Density and Higher Moments | 24 |
| 2.9.2 | Gel Fraction | 25 |
| 2.9.3 | Shear Modulus | 27 |
| 2.10 | Conclusions | 29 |
| 3 | Cross-Linked Directed Polymers | 31 |
| 3.1 | Introduction | 31 |
| 3.2 | Properties of a Single Chain | 33 |
| 3.2.1 | Radius of Gyration of an Uncrosslinked Chain | 33 |
| 3.2.2 | Properties of a Polymer Clamped in Space | 34 |
| 3.3 | Full Model: Cross-Linked Directed Polymers | 36 |
| 3.3.1 | Interactions | 36 |
| 3.3.2 | Disorder Average | 39 |
| 3.4 | Replica Calculation of the Free Energy | 40 |
| 3.4.1 | Disorder-Averaged Free Energy: First Steps | 40 |
| 3.4.2 | Introduction of the Replicated Density Field | 41 |
| 3.4.3 | Introduction of a Field Theory and Decoupling | 43 |
| 3.4.4 | Saddle Point Equation | 44 |
| 3.5 | Ansatz for the Order Parameter | 45 |
| 3.6 | The Saddle-Point Equation with Ansatz for Ω | 47 |

| | | |
|----------|--|-----------|
| 3.6.1 | Expansion to second order in Q | 48 |
| 3.6.2 | Sol-Gel Transition | 49 |
| 3.7 | The Equation for the Localization Length | 49 |
| 3.7.1 | Normalization of Length Scales | 49 |
| 3.7.2 | Result for the Distribution of Localization Lengths | 50 |
| 3.8 | Conclusions | 55 |
| 4 | Structure of Spider Silk Modeled as a Random Network | 57 |
| 4.1 | Introduction | 57 |
| 4.2 | Model | 60 |
| 4.2.1 | Unit Cell | 60 |
| 4.2.2 | Crystallite | 60 |
| 4.2.3 | Ensemble of Crystallites | 63 |
| 4.2.4 | Continuous Background | 64 |
| 4.3 | Scattering Function | 65 |
| 4.3.1 | Incoherent Part $G_1(\mathbf{q})$ | 66 |
| 4.3.2 | Coherent Part $G_2(\mathbf{q})$ | 67 |
| 4.3.3 | Scattering Amplitude of a Single Crystallite | 68 |
| 4.4 | Atomic Configuration of the Unit Cell | 69 |
| 4.4.1 | Unshifted Unit Cells | 69 |
| 4.4.2 | Possible Shifts inside the Unit Cell | 72 |
| 4.4.3 | Variations between Crystallites | 72 |
| 4.5 | Results | 73 |
| 4.5.1 | Experimental Scattering Function | 73 |
| 4.5.2 | Scattering Function from the Model | 74 |
| 4.6 | Conclusions | 79 |
| 5 | Aggregation and Gelation in Wet Granular Materials | 81 |
| 5.1 | Introduction | 81 |
| 5.2 | Models | 83 |
| 5.3 | Cooling Dynamics | 86 |
| 5.3.1 | Early Stage of Cooling | 87 |
| 5.3.2 | Late Stage of Cooling | 90 |
| 5.3.3 | Velocity Distribution | 95 |
| 5.3.4 | Partitioning of the Energy into Translational, Rotational and Internal Degrees of Freedom | 98 |
| 5.4 | Aggregation | 101 |
| 5.4.1 | Self-Similar Growth | 103 |
| 5.4.1.1 | Fractal Dimension of the Aggregates | 103 |
| 5.4.1.2 | Cluster Mass Distribution | 105 |
| 5.4.1.3 | Number of Clusters and Mean Cluster Mass | 106 |
| 5.4.2 | Properties of the Asymptotic Cluster | 107 |
| 5.4.2.1 | Fractal Dimension from Radius of Gyration | 108 |
| 5.4.2.2 | Fractal Dimension from Box Counting Algorithm | 110 |

| | | |
|----------|---|------------|
| 5.4.2.3 | Pair Correlation Function and Correlation Dimension | 114 |
| 5.4.2.4 | Coordination Number | 118 |
| 5.4.2.5 | Inertia Tensor and Spatial Extension | 122 |
| 5.5 | Conclusions | 123 |
| A | Properties of \perp- and \parallel-vectors | 125 |
| B | Calculations for the Randomly Cross-Linked Particle Model | 127 |
| B.1 | Calculation of the Replica Free Energy | 128 |
| B.1.1 | Emergence of the Replica Free Energy | 128 |
| B.1.2 | Introduction of the Replicated Density Field | 129 |
| B.2 | Hubbard-Stratonovich Transformation | 130 |
| B.2.1 | Applying the Hubbard-Stratonovich Transformation | 130 |
| B.2.2 | Meaning of the Order Parameter $\Omega(\hat{q})$ | 132 |
| B.3 | Expansion of the replica free energy to 3rd Order in Ω | 134 |
| B.4 | Fluctuations around the Saddle-Point Solution | 135 |
| B.5 | Replica Free Energy with Shear Deformations | 136 |
| B.5.1 | Preparing the Replica Free Energy for the Ansatz | 136 |
| B.5.2 | Expansion of the one-particle partition function \mathfrak{z} | 137 |
| B.5.3 | Insertion of the Order Parameter in \mathfrak{z} | 137 |
| B.5.4 | Insertion of the Order Parameter in $f_{n+1}\{\Omega\}$ | 145 |
| B.5.5 | Recomposing the Replica Free Energy | 146 |
| B.5.6 | Analysis of the quantity Ξ_{r,a^2} | 149 |
| B.6 | Auxiliary Calculations for the Results Section | 150 |
| B.6.1 | Distribution of the Number of Cross-Links | 150 |
| B.6.2 | Shear modulus | 152 |
| B.6.3 | Square of the expansion for $(1 - Q)$ | 153 |
| B.7 | The Order Parameter in the One Replica Sector | 154 |
| C | Calculations for the Directed Polymer Model | 157 |
| C.1 | Calculations for a Single Chain | 157 |
| C.1.1 | The Uncrosslinked Chain | 157 |
| C.1.2 | The Polymer Clamped in Space | 159 |
| C.2 | Examples of Single Chain Interactions | 160 |
| C.3 | Calculations for the Disorder-Averaged Free Energy | 162 |
| C.3.1 | Introduction of Replicas | 162 |
| C.3.2 | Introduction of the Replicated Density Field | 163 |
| C.3.3 | The Hubbard-Stratonovich Transformation | 164 |
| C.4 | The Saddle-Point Equation with Ansatz | 167 |
| C.4.1 | Expansion in Q to Infinite Order | 167 |
| C.4.2 | Expansion to Second Order in Q | 169 |
| C.5 | Obtaining the Equation for the Localization Length | 173 |
| C.5.1 | Normalization of Length Scales | 173 |
| C.5.2 | Laplace-Transformation of the Saddle Point Equation | 175 |

| | | |
|----------|--|------------|
| C.6 | Calculation of a Correlator | 177 |
| C.7 | The Average Cross-Link Density | 180 |
| D | Expressions for the Wet Granulates | 183 |
| D.1 | Infinite Sums similar to the Exponential Function | 183 |
| D.2 | Inverting the Equation $x^{-2}e^x = t$ | 183 |
| D.3 | Computation of the radial distribution function $g(r)$ | 184 |
| E | Further Results from the Spider Silk Model | 187 |
| E.1 | Effect of the Continuous Background | 187 |
| E.2 | Relevance of the Coherent Part of the Scattering Function | 187 |
| E.3 | Simplification of the Scattering Amplitude $A(\mathbf{q})$ | 188 |
| F | Used Symbols and Notation | 191 |
| F.1 | Notation | 191 |
| F.2 | Symbols for the Randomly Cross-Linked Particle Model | 192 |
| F.3 | Symbols for the Cross-Linked Directed Polymers | 194 |
| F.4 | Symbols for the Spider Silk Model | 197 |
| F.5 | Symbols for the Wet Granular System | 199 |
| | Bibliography | 201 |
| | Index | 211 |
| | Curriculum Vitae | 215 |

Introduction

Let us consider a large system, made up of many identical interconnected building blocks, so that close-by building blocks are preferably connected. Then, one may raise a number of questions, for example, under which conditions it is possible to go from one side of the system all the way to the other side, only using the connections (or equivalently that a “percolating” or “macroscopic” cluster is present). Or one may ask, what the structure of the clusters is, which the connections give rise to, for example how large the cluster are and whether or not they have a fractal structure.

These types of problems are known under the term *percolation theory* and despite their seemingly abstract and simple concepts, they are highly relevant for many problems in physics and for a wide range of applications in every day’s life. An example would be the spreading of forest fires: will a fire – starting at one tree – eventually span the whole forest, or will it be extinguished quickly only burning down a few trees? For this case, we imagine that our “building blocks” are the trees in the forest and we define them as connected if they are close enough so that a burning tree would ignite the other one. Then, the cluster sizes of this system will provide indications what the typical extent of the fire will be, and the presence of a macroscopic cluster gives information whether or not the fire may span the whole system. Further examples include filtering of fluids or gases through porous materials, e.g. movement of petroleum through fractured rock, the flow of electrical current through a random resistor network, or the spreading of diseases in a social network.

Another physically important example, treated in this thesis, is *polymerization*, where chemical bonds connect atoms or molecules and can thereby create large networks of macromolecules. To achieve this, one may start with a system of polymer chains (typically chains of connected carbon atoms) in a solvent and add a certain amount of cross-linkers (typically sulfur). Thereupon, one observes that the cross-links interconnect the chains, whereby larger and larger aggregates are formed. When the concentration of the cross-links exceeds a critical value, one can find an aggregate that spans the whole system. In this process the system undergoes a transition from a liquid (sol) to an amorphous solid (gel) phase, the so called “gelation transition”. Materials created in this way (like for example rubber) have interesting properties in terms of flexibility and toughness. It is important to note they have no lattice structure, in comparison to metals; instead they are *amorphous* because the aggregation mechanism always involves some randomness, and consequently they are translationally and rotationally invariant on a macroscopic scale.

The formation, structure and properties of aggregates created by random connections, and in particular the above-mentioned gelation transition are the topic of this thesis. In chapter 2 we will discuss **cross-linked Brownian particles** and particularly their elastic properties. When the cross-link density is large enough, such that a macroscopic cluster is present, the system exhibits a finite resistance to shear deformations. This mechanical resistance is quantified by the shear modulus, the focus of this chapter. We extend previous results, which were restricted to systems close to the gelation transition, to highly cross-linked gels.

In chapter 3 we investigate **cross-linked directed polymers**, where the polymer chains are aligned along a preferential direction. This anisotropic system is a simple model for describing polymers, highly stretched along one direction, or alternatively polymers in a nematic field. As cross-links are added to this system, we observe a gelation transition and investigate properties of the system close to that transition. Focussing on the distribution of localization lengths, we describe the extent of fluctuations of the chains about their mean positions. As we will see, the anisotropy of the system gives rise to a dependence of these structural properties on the distance from the boundaries in the preferred direction.

In chapter 4 we determine the microscopic structure of **spider-silk**, where small crystallites are interconnected by a network of chains. This composition of the material gives rise to outstanding mechanical properties, in particular the toughness and extensibility. We develop a statistical model for the crystallites; from that, the scattering function can be computed and be compared to experimental results, enabling us to determine structural, as well as statistical parameters of the system. We also investigate the role of coherent scattering from different crystallites and the importance of a background scattering density between crystallites.

And in chapter 5 we study the aggregation of **wet granular particles**. Here, the particles conjoin and develop clusters, because they are covered by a thin liquid film and may form “liquid bridges” between each other if the films touch each other and merge. These bonds induce an attractive and hysteretic interaction such that each bond rupture dissipates a fixed amount of energy. The system shows an interesting cooling behavior different from dry granulates, and a transition from fast cooling to a clustering state as soon as the particles’ kinetic energy is not sufficient to break the bonds anymore. As time proceeds, the clusters form larger and larger aggregates, which grow in a self-similar fashion. For all finite densities a percolation transition is observed, with a macroscopic cluster that is fractal on small length scales and compact on large length scales.

A number of technical points have been deferred to the appendices. In particular, the symbols and variables used in each chapter and a short remark on the notation can be found in Appendix F.

Highly Cross-Linked Brownian Particles

Contents

| | | |
|-------------|--|-----------|
| 2.1 | Introduction | 3 |
| 2.2 | Model: Randomly Cross-Linked Particles | 4 |
| 2.3 | Disorder Average | 6 |
| 2.4 | Replica Calculation of the Free Energy | 7 |
| 2.4.1 | Disorder-Averaged Free Energy and Introduction of Replicas | 7 |
| 2.4.2 | Introduction of the Replicated Density Fields | 9 |
| 2.4.3 | Free Energy of the Replicated Density Field | 11 |
| 2.4.4 | Introduction of a Field Theory and Decoupling | 12 |
| 2.4.5 | Universality and Connection to Other Models | 13 |
| 2.5 | Goldstone Fluctuations in General | 14 |
| 2.6 | Ansatz for the Order Parameter | 15 |
| 2.6.1 | Spontaneously Broken Translational Invariance | 17 |
| 2.6.2 | Goldstone Fluctuations and Ansatz for Ω | 18 |
| 2.6.3 | Physical Shear Deformations | 20 |
| 2.7 | Replica Free Energy with Shear Deformations | 21 |
| 2.8 | Returning to the Disorder-Averaged Free Energy | 23 |
| 2.9 | Results | 24 |
| 2.9.1 | Average Cross-Link Density and Higher Moments | 24 |
| 2.9.2 | Gel Fraction | 25 |
| 2.9.3 | Shear Modulus | 27 |
| 2.10 | Conclusions | 29 |

2.1 Introduction

Randomly cross-linked networks of molecules undergo a transition from a fluid to an amorphous solid state, as the number of cross-links is increased. By adding more and more permanent cross-links to the system, molecular clusters are gradually joined and in this process grow to larger and larger objects. Thereby the

system undergoes a transition from the **sol phase**, a complex fluid characterized by molecular clusters of various but finite sizes, to the **gel phase**, an amorphous solid where the constituents of the system have joined to at least one *macroscopic* (system spanning) cluster yielding a non-zero resistance to shear deformations. This *sol-gel transition* or *gelation transition* has been studied in recent years [Deam & Edwards, 1976; Ball & Edwards, 1980; Goldbart *et al.*, 1996; Panyukov & Rabin, 1996], and the structure as well as the elasticity [Mukhopadhyay *et al.*, 2004; Goldbart *et al.*, 2004] has been investigated. However, almost all theoretical studies have focussed on the *near-critical* region, where the analysis simplifies due to the existence of a small parameter – the distance to the transition point or, equivalently, the order parameter. In contrast, the *highly cross-linked* regime has hardly been investigated [Broderix *et al.*, 2002], yet it is particularly interesting in the application of random network models to glasses [Zallen, 1983].

Here we consider a particularly simple network which is built from spherical particles, connected by harmonic springs. This allows us to access the highly cross-linked regime and compute the shear modulus for *arbitrary* cross-link concentrations. It is found that the shear modulus is independent of all microscopic length scales of the model, such as the spring constant and the length scale that characterizes the localization of particles in the amorphous solid state. Instead, the shear modulus is completely determined by the *particle density* and the *density of cross-links*.

We approach the problem using equilibrium thermodynamics. The average over all cross-link configurations – the so called *disorder average* – is done with the help of the replica theory. Furthermore, to simplify the notation, energies are measured in units of $k_B T$.

The chapter is organized as follows: In Sec. 2.2 the model of randomly cross-linked particles and its interactions are presented and in Sec. 2.3 the disorder average is defined. In Sec. 2.4 we establish a field theory of the presented model and find a suitable order parameter. In Sec. 2.5 we have a short general look at Goldstone fluctuations, which are necessary to establish an Ansatz for the order parameter, done in Sec. 2.6. In Sec. 2.7 we apply this Ansatz to the field theory, and establish the connection of the result to the actual disorder-averaged free energy of the system in Sec. 2.8. The final results are presented in Sec. 2.9, and we will conclude in Sec. 2.10.

2.2 Model: Randomly Cross-Linked Particles

We consider a system of N identical particles at positions $\mathbf{r}_1, \dots, \mathbf{r}_N$ in a D -dimensional volume V . Permanent cross-links connect M randomly chosen pair of particles, such that a particular cross-link realization \mathcal{C} is specified by a list of M pairs of monomers $\mathcal{C} = \{(i_1, j_1), \dots, (i_M, j_M)\}$ (see Fig. 2.1). The cross-links are modeled by *harmonic springs*, so that the Hamiltonian for the cross-link interaction becomes:

This interaction is necessary to prevent the system from collapsing. Its strength λ has to be large enough to overcome the attractive interaction produced by the cross-links, and the function $U(\mathbf{x})$ has to fall off fast enough (at the end of Sec. 2.4.3 we will see, what that means specifically). Since λ already scales the strength of the interaction we take the freedom to require $\int_V U(\mathbf{x}) d\mathbf{x} = 1$.

The overall Hamiltonian is $H_{\mathcal{C}} = H_{\mathcal{C}}(\{\mathbf{r}_j\}) = H_{\text{Xlink}} + H_{\text{ev}}$. The index \mathcal{C} will remain as a reminder on variables that depend on the quenched disorder \mathcal{C} specifying the configuration of cross-links. All thermodynamic properties, including the elastic ones, can be obtained from the partition function

$$Z_{\mathcal{C}} = \int D\mathbf{r}_1 \cdots D\mathbf{r}_N e^{-H_{\mathcal{C}}(\{\mathbf{r}_j\})}. \quad (2.6)$$

Here we use the measure $D\mathbf{r}_j := d\mathbf{r}_j/V$ to make sure that $Z_{\mathcal{C}}$ is dimensionless.

This model became known as the **randomly (cross-)linked particle (RLP) model**, and is a very simple model for amorphous solids. As we see in the illustration 2.1 it can produce highly branched “molecule chains”. It is capable of predicting structural and mechanical properties [Broderix *et al.*, 2002; Mao *et al.*, 2007] and has been investigated with and without the use of replica technique. As we will see in Sec. 2.4.5, in the vicinity of the gelation transition, the model falls into the universality class of randomly cross-linked media.

2.3 Disorder Average

The average over all cross-link configurations \mathcal{C} is called the *disorder average* and denoted by $[\cdot]$. The disorder average of a quantity $\mathcal{A}_{\mathcal{C}}$ is calculated by:

$$[\mathcal{A}_{\mathcal{C}}] = \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \mathcal{A}_{\mathcal{C}} \cdot P(\mathcal{C}) \quad (2.7)$$

For the cross-link distribution $P(\mathcal{C})$, we use the Deam-Edwards distribution [Deam & Edwards, 1976]:

$$P_{\text{DE}}(\mathcal{C}) = \frac{1}{\mathcal{Z}_1} \frac{1}{M!} \left(\frac{\mu^2}{2N\phi} \right)^M Z_{\mathcal{C}} \quad (2.8)$$

$$\text{with } \phi = \frac{a^D (2\pi)^{D/2}}{V},$$

and with a normalization constant \mathcal{Z}_1 . On first glance it seems to be counter-intuitive that the probability for a cross-link configuration \mathcal{C} is proportional to the corresponding partition function $Z_{\mathcal{C}}$, but this characteristic is a very clever trick to make sure only particles close to each other get connected. In this regard, let us have a look at a cross-link configuration \mathcal{C} ; by writing the partition function (2.6) slightly differently, one can express the probability for this configuration as

$$P_{\text{DE}}(\mathcal{C}) \propto \left\langle e^{-\frac{1}{2a^2} \sum_{e=1}^M (\mathbf{r}_{i_e} - \mathbf{r}_{j_e})^2} \right\rangle_{H_{\text{ev}}},$$

where $\langle \cdot \rangle_{H_{\text{ev}}}$ is averaging with the statistical weight of the excluded volume interaction. In this form, one can see that configurations for which the distances $|\mathbf{r}_{i_e} - \mathbf{r}_{j_e}|$ are *small* have a *high probability*, while configurations that would involve some *long cross-links* have a *low probability*. For a detailed discussion of the Deam-Edwards distribution, see the work by Goldbart *et al.* [1996]; Broderix *et al.* [2002].

The parameter μ^2 determines the probability for a cross-link to be formed, and therefore controls the sol-gel transition. As we will see in Sec. 2.9.1, the average cross-link density is given by $[M]/N = \mu^2/2$ and the standard deviation, relative to the mean, vanishes in the thermodynamic limit. Thus, $\mu^2 = 2[M]/N$ is the *average coordination number*, i.e. the average number of particles to which a certain particle is connected.

2.4 Replica Calculation of the Free Energy

2.4.1 Disorder-Averaged Free Energy and Introduction of Replicas

From this model, we now would like to calculate the free energy, in order to access mechanical and structural properties. However, we are not interested in a specific cross-link configuration (and we cannot choose the cross-link configuration for 10^{23} particles by hand anyway), but only in *typical* properties of the system. Therefore we have to calculate the *disorder-averaged free energy*

$$[F_C] = -[\ln Z_C]. \quad (2.9)$$

In general it is very difficult to perform the disorder average, Eq. (2.7), of a logarithm. A successful way to perform the sum nevertheless is the replica trick: Instead of averaging the logarithm itself, we take advantage of a representation of the logarithm:

$$- [F_C] = [\ln Z_C] = \left[\lim_{n \rightarrow 0} \frac{Z_C^n - 1}{n} \right] = \lim_{n \rightarrow 0} \frac{[Z_C^n] - 1}{n}, \quad (2.10)$$

and thus we try to calculate

$$[Z_C^n] = \frac{1}{Z_1} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{1}{M!} \left(\frac{\mu^2}{2N\phi} \right)^M Z_C \cdot Z_C^n. \quad (2.11)$$

At first, let's have a look at the rightmost part of Eq. (2.11). At least for integer n , $Z_C \cdot Z_C^n$ can be written as a product $\prod_{\alpha=0}^n Z_C$. By doing that, we introduce $n+1$ **copies of the system with identical cross-link configuration** (one of

the copies, which we call the zeroth, stems from the Deam-Edwards distribution):

$$\begin{aligned}
Z_C \cdot Z_C^n &= \prod_{\alpha=0}^n \int \left(\prod_{j=1}^N D\mathbf{r}_j^{(\alpha)} \right) e^{-H_C(\{\mathbf{r}_j^{(\alpha)}\})} \\
&= \int \left(\prod_{j=1}^N \prod_{\alpha=0}^n D\mathbf{r}_j^{(\alpha)} \right) \exp \left(-\frac{1}{2a^2} \sum_{e=1}^M \sum_{\alpha=0}^n (\mathbf{r}_{i_e}^{(\alpha)} - \mathbf{r}_{j_e}^{(\alpha)})^2 - \sum_{\alpha=0}^n H_{\text{ev}}(\{\mathbf{r}_j^{(\alpha)}\}) \right) \\
&= \int \left(\prod_{j=1}^N D\hat{\mathbf{r}}_j \right) \exp \left(-\frac{1}{2a^2} \sum_{e=1}^M (\hat{\mathbf{r}}_{i_e} - \hat{\mathbf{r}}_{j_e})^2 - H_{\text{ev}}^{(n+1)} \right), \tag{2.12}
\end{aligned}$$

with $H_{\text{ev}}^{(n+1)} := \sum_{\alpha=0}^n H_{\text{ev}}(\{\mathbf{r}_j^{(\alpha)}\})$. Here we introduced the $n+1$ times replicated (and hence $D(n+1)$ -dimensional) vectors

$$\hat{\mathbf{r}}_j := (\mathbf{r}_j^{(0)}, \dots, \mathbf{r}_j^{(n)}). \tag{2.13}$$

This introduction of $n+1$ replicas of the system¹ is the main characteristic of replica theory and the reason for its name. One should keep in mind that the disorder is the same in all replicas; in particular M and i_e, j_e do not have replica indices (α). Assuming ergodicity, a possible way to imagine this replicated system is to see the particle conformations $\{\mathbf{r}_j^{(0)}\}, \{\mathbf{r}_j^{(1)}\}, \dots, \{\mathbf{r}_j^{(n)}\}$ as instances of the system taken at different times, with very long time differences in between. In this picture, the particle conformation in the zeroth replica, $\{\mathbf{r}_j^{(0)}\}$, corresponds to the particle conformation during cross-linking.

As we see in Appendix B.1.1, the sum over M in Eq. (2.11) can be performed and $[Z_C^n]$ can be written as:

$$[Z_C^n] = \frac{\mathcal{Z}_{n+1}}{\mathcal{Z}_1} \tag{2.14a}$$

$$\text{where } \mathcal{Z}_{n+1} = \int \left(\prod_{j=1}^N D\hat{\mathbf{r}}_j \right) \exp(-N\tilde{f}_{n+1}\{\hat{\mathbf{r}}_j\}) \tag{2.14b}$$

with the “**replica free energy**”

$$\tilde{f}_{n+1}\{\hat{\mathbf{r}}_j\} = -\frac{\mu^2}{2\phi N^2} \sum_{i,j=1}^N \exp\left(-\frac{(\hat{\mathbf{r}}_i - \hat{\mathbf{r}}_j)^2}{2a^2}\right) + \frac{\lambda}{2N} \sum_{\alpha=0}^n \sum_{i,j=1}^N U(\mathbf{r}_i^{(\alpha)} - \mathbf{r}_j^{(\alpha)}), \tag{2.14c}$$

which incorporates the degrees of freedom of the system, the replicated particle positions $\{\hat{\mathbf{r}}_1, \dots, \hat{\mathbf{r}}_N\}$. That means, the *disorder-averaged* $[Z_C^n]$ for a D -dimensional system can be expressed as a partition function \mathcal{Z}_{n+1} that involves $D(n+1)$ -dimensional vectors but *does not involve a disorder average* anymore. We will refer to \mathcal{Z}_{n+1} as the “**replica partition function**”.

\mathcal{Z}_1 was originally introduced as a mere normalization constant in Eq. (2.8); knowing that $\lim_{n \rightarrow 0} [Z_C^n] = 1$, we can confirm in Eq. (2.14a) that the notation for \mathcal{Z}_1 is chosen consistently.

¹or possibly only n replicas, if the cross-link distribution is not the Deam-Edwards distribution

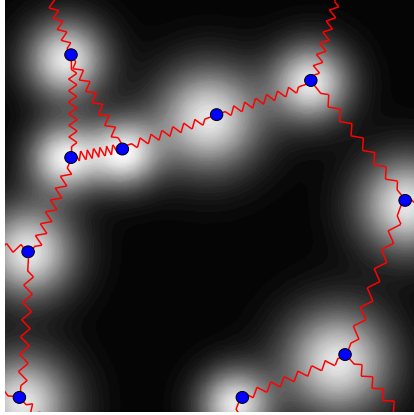


Figure 2.2: Exemplary illustration of a snapshot of the model as in Fig. 2.1, which only contains *localized* particles. Additionally, the density plot behind the snapshot illustrates, how the thermally averaged density as defined in Eq. (2.15) might look like.

2.4.2 Introduction of the Replicated Density Fields

Later we want to introduce a field theory, so sooner or later we have to think about a suitable order parameter. A crucial requirement is that it can distinguish between the liquid and the amorphous solid state. A first idea would be the thermally averaged density

$$\varrho_{\mathcal{C}}^{(\alpha)}(\mathbf{x}) := \frac{1}{N} \sum_{j=1}^N \left\langle \delta(\mathbf{x} - \mathbf{r}_j^{(\alpha)}) \right\rangle_{H_{\mathcal{C}}} . \quad (2.15)$$

In the liquid phase, when no macroscopic cluster is present, any particle – even if it is part of a small cluster of connected particles – has equal probability to be found anywhere in the sample; hence this order parameter would simply be a constant. In the gel phase, on the other hand, a macroscopic cluster is present, which is a cluster that contains an infinite number of particles as $N \rightarrow \infty$. A particle of that cluster can not pass through the entire system, because cross-linking to the cluster will restrict its movement to fluctuations around a mean position (as illustrated in Fig. 2.2). Hence we will refer to such a particle as a “**localized particle**”.

The emerging density fluctuations may be utilized to distinguish the gel from the sol phase *for a given cross-link configuration* \mathcal{C} . However, quantities we are interested in, should not be restricted to a certain cross-link configuration, but should be disorder averaged. And when averaging over all cross-link configurations, the macroscopic cluster can be embedded anywhere in the sample and hence the probability to find a particle at any place in the sample should be the same – as in the liquid phase.

To overcome this problem, we take advantage of the fact that the cross-link configuration is the same for all replicas and hence the position of a localized particle should be correlated among the replicas: In the context of gelation, the crucial

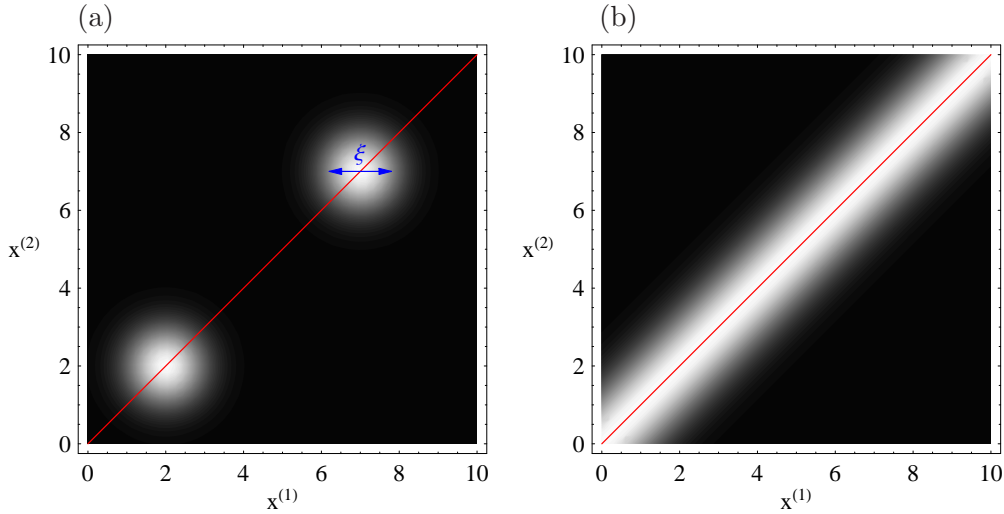


Figure 2.3: For one spatial dimension, the replicated density $O(x^{(1)}, x^{(2)})$, the probability to find a particle at $x^{(1)}$ in replica 1 and at $x^{(2)}$ in replica 2, is illustrated; white means high values. The red line is the angle bisector $x^{(1)} = x^{(2)}$. (a) Example with two localized particles fluctuating around the mean positions $R_1 = 2$ and $R_2 = 7$. The blue arrow (\leftrightarrow) represents the localization length. (b) For the disorder average, we have to integrate the mean positions over the system size. Thus, $O(x^{(1)}, x^{(2)})$ becomes a rectilinear ridge along the axis $x^{(1)} = x^{(2)}$.

question is, whether or not $|\mathbf{r}_j^{(\alpha)} - \mathbf{r}_j^{(\beta)}|$, with $\alpha \neq \beta$, stays finite for a certain fraction of particles. With that in mind, we define the replica density:

$$O(\hat{x}) := \frac{1}{N} \sum_{j=1}^N \delta(\hat{x} - \hat{r}_j) = \frac{1}{N} \sum_{j=1}^N \delta(\mathbf{x}^{(0)} - \mathbf{r}_j^{(0)}) \cdots \delta(\mathbf{x}^{(n)} - \mathbf{r}_j^{(n)}), \quad (2.16)$$

which is the joint probability to find a particle at $\mathbf{x}^{(0)}$ in replica 0, and the same particle at $\mathbf{x}^{(1)}$ in replica 1, ... and at $\mathbf{x}^{(n)}$ in replica n . A thermally averaged example of $O(\hat{x})$, is illustrated in Fig. 2.3a for the case of one spatial dimension and two replicas: A *localized* particle which is found at position \mathbf{x} in one replica, will have a higher probability to be close to that position also in other replicas. The extent of the particle fluctuations, which we call **localization length**, is also constituted by the length scale of these correlations.

As before, after performing the disorder average, the *mean positions* have equal probability to be anywhere in the sample; however, a particle of the macroscopic cluster still has a high probability to be found at *close-by* positions among the different replicas. This situation is illustrated in Fig. 2.3b, again for one dimension and two replicas: Due to translational invariance, the *disorder-averaged* replicated density does not depend on the mean position $x_{\text{cm}} = (x^{(1)} + x^{(2)})/2$, but only on the difference $(x^{(1)} - x^{(2)})/2$ between the replicas. That latter dependence still represents the extent of the particle fluctuations ξ . In Sec. 2.6 we are going to find

an Ansatz for the replicated density field and mathematically formulate the aspects introduced here.

For completeness, we also define the replicated density in Fourier space:

$$O(\hat{q}) := \frac{1}{N} \sum_{j=1}^N \exp(i\hat{q}\hat{r}_j), \quad (2.17a)$$

$$O^{(\alpha)}(\mathbf{q}) := \frac{1}{N} \sum_{j=1}^N \exp(i\mathbf{q}\mathbf{r}_j^{(\alpha)}) = O(\mathbf{0}, \dots, \mathbf{0}, \underset{\uparrow}{\mathbf{q}}, \mathbf{0}, \dots, \mathbf{0}). \quad (2.17b)$$

The first definition is the full Fourier transform of Eq. (2.16) and contains correlations between replicas. The latter definition is the ordinary Fourier density of the particle configuration of a given replica α .

2.4.3 Free Energy of the Replicated Density Field

Now we can go back to Eq. (2.14c) and express the replica free energy in terms of the density fields as introduced in the last section. The calculation is straightforward and done in Appendix B.1.2:

$$\tilde{f}_{n+1}\{\hat{r}_j\} = f_0 - \phi^n \frac{\mu^2}{2} \sum_{\hat{q} \neq \hat{0}} |O(\hat{q})|^2 \Delta(\hat{q}) + \frac{n_0 \lambda}{2} \sum_{\alpha=0}^n \sum_{\mathbf{q} \neq \mathbf{0}} |O^{(\alpha)}(\mathbf{q})|^2 U(\mathbf{q}) \quad (2.18)$$

with the mean particle density $n_0 = N/V$ and the simplifying definitions

$$f_0 := -\phi^n \frac{\mu^2}{2} + (n+1) \frac{\lambda n_0}{2} \quad (2.19)$$

$$\Delta(\hat{q}) := \exp\left(-\frac{\hat{q}^2 a^2}{2}\right) \quad (2.20)$$

The last term of Eq. (2.18) solely comes from the excluded volume interaction. On account of this interaction, density fluctuations cost energy in the same manner in each replica.

In the middle term, the sum also goes over \hat{q} -vectors whose components may involve *different* replicas; hence it comprises the correlations discussed in the previous section. However, at this point it is difficult to see what the immediate effect of this term is. To get a better understanding and to enable further calculations, we introduce different *sets* of \hat{q} -vectors:

We define the **one replica sector** (1RS) as the set of \hat{q} -vectors with exactly one of the $n+1$ components non-zero:

$$1RS = \{\hat{q} \neq \hat{0} \mid \hat{q} = (\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}^{(\alpha)}, \mathbf{0}, \dots, \mathbf{0}) \text{ with } 0 \leq \alpha \leq n\}. \quad (2.21)$$

Complementary, the **higher replica sector** (HRS) is the set of \hat{q} -vectors with at least two replica entries non-zero, and therefore all (non-zero) \hat{q} -vectors which are not in the 1RS:

$$HRS = \{\hat{q} \neq \hat{0} \mid \hat{q} \notin 1RS\}, \quad (2.22)$$

and for the sake of completeness the **zero replica sector** (0RS) only contains $\hat{q} = \hat{0}$:

$$\text{0RS} = \{\hat{0}\} \quad (2.23)$$

With these definitions, Eq. (2.18) becomes:

$$\tilde{f}_{n+1}\{\hat{r}_j\} = f_0 - \phi^n \frac{\mu^2}{2} \sum_{\hat{q} \in \text{HRS}} |\text{O}(\hat{q})|^2 \Delta(\hat{q}) + \frac{1}{2} \sum_{\hat{q} \in \text{1RS}} |\text{O}(\hat{q})|^2 \tilde{\lambda}(\hat{q}) \quad (2.24a)$$

$$\text{with } \tilde{\lambda}(\hat{q}) := n_0 \lambda U(\hat{q}) - \phi^n \mu^2 \Delta(\hat{q}) . \quad (2.24b)$$

The interpretation of $\tilde{\lambda}(\hat{q})$ is straightforward: In the 1RS (which accounts for density fluctuations) the repulsive excluded volume interaction counteracts the attractive cross-links, see (2.24b). Density fluctuations of wavelength \hat{q} cost the more energy, the higher $\tilde{\lambda}(\hat{q})$ is. As mentioned in the beginning, the strength of excluded volume interaction λ has to be large enough to ensure $\tilde{\lambda}(\hat{q}) > 0$ for all $\hat{q} \in \text{1RS}$ to overcome the attractive cross-link interaction;² otherwise density fluctuations $\text{O}(\hat{q})$ of the corresponding wavelength \hat{q} will be energetically favored and the system will become unstable.

2.4.4 Introduction of a Field Theory and Decoupling

It is important to keep in mind that the replicated density $\text{O}(\hat{q})$ is just a simplifying notation. No coarse graining or simplification has been made and in particular, the degrees of freedom of the replica free energy in Eq. (2.24) are still the particle positions $\{\hat{r}_j\}$. Here we will convert the formalism into a field theory, *i.e.* degrees of freedom shall be constituted by a fluctuating density field, rather than by the particle positions.

In the current state, Eq. (2.24a), the density only appears as a quadratic form. If we somehow could make it linear, *i.e.* $f\{\hat{r}_j\} \propto \sum_{\hat{q}} a(\hat{q}) \text{O}(\hat{q})$, the partition function would decouple with respect to the particles:

$$\int \left(\prod_{j=1}^N D\hat{r}_j \right) \exp \left(- \sum_{\hat{q}} a(\hat{q}) \sum_j e^{i\hat{q}\hat{r}_j} \right) = \left(\int D\hat{r} \exp \left(- \sum_{\hat{q}} a(\hat{q}) e^{i\hat{q}\hat{r}} \right) \right)^N \quad (2.25)$$

A way to linearize the quadratic form is the so called **Hubbard-Stratonovich transformation**. The basic idea is that the linear term of a Gaussian integral becomes quadratic after performing the integral:

$$\exp(k^2) \propto \int_{-\infty}^{\infty} dx \exp(-x^2 + kx) . \quad (2.26)$$

The cost for this linearization is the introduction of the integral over a new variable, in this case x . We can make this transformation for each \hat{q} -value³ and obtain the

²that requires that $U(\hat{q})$ does not fall off faster than $\Delta(\hat{q}) = \exp(-a^2 \hat{q}^2/2)$ for $|\hat{q}| \rightarrow \infty$.

³and we have to take into account the correct prefactors and that $\text{O}(\hat{q})$ is complex

following result (see Appendix B.2.1 for the calculation):

$$\mathcal{Z}_{n+1} = \exp(-Nf_0) \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\}) \quad (2.27a)$$

with the *replica free energy*, dependent on the field Ω

$$f_{n+1}\{\Omega\} = \frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) |\Omega(\hat{q})|^2 + \frac{1}{2} \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) |\Omega(\hat{q})|^2 - \ln \mathfrak{z}, \quad (2.27b)$$

with an *effective one-particle partition function*

$$\mathfrak{z} = \int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}} + i \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}}\right), \quad (2.27c)$$

and a constant contribution f_0 as in Eq. (2.19). The field $\Omega(\hat{q})$ runs through every possible configuration, via the newly introduced integral in Eq. (2.27a), with the measure $\mathcal{D}\Omega \propto \prod_{\hat{q}} d\Omega(\hat{q})$. The interpretation of the field $\Omega(\hat{q})$ is the same as the original density $O(\hat{q})$, since the difference between them vanishes in the thermodynamic limit (as seen in Appendix B.2.2). Specifically $\Omega(\hat{q}) = O(\hat{q})$ for $\hat{q} \in \text{HRS}$ and $\Omega(\hat{q}) = iO(\hat{q})$ for $\hat{q} \in \text{IRS}$. Furthermore, as anticipated in Eq. (2.25), the integral over all particle positions is reduced to a single integral over one “effective” particle in Eq. (2.27c).

This is an important stage of the calculation, because **so far no approximation has been made**. In this work we are not interested in density fluctuations which would be caused by a finite bulk modulus. Therefore, we now take the limit of infinitely strong excluded volume interaction, $\lambda \rightarrow \infty$ and thus $\tilde{\lambda}(\hat{q}) \rightarrow \infty$, to prevent any density fluctuations, i.e. $\Omega(\hat{q} \in \text{IRS}) = 0$. The resulting replica free energy then has a simple form, which only involves the higher replica sector:

$$f_{n+1}\{\Omega\} = \frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) |\Omega(\hat{q})|^2 - \ln \mathfrak{z}, \quad (2.28a)$$

$$\text{with } \mathfrak{z} = \int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}}\right) \quad (2.28b)$$

2.4.5 Universality and Connection to Other Models

For many systems in statistical mechanics it is possible to establish a Landau-Wilson free energy, which depends on a coarse-grained order parameter. A famous example is the Ising model with the order parameter being the coarse-grained magnetization [Goldenfeld, 1992, Chapter 5]. If this order parameter vanishes as a phase transition is approached, one often expands the Landau-Wilson free energy to lowest necessary order.

As discussed before in Sec. 2.4.2, the replicated density $\Omega(\hat{q})$ is an order parameter that can distinguish between the solid and liquid phase. In the liquid phase $\Omega(\hat{q} \neq \hat{0}) = 0$, and close to the transition in the gel phase, we may suspect $\Omega(\hat{q} \neq \hat{0})$ to be small, since most of the particles are delocalized and do not give rise to correlations between replicas. Hence we can establish a Ginzburg-Landau like expansion of our model. This expansion of $f_{n+1}\{\Omega\}$, Eq. (2.28), can easily be done (see Appendix B.3). To lowest necessary order in Ω and \hat{q} (\hat{q} corresponding to gradients in real-space representation), the replica free energy is:

$$f_{n+1}\{\Omega\} = \sum_{\hat{q} \in \text{HRS}} \left(-A\varepsilon + \frac{B}{2}\hat{q}^2 \right) |\Omega(\hat{q})|^2 - C \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Omega(\hat{q}_1)\Omega(\hat{q}_2)\Omega(\hat{q}_3)\delta_{\hat{q}_1+\hat{q}_2+\hat{q}_3, \hat{0}} + \mathcal{O}(\Omega^4, \Omega^3\hat{q}^2, \Omega^2\hat{q}^4) \quad (2.29)$$

with

$$\varepsilon = \mu^2 - 1, \quad A = \frac{\mu^2}{2}, \quad B = \left(\mu^4 - \frac{\mu^2}{2} \right) a^2, \quad C = \frac{\mu^6}{6}. \quad (2.30)$$

In this simple form (2.29), one can already understand some basic properties of the system: the **sol-gel transition** happens when the parameter $\varepsilon = \mu^2 - 1$ changes sign: For $\varepsilon < 0$, $\Omega(\hat{q}) = 0$ is a stable solution, since the prefactor of the quadratic term is positive and any small deviation from this solution costs more energy. For $\varepsilon \gtrsim 0$, this solution becomes unstable and small deviations from $\Omega(\hat{q}) = 0$ are energetically favorable, whereby the cubic term is necessary to stabilize the new solution. The gradient term ($\propto \hat{q}^2 |\Omega(\hat{q})|^2$) is the lowest order that involves spatial dependencies; close to the transition, the relevant length scales of the system (like the above-mentioned localization length) typically diverge. Hence $\Omega(\hat{q})$ should be very small for large $|\hat{q}|$, which justifies neglecting higher orders of \hat{q} close to the transition.

Eq. (2.29) is a *universal* free energy of an incompressible random system close to the sol-gel transition: This form can also be obtained by pure symmetry considerations and can be used as a phenomenological starting point for the replica free energy [Peng *et al.*, 1998]. With the *randomly cross-linked particle model* one has a very simple *microscopic* model that can reproduce this phenomenological free energy and can provide values for the constants A , B and C . Other models, like cross-linked semiflexible chains, also reproduce Eq. (2.29) with different values for A , B and C [Goldbart *et al.*, 2004; Mukhopadhyay *et al.*, 2004]. Later we will perform this expansion to *infinite* order to see how this model behaves far from the gelation transition.

2.5 Goldstone Fluctuations in General

In the next section, we will see that the saddle point solution for $\Omega(\hat{q})$ has a **continuous symmetry**; specifically that means that there is a *whole family* of saddle-point

solutions, which are parameterized by *homogeneous translations* of the entire system between the replicas. We want to establish an Ansatz for the order parameter, which allows small *long-wavelength* and *low-energy* fluctuations (so called **Goldstone fluctuations**) around the saddle point.

Before getting to this Ansatz, it is helpful to have a rough understanding of Goldstone fluctuations in general (see also e.g. [Goldenfeld, 1992, chapter 11] or [Goldstone *et al.*, 1962]). In this regard, let us first consider these Goldstone modes for an easier model, the so called *XY-model*: In that model, two-dimensional spins \mathbf{s}_i of unit length can rotate in a plane and reside on a D -dimension lattice (where D can be different from 2); they interact such that neighboring spins (“ $\langle \rangle$ ”) preferably align:

$$H(\{\mathbf{s}_i\}) = -J \sum_{\langle i,j \rangle} \mathbf{s}_i \cdot \mathbf{s}_j = -J \sum_{\langle i,j \rangle} \cos(\theta_i - \theta_j), \quad (2.31)$$

where θ_i is the angle of spin i with respect to an arbitrary axis. $J > 0$ is the strength of the interaction or can be interpreted as the inverse temperature. One can easily see that in the ground state, all spins point in the same direction.

The important feature for us is that the system has a **continuous symmetry**, i.e. all spins can continuously be rotated at the same time without energy cost. Or in other words, the transformation $\theta_i \rightarrow \theta_i + \phi$ for all i and arbitrary ϕ leaves the Hamiltonian (2.31) invariant.

For $D > 2$, **spontaneous symmetry breaking** occurs when reducing the temperature below a critical value T_c [Goldenfeld, 1992]. In that case, the system chooses a direction, along which all spins preferably align, yielding a non-zero magnetization $\mathbf{m} = \frac{1}{N} \sum_{i=1}^N \mathbf{s}_i \neq \mathbf{0}$ (see Fig. 2.4, left). On account of the above-mentioned continuous symmetry of the Hamiltonian, this direction can be rotated *without energy cost* (for example by applying an infinitesimal field).

What about the case that the axis of magnetization is *slowly varying in space* (Fig. 2.4, right)? Reasoning by continuity, one might expect that the energy associated with these **long-wavelength fluctuations** of the angle of the spins have **low energy cost**. Indeed one finds that in the long-wavelength limit, the cost of these fluctuations vanishes with the wave vector \mathbf{k} like $\mathcal{O}(\mathbf{k}^2)$.

These long-wavelength and low-energy modes associated with a continuous symmetry are the **Goldstone modes**. They are a general phenomenon described by *Goldstone’s theorem*, which states that in the presence of a spontaneously broken continuous symmetry there are long-wavelength excitations, whose energy vanishes continuously in the limit $\mathbf{k} \rightarrow \mathbf{0}$.

2.6 Ansatz for the Order Parameter

In Sec. 2.4 we calculated the disorder-averaged free energy $[F_C] = -[\ln Z_C] = -\lim_{n \rightarrow 0} (\mathcal{Z}_{n+1} - \mathcal{Z}_1)/(n\mathcal{Z}_1)$ with help of the replica technique. The replica partition function $\mathcal{Z}_{n+1} \propto \int \mathcal{D}\Omega e^{-Nf_{n+1}\{\Omega\}}$ is represented as a functional integral over the order-parameter field Ω .

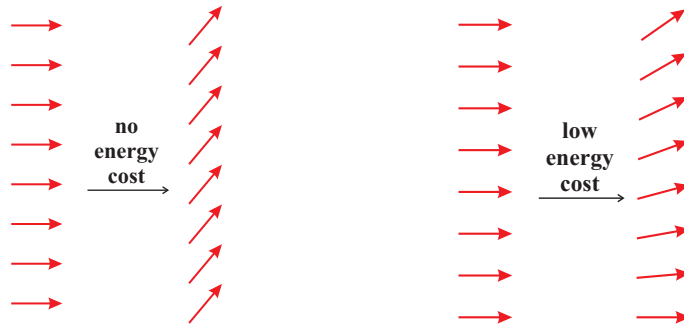


Figure 2.4: Illustration of the XY -model. Only one spatial dimension of the lattice is shown and spins are depicted as red arrows (\rightarrow). Left: A rotation of all spins by a certain angle is an exact symmetry of the Hamiltonian (2.31) and has no energy cost. Right: If the rotation is slowly varying in space, *i.e.* a long-wavelength fluctuation, the energy is arbitrarily low.

As seen in Appendix B.2.2, in the thermodynamic limit this order parameter equals the replicated density $O(\hat{q})$ already addressed in Sec. 2.4.2. The real-space representation of the order parameter

$$\Omega(\hat{x}) = \frac{1}{N} \sum_{i=1}^N \delta(\mathbf{x}^{(0)} - \mathbf{r}_i^{(0)}) \cdots \delta(\mathbf{x}^{(n)} - \mathbf{r}_i^{(n)}) \quad (2.32)$$

quantifies the probability of finding a particle at position $\mathbf{x}^{(0)}$ in replica 0, at $\mathbf{x}^{(1)}$ in replica 1, etc.

In previous work, [Castillo *et al.* \[1994\]](#) have shown that the **saddle-point solution** for the order parameter, obtained by setting $\delta f_{n+1}\{\Omega\}/\delta\Omega = 0$, has the following simple form⁴:

$$\bar{\Omega}_{\text{sp}}(\hat{x}) = \frac{1-Q}{V^{n+1}} + \frac{Q}{V} \int_V d\mathbf{y} \int_0^\infty \frac{d\xi^2 \mathcal{P}(\xi^2)}{(2\pi\xi^2)^{D(n+1)/2}} \exp\left(-\sum_{\alpha=0}^n \frac{(\mathbf{x}^{(\alpha)} - \mathbf{y})^2}{2\xi^2}\right), \quad (2.33)$$

with self-consistent equations for Q and $\mathcal{P}(\xi^2)$ given in [[Castillo *et al.*, 1994](#); [Broderix *et al.*, 2002](#)]. This form has a very intuitive interpretation, since the following premises, applied to Eq. (2.32), *also* reproduce (2.33): A fraction of monomers $(1-Q)$ is delocalized and can be found anywhere in the sample with equal probability. The remaining fraction Q is the “infinite” cluster, in which each particle i performs Gaussian fluctuations about a replica-independent mean position \mathbf{y}_i with localization length ξ_i . (The situation is qualitatively depicted in Fig. 2.3a in Sec. 2.4.2.) The mean positions \mathbf{y}_i are randomly distributed (Fig. 2.3b), and the localization lengths ξ follow the distribution $\mathcal{P}(\xi^2)$.

⁴[Castillo *et al.* \[1994\]](#) use a slightly different model of cross-linked chains, which belongs to the same universality class as the presented model. The saddle-point value obtained for our model is the same.

2.6.1 Spontaneously Broken Translational Invariance

The effective replicated Hamiltonian is invariant under uniform translations of all particles, **separately in each replica**, see e.g. Eq. (2.14c), since the particle positions only appear in terms of the form $(\mathbf{r}_i^{(\alpha)} - \mathbf{r}_j^{(\alpha)})$; i.e., the transformation

$$\mathbf{r}_i^{(\alpha)} \rightarrow \mathbf{r}_i^{(\alpha)} + \mathbf{u}^{(\alpha)} \quad (2.34)$$

is an exact symmetry of the Hamiltonian.⁵

Now, let us have a look at the saddle point solution (2.33): In the amorphous solid phase, $Q > 0$, this symmetry is spontaneously broken, and the transformation (2.34), in general, yields a different order parameter; the only symmetries remaining are **common displacements of all replicas**, i.e., $\mathbf{r}_i^{(\alpha)} \rightarrow \mathbf{r}_i^{(\alpha)} + \mathbf{u}$, reflecting the macroscopic translational invariance of the amorphous solid phase. (For a detailed discussion see [Goldbart *et al.*, 2004; Mukhopadhyay *et al.*, 2004].) Consequently, there is a **whole family of order parameters** which are all related by **replica-dependent, spatially uniform translations**. All of them give rise to the same free energy and the system will “spontaneously” choose one of them. To distinguish common and relative displacements of the replicas, it is helpful to split up the $D(n+1)$ -dimensional hatted vectors \hat{x} into D -dimensional longitudinal ones

$$\mathbf{x}_{\parallel} = \frac{1}{\sqrt{n+1}} \sum_{\alpha=0}^n \mathbf{x}^{(\alpha)}, \quad (2.35)$$

and Dn -dimensional transverse ones

$$x_{\perp} = \hat{x} - \frac{1}{\sqrt{n+1}} (\mathbf{x}_{\parallel}, \dots, \mathbf{x}_{\parallel}). \quad (2.36)$$

Properties of this division into \parallel - and \perp -vectors can be found in Appendix A. A very useful relation to keep in mind is

$$\hat{x} \cdot \hat{y} = \mathbf{x}_{\parallel} \cdot \mathbf{y}_{\parallel} + x_{\perp} \cdot y_{\perp}. \quad (2.37)$$

When considering a parallel-vector like \mathbf{x}_{\parallel} , one should keep in mind that it is not exactly the *center of mass* $\mathbf{x}_{\text{cm}} = \frac{1}{n+1} \sum_{\alpha} \mathbf{x}^{(\alpha)}$ of its replica components, but $\mathbf{x}_{\parallel} = \sqrt{n+1} \mathbf{x}_{\text{cm}}$. Only this definition enables the simple relation (2.37). As a consequence, \parallel -vectors reside in a different volume $V_{\parallel} := V(n+1)^{D/2}$, rather than V . For consistency, we also define a parallel-vector $\mathbf{R}_{\parallel} := \sqrt{n+1} \mathbf{R}$ for any *non-replicated* vector \mathbf{R} .

With the help of these definitions we can parameterize the manifold of symmetry-related saddle points by u_{\perp} :

$$\bar{\Omega}_{u_{\perp}}(\hat{x}) = \frac{1-Q}{V^{n+1}} + \frac{Q}{V_{\parallel}} \int_0^{\infty} \frac{d\xi^2 \mathcal{P}(\xi^2)}{(2\pi\xi^2)^{Dn/2}} \exp\left(-\frac{(x_{\perp} - u_{\perp})^2}{2\xi^2}\right), \quad (2.38)$$

⁵The effective replicated Hamiltonian is also invariant under uniform rotations, separately in each replica. However the spontaneous breaking of this symmetry does not give rise to Goldstone modes and is not discussed further here.

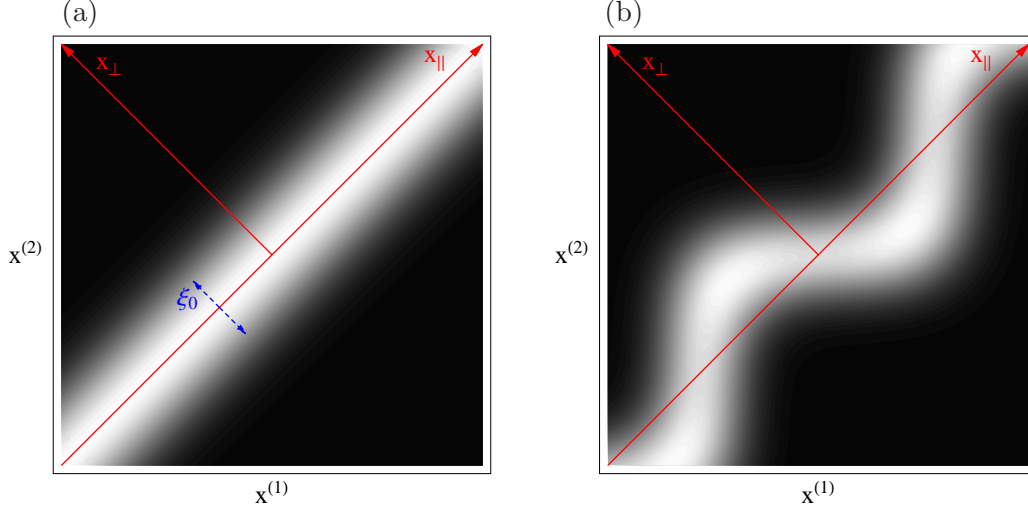


Figure 2.5: Ansatz for the order parameter $\Omega(x^{(1)}, x^{(2)})$ in real space for two replicas in one dimension, as Fig. 2.3. To simplify, we have assumed a fixed localization length ξ_0 in this figure. $\Omega(x^{(1)}, x^{(2)})$ is the probability of finding a particle at position $x^{(1)}$ in replica 1 and at $x^{(2)}$ in replica 2. For the rotated coordinate system $x_{\parallel} = (x^{(2)} + x^{(1)})/\sqrt{2}$ and $x_{\perp} = (x^{(2)} - x^{(1)})/2$. In (a) the displacement vector $u_{\perp} \equiv 0$, and in (b) $u_{\perp} = \sin x_{\parallel}$. (Note, however, that for physical systems typically $\xi \ll$ wavelength of u_{\perp} .)

where $u_{\perp} = \hat{0}$ yields Eq. (2.33). The situation is illustrated in Fig. 2.5a, where the order parameter is shown for two replicas in one space-dimension. The order parameter does not depend on \mathbf{x}_{\parallel} , so that it is represented by a rectilinear ridge having its maximum height along the axis $x_{\perp} = \hat{0}$ and having width ξ . The family of symmetry-related solutions is generated by rigidly translating the ridge to a new position: $x_{\perp} = u_{\perp}$. (The delocalized particles contribute only an unimportant constant background $(1 - Q)/V^{n+1}$, which has been ignored in the figure.)

2.6.2 Goldstone Fluctuations and Ansatz for Ω

Now, we want to establish an **Ansatz for the order parameter** that allows for special **fluctuations around the saddle point**. These fluctuations are low-energy and long-wavelength excitations (*i.e.* Goldstone modes) and are constructed by particle displacements $u_{\perp}(\mathbf{x}_{\parallel})$ that depend on the position in the sample [Mukhopadhyay *et al.*, 2004; Goldbart *et al.*, 2004]. These *local* deformations may be generated by thermal fluctuations in the system. The Ansatz developed in this section was already briefly presented in [Ulrich *et al.*, 2006].

For the more general Ansatz, we make the same basic premises that reproduced the saddle point solution (2.33): The fraction $(1 - Q)$ of particles is delocalized, and a particle j of the remaining fraction Q (the *gel fraction*) performs Gaussian fluctuations around a mean position $\mathbf{R}_j^{(\alpha)}$. Now, however, these mean positions

depend on the replica and may be shifted by a spatially dependent value $\mathbf{u}_\perp^{(\alpha)}(\mathbf{x}_\parallel)$ from one another between the replicas, such that $\mathbf{R}_j^{(\alpha)} = \mathbf{R}_j + \mathbf{u}_\perp^{(\alpha)}(\mathbf{R}_{j\parallel})$. Hence the probability to find particle j at position \mathbf{x} in replica α is

$$\begin{aligned} \langle \delta(\mathbf{x} - \mathbf{r}_j^{(\alpha)}) \rangle &= \\ &= \begin{cases} \frac{1}{(2\pi\xi_j^2)^{D/2}} \exp\left(-\frac{(\mathbf{x} - (\mathbf{R}_j + \mathbf{u}_\perp^{(\alpha)}(\mathbf{R}_{j\parallel})))^2}{2\xi_j^2}\right) & \text{if particle } j \text{ localized (loc.)} \\ 1/V & \text{if particle } j \text{ delocalized (deloc.)} \end{cases} \end{aligned} \quad (2.39)$$

and thus, for all replicas, the joint probability density to find a particle at $\mathbf{x}^{(0)}$ in replica (0), at $\mathbf{x}^{(1)}$ in replica (1), etc. is

$$\langle \delta(\hat{x} - \hat{r}_j) \rangle = \begin{cases} \frac{1}{(2\pi\xi_j^2)^{D(n+1)/2}} \exp\left(-\frac{(\hat{x} - \hat{R}_j)^2}{2\xi_j^2}\right) & \text{if } j \text{ loc.} \\ V^{-(n+1)} & \text{if } j \text{ deloc.} \end{cases} \quad (2.40)$$

with $\hat{R}_j = (\mathbf{R}_j^{(0)}, \dots, \mathbf{R}_j^{(n)}) = (\mathbf{R}_j + \mathbf{u}_\perp^{(0)}(\mathbf{R}_{j\parallel}), \dots, \mathbf{R}_j + \mathbf{u}_\perp^{(n)}(\mathbf{R}_{j\parallel}))$. The Fourier transform of this distribution is given by:

$$\begin{aligned} \langle \exp(i\hat{q}\hat{r}_j) \rangle &= \\ &= \int d\hat{x} \exp(i\hat{q}\hat{x}) \langle \delta(\hat{x} - \hat{r}_j) \rangle = \begin{cases} \exp\left(-\frac{\hat{q}^2\xi_j^2}{2} + i\hat{q}\hat{R}_j\right) & \text{if } j \text{ loc.} \\ \delta_{\hat{q},\hat{0}} & \text{if } j \text{ deloc.} \end{cases} \end{aligned} \quad (2.41)$$

Hence, we get for the averaged order parameter:

$$\begin{aligned} \langle \Omega(\hat{q}) \rangle &= \frac{1}{N} \sum_{j=1}^N \langle \exp(i\hat{q}\hat{r}_j) \rangle \\ &= \frac{1}{N} \sum_{j \text{ loc.}} \exp\left(-\frac{\hat{q}^2\xi_j^2}{2} + i \sum_{\alpha=0}^n \mathbf{q}^{(\alpha)} \cdot (\mathbf{R}_j + \mathbf{u}_\perp^{(\alpha)}(\mathbf{R}_{j\parallel}))\right) + \frac{1}{N} \cdot \mathcal{N}(1-Q)\delta_{\hat{q},\hat{0}} \end{aligned} \quad (2.42)$$

Now, to incorporate the disorder average, we assume a uniform distribution of the *undeformed* mean positions \mathbf{R}_j . We also define the distribution for the localization lengths

$$\mathcal{P}(\xi^2) = \frac{1}{QN} \sum_{j \text{ loc.}} \delta(\xi^2 - \xi_j^2) \quad (2.43)$$

and get:

$$\begin{aligned} \bar{\Omega}_u(\hat{q}) - (1-Q)\delta_{\hat{q},\hat{0}} &= \frac{1}{N} \sum_{j \text{ loc.}} \exp\left(-\frac{\hat{q}^2\xi_j^2}{2}\right) \int \frac{d\mathbf{R}_j}{V} \exp\left(i \sum_{\alpha=0}^n \mathbf{q}^{(\alpha)} \cdot (\mathbf{R}_j + \mathbf{u}_\perp^{(\alpha)}(\mathbf{R}_{j\parallel}))\right) \\ &= \frac{Q}{V_\parallel} \int_0^\infty d\xi^2 \mathcal{P}(\xi^2) \exp\left(-\frac{\hat{q}^2\xi^2}{2}\right) \int d\mathbf{x}_\parallel \exp(i\mathbf{q}_\parallel \mathbf{x}_\parallel + i\mathbf{q}_\perp \cdot \mathbf{u}_\perp(\mathbf{x}_\parallel)) \end{aligned} \quad (2.44)$$

which can be written in a simple way:

$$\bar{\Omega}_u(\hat{q}) = (1 - Q)\delta_{\hat{q},\hat{0}} + Q\omega(\hat{q}) \int D\mathbf{x}_{\parallel} \exp(i\mathbf{q}_{\parallel}\mathbf{x}_{\parallel} + iq_{\perp}u_{\perp}(\mathbf{x}_{\parallel})) \quad (2.45a)$$

$$\text{with } \omega(\hat{q}) = \int_0^{\infty} d\xi^2 \mathcal{P}(\xi^2) \exp\left(-\frac{\hat{q}^2\xi^2}{2}\right) \quad (2.45b)$$

and with the dimensionless measure $D\mathbf{x}_{\parallel} = d\mathbf{x}_{\parallel}/V_{\parallel}$. In real space this result transforms into

$$\bar{\Omega}_u(\hat{x}) = \frac{1 - Q}{V^{n+1}} + Q \int_0^{\infty} \frac{d\xi^2 \mathcal{P}(\xi^2)}{(2\pi\xi^2)^{D(n+1)/2}} \times \int D\mathbf{y}_{\parallel} \exp\left(-\frac{(\mathbf{y}_{\parallel} - \mathbf{x}_{\parallel})^2 + (x_{\perp} - u_{\perp}(\mathbf{y}_{\parallel}))^2}{2\xi^2}\right). \quad (2.46)$$

This form is illustrated in Fig. 2.5b: Unlike the homogeneous translations in Eq. (2.38) and Fig. 2.5a, the ridge is not translated rigidly, but deformed by the position-dependent vector $u_{\perp}(\mathbf{x}_{\parallel})$. These excitations *do* cost energy; only in the long-wavelength limit do we expect the energy of the excitation to vanish with an inverse power of the wavelength.

We only consider **pure shear deformations**, and hence require that $|\det(\delta_{\tau\nu} + \partial_{\tau}u_{\perp}^{(\alpha,\nu)})| = 1$. This constraint is consistent with taking the excluded volume interaction to be large, which makes the system incompressible. For small deformations, the constraint simplifies to $\partial_{\nu}u_{\perp}^{(\alpha,\nu)} = 0$.

It is important to note that the deformations in the Ansatz, Eq. (2.45), represent **small fluctuations around the saddle-point solution** (2.33). As we see in Appendix B.4, in the limit of *small deformations* u_{\perp} and *long wavelengths* (i.e. small \hat{q}), the Ansatz can be expressed as

$$\bar{\Omega}_u(\hat{q}) = \bar{\Omega}_{\text{sp}}(\hat{q}) + \delta\Omega_u(\hat{q}), \quad (2.47)$$

with a small $\delta\Omega_u(\hat{q}) = \mathcal{O}(q_{\perp}u_{\perp})$. Consequently, for $u_{\perp}(\mathbf{x}_{\parallel}) \equiv \hat{0}$, the Ansatz reproduces the saddle-point solution (2.33) (here in Fourier representation):

$$\bar{\Omega}_{u\equiv 0}(\hat{q}) \equiv \bar{\Omega}_{\text{sp}}(\hat{q}) = (1 - Q)\delta_{\hat{q},\hat{0}} + Q\omega(\hat{q})\delta_{\mathbf{q}_{\parallel},\mathbf{0}}. \quad (2.48)$$

2.6.3 Physical Shear Deformations

Here we examine what the *physical meaning* of the shear deformation field $u_{\perp}(\mathbf{x}_{\parallel})$ is. Let us first assume, there is a particle at position \mathbf{x} in the unsheared state ($u_{\perp} = \hat{0}$, see Fig. 2.6, left). If there is a non-zero $u_{\perp}(\mathbf{x}_{\parallel})$, this particle will be moved to the position $\mathbf{x} + \mathbf{u}_{\perp}^{(0)}(\mathbf{x}_{\parallel})$ in the zeroth replica. In this replica cross-linking takes place,

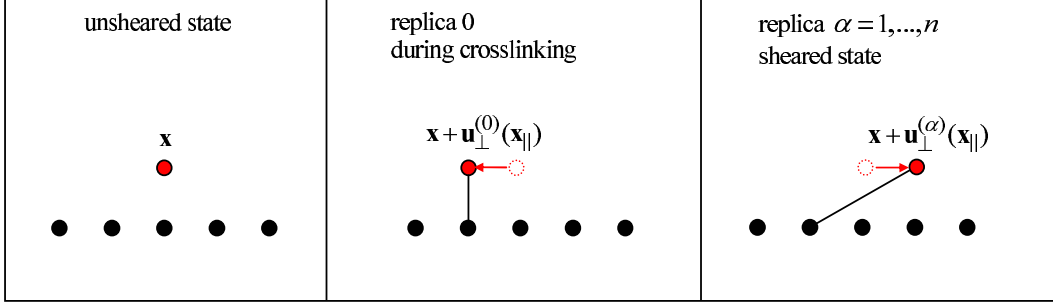


Figure 2.6: Descriptive illustration of the deformation vector $u_{\perp}(\mathbf{x}_{\parallel})$. Left: In the unsheared state, there is a particle at position \mathbf{x} . Center: In the zeroth replica the deformation causes the particle to be at the position $\mathbf{x} + \mathbf{u}_{\perp}^{(0)}(\mathbf{x}_{\parallel})$. It will preferably be cross-linked to particles close to that position. Right: In replicas $\alpha = 1, \dots, n$, that particle is moved to $\mathbf{x} + \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel})$, but still cross-linked to particles that were close in replica (0).

hence this particle will preferably be cross-linked to particles close to that position (Fig. 2.6, center). The measurement is performed in replicas $\alpha = 1, \dots, n$, in which the particle is moved to the position $\mathbf{x} + \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel})$ (Fig. 2.6, right). However, the physically relevant deformation is the displacement of this particle relative to the state during *cross-linking*, $\mathbf{u}_{\text{phys}}^{(\alpha)}(\mathbf{x}_{\parallel}) = \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel}) - \mathbf{u}_{\perp}^{(0)}(\mathbf{x}_{\parallel})$. Hence it is useful to introduce the *physical* deformation field

$$\mathbf{u}_{\text{phys}}^{(\alpha)}(\mathbf{x}_{\parallel}) := \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel}) - \mathbf{u}_{\perp}^{(0)}(\mathbf{x}_{\parallel}), \quad \alpha = 0, \dots, n \quad (2.49)$$

which has n non-zero components and contains the same information as $u_{\perp}(\mathbf{x}_{\parallel})$.

One can obtain a useful relation between the two fields:

$$\sum_{\alpha=0}^n \left(\frac{\partial u_{\perp}^{(\alpha, \nu)}}{\partial x_{\parallel}^{(\tau)}} \right)^2 = \sum_{\alpha=1}^n \left(\frac{\partial u_{\text{phys}}^{(\alpha, \nu)}}{\partial x_{\parallel}^{(\tau)}} \right)^2 - \frac{1}{n+1} \left(\sum_{\alpha=1}^n \frac{\partial u_{\text{phys}}^{(\alpha, \nu)}}{\partial x_{\parallel}^{(\tau)}} \right)^2, \quad (2.50)$$

which can easily be proved by plugging in the definition of $\mathbf{u}_{\text{phys}}^{(\alpha)}(\mathbf{x}_{\parallel})$, Eq. (2.49). In the above equation $\nu, \tau = 1, \dots, D$ are spatial indices. We note that the sum of the last term is of the form $(\sum_{\alpha=1}^n \dots)^2 = \mathcal{O}(n^2)$ and therefore we can write in the limit $n \rightarrow 0$:

$$\boxed{\sum_{\alpha=0}^n \left(\frac{\partial u_{\perp}^{(\alpha, \nu)}}{\partial x_{\parallel}^{(\tau)}} \right)^2 = \sum_{\alpha=1}^n \left(\frac{\partial u_{\text{phys}}^{(\alpha, \nu)}}{\partial x_{\parallel}^{(\tau)}} \right)^2 + \mathcal{O}(n^2)} \quad (2.51)$$

2.7 Replica Free Energy with Shear Deformations

Now we will insert the Ansatz (2.45) into the general expression (2.28) for the free energy and thereby allow for physically motivated fluctuations (namely the local

shear displacements $u_{\perp}(\mathbf{x}_{\parallel})$ around the saddle point:

$$\frac{Z_{n+1}}{\exp(-Nf_0)} = \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\}) \approx \int \mathcal{D}u \exp(-Nf_{n+1}\{\bar{\Omega}_u\}) \quad (2.52)$$

The integral $\int \mathcal{D}u$ goes over all possibilities of the local shear field, separately in all replicas.

In this approach, we will only consider small, long-wavelength distortions, so that it is sufficient to keep u_{\perp} only to the lowest order, $(\partial u_{\perp}/\partial \mathbf{x}_{\parallel})^2$. Higher orders in u_{\perp} and higher-order derivatives will be neglected. The calculation is somewhat intricate and can be found in Appendix B.5. The result for the replica free energy can be written in a very concise way as:

$$Nf_{n+1}\{\bar{\Omega}_u\} = Nf_{n+1}^{\text{sp}} + \frac{G}{2} \int d\mathbf{x}_{\parallel} \sum_{\alpha=0}^n \sum_{\tau,\nu=1}^D \left(\frac{\partial u_{\perp}^{(\alpha,\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 + \mathcal{O}(n^2), \quad (2.53a)$$

where the first term, f_{n+1}^{sp} , is the **saddle-point value** of $f_{n+1}\{\Omega\}$:

$$f_{n+1}^{\text{sp}} = n \left\{ \frac{D}{2} \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) + \frac{\mu^2 Q^2}{2} \Xi_{2,a^2/2} - \exp(-\mu^2 Q) \sum_{r=2}^{\infty} \frac{(\mu^2 Q)^r}{r!} \Xi_{r,a^2} \right\}, \quad (2.53b)$$

and the second term describes the cost of **fluctuations (i.e. shear deformations) around the saddle point**, with the shear modulus

$$G = \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) n_0. \quad (2.53c)$$

For the saddle-point value Eq. (2.53b), Ξ_{r,a^2} is a simplifying definition that depends on the microscopic parameters of the system:

$$\Xi_{r,a^2} := \int d\xi_1^2 \mathcal{P}(\xi_1^2) \cdots d\xi_r^2 \mathcal{P}(\xi_r^2) \ln \left(\left(\frac{V}{(2\pi)^{D/2}} \right)^{r-1} \left(\frac{\prod_{\gamma=1}^r (a^2 + \xi_{\gamma}^2)^{-1}}{\sum_{\gamma=1}^r (a^2 + \xi_{\gamma}^2)^{-1}} \right)^{D/2} \right). \quad (2.54a)$$

In the thermodynamic limit, when $V^{1/D} \gg \xi, a$, the dependence on ξ and a vanishes, and Ξ_{r,a^2} reduces to (see Eq. (B.101) in Appendix B.5.6):

$$\Xi_{r,a^2} = (r-1) \ln N + \mathcal{O}(N^0). \quad (2.54b)$$

One has to keep in mind, however, that for some purposes the full form (2.54a) is needed, for example, for determining the distribution of localization lengths $\mathcal{P}(\xi^2)$

as e.g. in [Goldbart *et al.*, 1996, Sec. VI C]. For all our purposes, however, the simple Eq. (2.54b) is sufficient. We plug it into Eq. (2.53b) and use (B.93) for the sum over r . With that, f_{n+1}^{SP} becomes very simple:

$$f_{n+1}^{\text{SP}} = n \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) \left(\frac{D'}{2} - \ln N + \mathcal{O}(N^0) \right) \quad (2.55)$$

It is noteworthy that this saddle point solution of the free energy *per particle* f_{n+1}^{SP} is not intensive, but proportional to $\ln N$. This point has already been discussed in [Goldbart *et al.*, 1996]: it is most probably coming from the fact that for a given cross-link configuration \mathcal{C} , certain particles are indistinguishable (e.g. if particles 1, 2 and 3 are connected as a triangle, all three of those particles can be interchanged yielding the same configuration). Therefore, if we insist on an intensive free energy, we have to normalize the partition function by a prefactor taking into account this indistinguishability. However, until now, this could not be done for this or similar models and does not matter if one considers $\partial f / \partial Q = 0$.

2.8 Returning to the Disorder-Averaged Free Energy

In the last section, we found the *replica free energy* $f_{n+1}\{\bar{\Omega}_u\}$ and its dependence on the shear deformations $u_\perp(\mathbf{x}_\parallel)$, see Eqs. (2.53). Here we want to establish the connection to the “real” *disorder-averaged free energy* $[F_{\mathcal{C}}]$. With Eqs. (2.14a) and (2.10), we can relate the replicated partition function \mathcal{Z}_{n+1} to the disorder-averaged free energy:

$$\begin{aligned} -[F_{\mathcal{C}}] &= \lim_{n \rightarrow 0} \frac{[\mathcal{Z}_{\mathcal{C}}^n] - 1}{n} = \lim_{n \rightarrow 0} \frac{\mathcal{Z}_{n+1}/\mathcal{Z}_1 - 1}{n} = \frac{1}{\mathcal{Z}_1} \lim_{n \rightarrow 0} \frac{\mathcal{Z}_{n+1} - \mathcal{Z}_1}{n} \\ &= \frac{1}{\mathcal{Z}_1} \lim_{n \rightarrow 0} \frac{\partial \mathcal{Z}_{n+1}}{\partial n} = \lim_{n \rightarrow 0} \frac{\partial \ln \mathcal{Z}_{n+1}}{\partial n} \end{aligned} \quad (2.56)$$

and hence:

$$\boxed{[F_{\mathcal{C}}] = - \lim_{n \rightarrow 0} \frac{\partial}{\partial n} \ln \mathcal{Z}_{n+1} .} \quad (2.57)$$

Applying this relation to \mathcal{Z}_{n+1} from Eqs. (2.52) and (2.53a), we find:

$$\begin{aligned} [F_{\mathcal{C}}] &= N \lim_{n \rightarrow 0} \frac{\partial}{\partial n} (f_0 + f_{n+1}^{\text{SP}}) \\ &\quad + \lim_{n \rightarrow 0} \frac{\partial}{\partial n} \ln \int \mathcal{D}u \exp \left(-\frac{G}{2} \sum_{\alpha} \sum_{\nu, \tau} \int d\mathbf{x}_\parallel \left(\frac{\partial u_{\perp}^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_{\parallel}^{(\tau)}} \right)^2 \right) . \end{aligned} \quad (2.58)$$

The only degrees of freedom left are constituted by the displacement field $u_\perp(\mathbf{x}_\parallel)$. However, u_\perp contains the mathematically unpleasant constraint that the sum over all replica components has to be zero. With Eq. (2.51), we can relate u_\perp to the

physical shear deformation \hat{u}_{phys} and write the integral $\int \mathcal{D}u$ as an integral over the n components of \hat{u}_{phys} :

$$\begin{aligned}
[F_{\mathcal{C}}] &= N \lim_{n \rightarrow 0} \frac{\partial(f_0 + f_{n+1}^{\text{SP}})}{\partial n} \\
&= - \lim_{n \rightarrow 0} \frac{\partial}{\partial n} \ln \left\{ \int \left(\prod_{\alpha=1}^n \mathcal{D}\mathbf{u}_{\text{phys}}^{(\alpha)} \right) \exp \left(-\frac{G}{2} \sum_{\alpha=1}^n \sum_{\nu, \tau} \int d\mathbf{x}_{\parallel} \left(\frac{\partial u_{\text{phys}}^{(\alpha, \nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 \right) \right\} \\
&= - \lim_{n \rightarrow 0} \frac{\partial}{\partial n} \ln \left(\left\{ \int \mathcal{D}\mathbf{u}_{\text{phys}} \exp \left(-\frac{G}{2} \sum_{\nu, \tau} \int d\mathbf{x}_{\parallel} \left(\frac{\partial u_{\text{phys}}^{(\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 \right) \right\}^n \right).
\end{aligned} \tag{2.59}$$

We can get n -th power of the curly bracket out of the logarithm, and obtain:

$$[F_{\mathcal{C}}] = N \lim_{n \rightarrow 0} \frac{\partial(f_0 + f_{n+1}^{\text{SP}})}{\partial n} - \ln Z^{\text{el}} \tag{2.60a}$$

$$\text{with } Z^{\text{el}} = \int \mathcal{D}\mathbf{u}_{\text{phys}} \exp \left(-\frac{G}{2} \sum_{\nu, \tau} \int d\mathbf{x} \left(\frac{\partial u_{\text{phys}}^{(\nu)}(\mathbf{x})}{\partial x^{(\tau)}} \right)^2 \right) \tag{2.60b}$$

Hereby, Z^{el} is the partition function of the energy contribution of an incompressible medium [Landau & Lifshitz, 1965]. It does not involve replicas anymore, and thus we can confirm that G indeed plays the role of the shear modulus. It is analyzed in more detail in Sec. 2.9.3.

We can see in Eq. (2.60a) that, in addition to the elastic contribution, the *linear order in n* of the saddle point solution f_{n+1}^{SP} adds to the disorder-averaged free energy per particle.

2.9 Results

2.9.1 Average Cross-Link Density and Higher Moments

Given the Deam-Edwards distribution Eq. (2.8), it is possible to calculate expectation values of the form $[M(M-1)\cdots(M-J+1)]$ with the total number of cross-links M in the system and the disorder average $[\cdot]$. In Appendix B.6.1 we find that

$$[M(M-1)\cdots(M-J+1)] = \left(\frac{\mu^2 N}{2} \right)^J. \tag{2.61}$$

Of particular importance is of course the case $J=1$ and $J=2$, from which the first two moments can be calculated. For those two cases, the above equation yields:

$$[M] = \frac{\mu^2 N}{2} \tag{2.62a}$$

$$[M(M-1)] = \left(\frac{\mu^2 N}{2} \right)^2. \tag{2.62b}$$

Thus for any choice of μ^2 , the cross-link density is:

$$\boxed{\frac{[M]}{N} = \frac{\mu^2}{2}} \quad (2.63)$$

and with two particles involved in every cross-link, the parameter $\mu^2 = 2[M]/N$ is the **average coordination number**.

The fluctuations of the number of cross-links are calculated from Eqs. (2.62) with

$$\begin{aligned} \Delta M^2 &:= [M^2] - [M]^2 = [M(M-1)] + [M] - [M]^2 \\ &= \left(\frac{\mu^2 N}{2}\right)^2 + \frac{\mu^2 N}{2} - \left(\frac{\mu^2 N}{2}\right)^2 = \frac{\mu^2 N}{2}, \end{aligned} \quad (2.64)$$

and hence

$$\boxed{\Delta M = \sqrt{[M^2] - [M]^2} = \sqrt{\mu^2 N/2}.} \quad (2.65)$$

As one can easily see, for any finite μ^2 , the fluctuations $\Delta M/[M]$ relative to the mean number of cross-links vanish like $N^{-1/2}$.

In principle, all moments can be calculated very easily using the formula (2.61). Indeed, it turns out that all moments are equal to those of the Poisson distribution with parameter $\mu^2 N/2$. [Broderix *et al.*, 2002, sec. 3.1] also found a convergence of the cross-link distribution towards a Poisson distribution for the randomly cross-linked particle model with a slightly different cross-link distribution, without the use of the replica technique. Instead of the parameter μ^2 , in that work the probability for a certain cross-link is solely determined by the typical cross-linking length.

2.9.2 Gel Fraction

In order to obtain a relation for the gel fraction Q , we need to find the minimum of the free energy with respect to Q . Therefore, we calculate

$$\frac{\partial}{\partial Q} \frac{\partial f_{n+1}^{\text{SP}}}{\partial n} = -(-\mu^2 \exp(-\mu^2 Q) + \mu^2 - \mu^2 Q) \ln N \stackrel{!}{=} 0, \quad (2.66)$$

which becomes an implicit equation for Q :

$$\boxed{1 - Q = e^{-\mu^2 Q}} \quad (2.67)$$

This equation is well known and has been derived in many contexts like random graph percolation, random nets and theory of epidemics [Solomonoff & Rapoport, 1951; Landau, 1952; Erdős & Rényi, 1960, 1961], as well as similar gelling systems [Broderix *et al.*, 2002; Goldbart *et al.*, 1996; Castillo *et al.*, 1994].

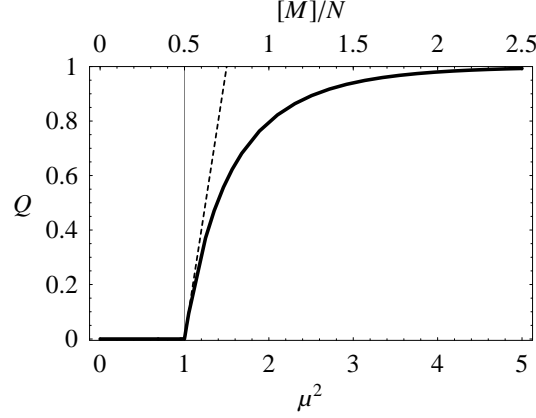


Figure 2.7: Fraction of localized particles Q , depending on the average coordination number. The infinite cluster forms at the critical value $\mu_c^2 = 1$. The full line is the exact solution of Eq. (2.67) and the dashed line the approximation (2.69).

Given that the relation for the gel fraction (2.67) is already well known, we will only briefly discuss it here: It has a trivial solution $Q = 0$, which is stable for $\mu^2 \leq 1$ and unstable for $\mu^2 > 1$. At $\mu^2 = \mu_c^2 = 1$, there is a second order transition to a gel state $Q > 0$, as for $\mu^2 > 1$ a further, nontrivial stable solution emerges. This is the *sol-gel transition*, already briefly mentioned in Sec. 2.4.5.

In Fig. 2.7, the stable branch of the gel fraction Q is plotted versus the cross-link density. At a critical value $\mu_c^2 = 1$, one can see the transition from a sol phase, with all particles delocalized ($Q = 0$), to a gel phase, where certain particles are localized ($Q > 0$).

Close to that transition, Eq. (2.67) can be expanded for $\mu^2 \approx 1$, $Q \ll 1$ [Peng *et al.*, 1998; Goldbart *et al.*, 2004; Mukhopadhyay *et al.*, 2004]:

$$\begin{aligned}
 1 - Q &= 1 - \mu^2 Q + \frac{1}{2} \mu^4 Q^2 + \mathcal{O}(Q^3) \\
 -1 &= -\mu^2 + \frac{1}{2} \mu^4 Q + \mathcal{O}(Q^2) \\
 Q &= 2(\mu^2 - 1) + \mathcal{O}(Q^2)
 \end{aligned} \tag{2.68}$$

And with $\varepsilon = \mu^2 - 1$, the distance from the critical point:

$$Q = 2\varepsilon + \mathcal{O}(\varepsilon^2) \tag{2.69}$$

In this work we are interested in the behavior for *arbitrary* μ^2 . In this regard, we rewrite Eq. (2.67) as:

$$\begin{aligned}
 1 &= (1 - Q)e^{\mu^2 Q} \\
 -\mu^2 e^{-\mu^2} &= \mu^2(Q - 1)e^{\mu^2(Q-1)}.
 \end{aligned} \tag{2.70}$$

This form can be written in terms of the Lambert function $W(x)$, which is the solution of the equation $x = W(x)e^{W(x)}$, see [Corless *et al.*, 1996], and thus we get:

$$\begin{aligned}\mu^2(Q-1) &= W(-\mu^2 e^{-\mu^2}) \\ Q &= 1 + \frac{W(-\mu^2 e^{-\mu^2})}{\mu^2}.\end{aligned}\quad (2.71)$$

With the help of the expansion of the Lambert function,

$$W(-x) = -\sum_{k=1}^{\infty} \frac{k^{k-1}}{k!} x^k \quad \text{for } |x| < \frac{1}{e}, \quad (2.72)$$

we get

$$\boxed{Q = 1 - e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k \quad \text{for all } \mu^2 > 1.} \quad (2.73)$$

2.9.3 Shear Modulus

In Eq. (2.60), we identified the shear modulus G . Restoring units of energy, it becomes:

$$G = \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) n_0 k_B T \quad (2.74)$$

As one would expect, it vanishes in the liquid phase ($\mu^2 < 1$, $Q = 0$). In the solid phase ($\mu^2 > 1$), it increases linearly with particle density n_0 and depends only on the cross-link density, given the gel fraction $Q = Q(\mu^2)$ (derived in the previous section 2.9.2).

Close to the gel transition $\mu^2 \gtrsim 1$, the shear modulus can be expanded for small Q :

$$\begin{aligned}G &= \left(1 - \mu^2 Q + \frac{(\mu^2 Q)^2}{2!} - \frac{(\mu^2 Q)^3}{3!} + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) n_0 k_B T + \mathcal{O}(Q^4) \\ &= \left(\frac{\mu^2(\mu^2 - 1)Q^2}{2} - \frac{\mu^6 Q^3}{6} \right) n_0 k_B T + \mathcal{O}(Q^4),\end{aligned}\quad (2.75)$$

and with Eq. (2.69), we can write the shear modulus in terms of $\varepsilon = \mu^2 - 1$, resulting in a power-law behavior (see Fig. 2.8a):

$$\boxed{G = \frac{2}{3} n_0 k_B T \varepsilon^3 + \mathcal{O}(\varepsilon^4).} \quad (2.76)$$

In the **highly cross-linked regime**, we can also write the shear modulus as a power series. To this end, we start from Eq. (2.74) again and use the (exact) relation (2.67)

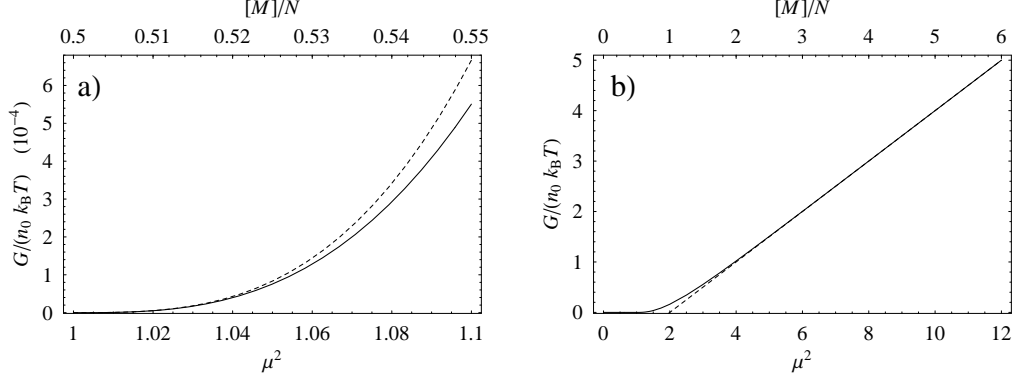


Figure 2.8: Shear modulus G divided by the particle density n_0 and $k_B T$, depending on the average coordination number μ^2 . The solid line shows the exact expression (2.74), the dashed lines are the approximations valid (a) close to the sol-gel transition, Eq. (2.76), and (b) for highly connected systems, Eq. (2.79).

to get rid of the exponential:

$$\begin{aligned} \frac{G}{n_0 k_B T} &= (1 - Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \\ &= \frac{\mu^2}{2} - 1 + (1 - Q) - \frac{\mu^2}{2} (1 - Q)^2. \end{aligned} \quad (2.77)$$

The expansion for $(1 - Q)$ is already given in Eq. (2.73). In Appendix B.6.2, we find an expression for the square of this infinite sum, $(1 - Q)^2$, with which we easily obtain a relation for $(1 - Q) - \frac{\mu^2}{2} (1 - Q)^2$, see Eq. (B.118). With that, the shear modulus Eq. (2.77) can be written in a series in $\mu^2 e^{-\mu^2}$,

$$\frac{G}{n_0 k_B T} = \frac{\mu^2}{2} - 1 + e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-2}}{k!} (\mu^2 e^{-\mu^2})^k, \quad (2.78)$$

which can be used to compute G for all $\mu^2 > 1$. In the highly cross-linked regime $\mu^2 \gg 1$, G is approximated by (see Fig. 2.8b):

$$G = \frac{1}{2} n_0 k_B T (\mu^2 - 2) + \mathcal{O}(e^{-\mu^2}). \quad (2.79)$$

The scaling $G \sim \varepsilon^3$ close to the critical point has been derived previously [Castillo & Goldbart, 1998; Mukhopadhyay *et al.*, 2004; Goldbart *et al.*, 2004]; the critical exponent is a result of mean field theory and expected to be modified by interacting fluctuations below $D = 6$ [Peng & Goldbart, 2000]. In the highly cross-linked limit, fluctuations are presumably weak so that our result is expected to hold even beyond the Gaussian expansion around mean field theory. The result we get is reminiscent of the classical theory of rubber elasticity, $G = n_s k_B T$, with the density of strands n_s being replaced by the density of cross-links: $\frac{1}{2} \mu^2 n_0 = [M]/V$.

In order to compare the calculated shear modulus with experimental results, note that, if the cross-link length is small compared to the particle size, it is unlikely for one monomer in $D = 3$ ($D = 2$) dimensions to be connected to more than 12 (6) other monomers. Therefore, average coordination numbers $\mu^2 > 12$ (> 6) are inaccessible in these systems. Only if long tethers can form, reaching further than the nearest neighbor, higher average coordination numbers μ^2 are possible.

2.10 Conclusions

We have determined the shear modulus G of a randomly cross-linked system. It only depends on the particle density $n_0 = N/V$ and the average coordination number μ^2 . Interestingly, neither the distribution of localization lengths $\mathcal{P}(\xi^2)$, defined in Eq. (2.43), nor the typical cross-link length a influence the macroscopic behavior of the material. This result is consistent with the observation that the shear modulus determines the response of the system to a shear deformation on the longest scales: The wavelength of the shear deformation has to be larger than all microscopic length scales. Furthermore, one observes that $G \propto k_B T$, *i.e.* the shear elasticity is purely entropic. This is a consequence of our simple model with entropic springs to account for the cross-links. In more complex systems, *e.g.* covalently bonded random networks, one expects to find additional energetic contributions to G in the highly cross-linked regime. In any case, the bulk modulus should depend on the strength of the excluded-volume interaction, and hence it is not expected to be as universal as the shear modulus.

It is also noteworthy that the shear modulus does not depend on the spatial dimension D , apart from the fact that the maximal coordination numbers depend on dimensionality and determine the physical range of μ^2 . Again, this could be expected, because we only consider a Gaussian expansion around the saddle point; interactions between fluctuations will most likely cause a dependence on dimensionality.

Cross-Linked Directed Polymers

Contents

| | | |
|------------|--|-----------|
| 3.1 | Introduction | 31 |
| 3.2 | Properties of a Single Chain | 33 |
| 3.2.1 | Radius of Gyration of an Uncrosslinked Chain | 33 |
| 3.2.2 | Properties of a Polymer Clamped in Space | 34 |
| 3.3 | Full Model: Cross-Linked Directed Polymers | 36 |
| 3.3.1 | Interactions | 36 |
| 3.3.2 | Disorder Average | 39 |
| 3.4 | Replica Calculation of the Free Energy | 40 |
| 3.4.1 | Disorder-Averaged Free Energy: First Steps | 40 |
| 3.4.2 | Introduction of the Replicated Density Field | 41 |
| 3.4.3 | Introduction of a Field Theory and Decoupling | 43 |
| 3.4.4 | Saddle Point Equation | 44 |
| 3.5 | Ansatz for the Order Parameter | 45 |
| 3.6 | The Saddle-Point Equation with Ansatz for Ω | 47 |
| 3.6.1 | Expansion to second order in Q | 48 |
| 3.6.2 | Sol-Gel Transition | 49 |
| 3.7 | The Equation for the Localization Length | 49 |
| 3.7.1 | Normalization of Length Scales | 49 |
| 3.7.2 | Result for the Distribution of Localization Lengths | 50 |
| 3.8 | Conclusions | 55 |

3.1 Introduction

In the previous chapter we investigated a system of cross-linked molecular networks. Neither the particles nor the cross-linking gave rise to a preferred direction, hence on a macroscopic scale, the system was isotropic. In many cases, however, the experimental setup induces a **preferred direction of the polymer chains**, like e.g. for so called polymer brushes, where strands are mounted perpendicular to a bottom plate, or DNA strands which are highly stretched along one direction. For polymer chains in a nematic solvent or nematic polymers [Kamien *et al.*, 1992], the alignment and hence the emergence of a preferred direction can also be due

to spontaneous symmetry breaking, whereby the chains align in an arbitrary but uniform direction, when e.g. the temperature is decreased below a critical value. Hence, the field of *directed polymers* has become an active area of research during the last twenty years.

This problem of directed chains is also relevant for the behavior of type-II superconductors [Kamien *et al.*, 1992; Marchetti & Nelson, 1993]. A magnetic field, larger than a critical value H_{c1} , penetrates the superconductor in form of *flux lines*, which are preferably aligned along the field and each of which is carrying one quantum of magnetic flux. Due to their mutual repulsion, these flux lines can form various lattice or glassy structures in the transverse plane. In this respect, the flux lines are similar to directed polymer chains with excluded volume interaction.

Recently, there has been an increasing interest in **cross-linked** polymer brushes with promising applications. This is due to the fact that cross-linking can influence chemical and mechanical stability, permeability and swelling characteristics of the polymer brushes [Huang *et al.*, 2001; Loveless *et al.*, 2006]. For example Ionov *et al.* [2004] investigated binary polymer brushes, consisting of both hydrophobic and hydrophilic chains; they managed to cross-link the hydrophobic components in the extended or collapsed state, depending on the position of the sample; thereby they could locally, on a μm -scale, tailor the properties of the material. On the other hand, amphiphilic block copolymers (*i.e.* polymer chains with one hydrophobic and one hydrophilic end) in an aqueous solution are known to form structures like micelles, lipid bilayers, or liposomes, whose interior comprises brush-like structures. For micelles Iijima *et al.* [1999] and Xu *et al.* [2004] also found that cross-linking can be used to achieve stabilization. Stable micelles are of great importance for pharmaceutical applications, since they can be utilized as carriers for targeted drug delivery.

In this chapter we investigate a simple model of flexible cross-linked directed polymers, embedded between two parallel surfaces (see Fig. 3.1). The polymer chains are modeled as functions $\mathbf{r}(z)$, which are preferably aligned along the distinguished direction z . This preferential alignment may be interpreted as a mechanical stretching of the chains along the z -direction, or as the interaction with a nematic field. We will study the effect of permanent cross-links on this simple structure and observe a gelation transition as the cross-links concentration is increased above a critical value, similar to the RLP model.

In the following section 3.2, we will discuss the behavior of one single polymer chain. The full model, comprising a large ensemble of these chains, is introduced in Sec. 3.3. In Sec. 3.4 we develop a Landau-Wilson free energy, with the replicated density being the order parameter. In Sec. 3.5 an Ansatz for the order parameter is developed, which is shown to solve the saddle-point equation in Sec. 3.6; the sol-gel transition, a consequence of the saddle-point equation, is investigated there (see Sec. 3.6.2). The distribution for the localization lengths is derived in Sec. 3.7, and we will conclude in Sec. 3.8.

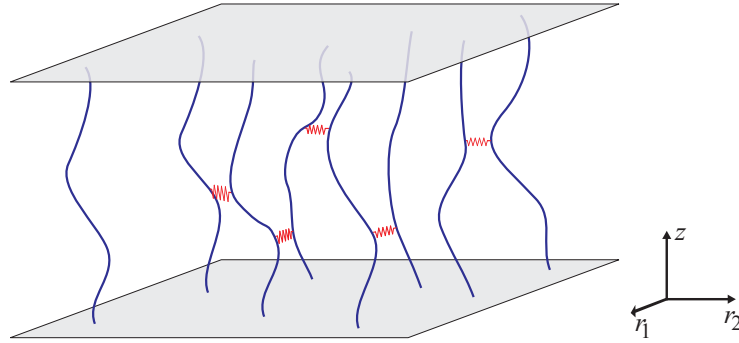


Figure 3.1: Illustration of the cross-linked directed polymer model. The polymers are depicted as blue lines ($\color{blue}{\text{~}}\color{blue}{\text{~}}\color{blue}{\text{~}}$). They are preferably aligned along the z -axis and are randomly cross-linked ($\color{red}{\text{~}}\color{red}{\text{~}}\color{red}{\text{~}}$). We will refer to the directions *perpendicular* to the axis of alignment as the *transverse* or *in-plane* direction (here r_1 and r_2).

3.2 Properties of a Single Chain

Before starting with the system of cross-linked directed polymers, it is illustrative to investigate the behavior of one single polymer chain. Such a chain is modeled as a D -dimensional vector $\mathbf{r}(z)$ that depends on an additional dimension, the system height z . We will have a look at fluctuations of a fully uncrosslinked chain (Sec. 3.2.1) as well as a chain clamped to a certain point in space (Sec. 3.2.2).

3.2.1 Radius of Gyration of an Uncrosslinked Chain

Consider a single free polymer chain $\mathbf{r}(z)$ only subject to the alignment interaction

$$H_{\text{align},1} = \frac{\sigma}{2} \int_0^L dz \dot{\mathbf{r}}^2(z), \quad (3.1)$$

which penalizes deviations from the alignment along the z -axis. Throughout this chapter about directed polymers, the index $_1$ on any Hamiltonian specifies that it is acting only on one chain, to avoid confusion with Hamiltonians acting on the whole system of N chains, investigated later. The probability for a conformation $\mathbf{r}(z)$ is given by the Boltzmann weight $e^{-H_{\text{align},1}}$. Here we want to calculate the average extension of this chain in \mathbf{r} -direction, or in other words, its **transverse radius of gyration**:

$$R_g^2 := \left\langle \frac{1}{2L^2} \int_0^L dz_1 dz_2 (\mathbf{r}(z_1) - \mathbf{r}(z_2))^2 \right\rangle_{H_{\text{align},1}}. \quad (3.2)$$

The average $\langle \cdot \rangle_{H_{\text{align},1}}$ is represented by a functional integral over all chain conformations. For an arbitrary observable \mathcal{A} ,

$$\langle \mathcal{A} \rangle_{H_{\text{align},1}} = \int \mathcal{D}\mathbf{r}(z) \mathcal{A} e^{-H_{\text{align},1}}. \quad (3.3)$$

The measure $\mathcal{D}\mathbf{r}(z)$ of all functional integrals in this thesis is normalized such that

$$\int \mathcal{D}\mathbf{r}(z) e^{-H_{\text{align},1}} = 1 \quad (3.4)$$

That is why Eq. (3.3) does not need a normalization. In Appendix C.1.1, we solve Eq. (3.2) by a discretization of the functional integral and get for the radius of gyration:

$$\boxed{R_g^2 = \frac{D}{6} \frac{L}{2\sigma}} \quad (3.5)$$

We will see that the quantity

$$\boxed{\ell^2 := \frac{L}{2\sigma} \propto R_g^2} \quad (3.6)$$

will be of great importance in the full system of *cross-linked* polymer chains, which we investigate later.

3.2.2 Properties of a Polymer Clamped in Space

As a next step we calculate the properties of a single polymer chain which is clamped to a fixed point $(\mathbf{0}, z_c)$ in space, see Fig. 3.2, left. Hence the chain $\mathbf{r}(z)$ is subject to the following two interactions $H_1 = H_{\text{align},1} + H_{\text{clamp},1}$ with:

$$H_{\text{align},1} = \frac{\sigma}{2} \int_0^L dz \dot{\mathbf{r}}^2(z) \quad (3.7)$$

$$H_{\text{clamp},1} = \frac{1}{2a^2} (\mathbf{r}(z_c) - \mathbf{0})^2 \quad (3.8)$$

To investigate properties of this simple system, we introduce the partition function with an auxiliary field \mathbf{b} :

$$Z = \int \mathcal{D}\mathbf{r}(z) \exp(-H_1 + \mathbf{b} \cdot \mathbf{r}(z_0)) , \quad (3.9)$$

so that we can easily compute the mean fluctuations of the chain at height z_0 :

$$\nabla_{\mathbf{b}}^2 \ln Z = \nabla_{\mathbf{b}} \left(\frac{\nabla_{\mathbf{b}} Z}{Z} \right) = \frac{\nabla_{\mathbf{b}}^2 Z}{Z} - \frac{(\nabla_{\mathbf{b}} Z)^2}{Z^2} = \langle \mathbf{r}^2(z_0) \rangle - \langle \mathbf{r}(z_0) \rangle^2 =: \Delta \mathbf{r}^2(z_0) . \quad (3.10)$$

The functional integral in Eq. (3.9) can be solved by a discretization, similar to one in the previous section. In Appendix C.1.2 we find:

$$\ln Z = \frac{\mathbf{b}^2}{2} \left(\frac{|z_c - z_0|}{\sigma} + a^2 \right) + \ln \left(\frac{(2\pi a^2)^{D/2}}{A} \right) . \quad (3.11)$$

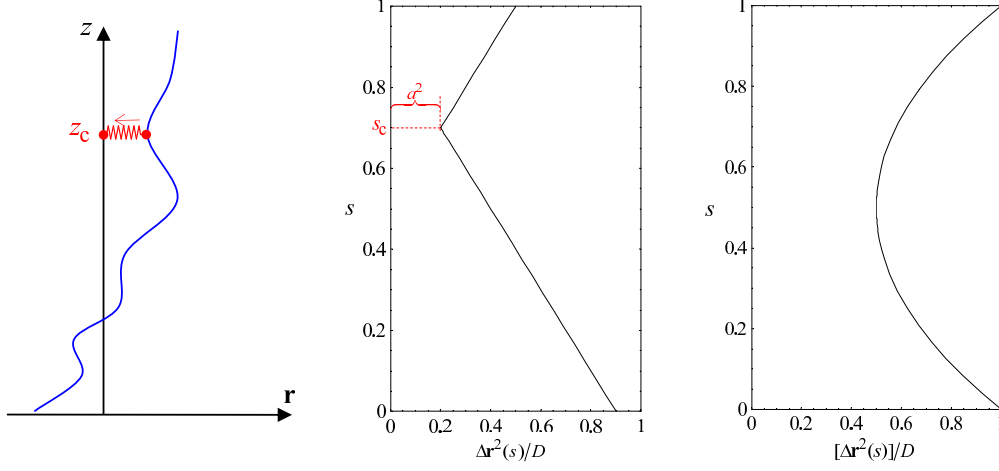


Figure 3.2: Left: Illustration of the clamped polymer chain. The polymer (λ) is clamped (wavy) to the z -axis at a certain height z_c . Center: Plot of the size of the fluctuations $\Delta \mathbf{r}^2(s)$, see Eq. (3.12), exemplarily for the case $a^2 = 0.2$, $s_c = 0.7$. Right: Disorder-averaged fluctuations $[\Delta \mathbf{r}^2(s)] \propto s^2 + (1 - s)^2$, according to Eq. (3.14) for the case $a^2 = 0$.

With that result we can apply Eq. (3.10) to get the fluctuations of the chain (for simplification we change z_0 to z):

$$\Delta \mathbf{r}^2(z) = D \cdot \left(\frac{|z_c - z|}{\sigma} + a^2 \right) = D \cdot \left(\frac{L}{\sigma} |s_c - s| + a^2 \right). \quad (3.12)$$

On the right hand side of the above equation, we introduced the normalized system height $s = z/L$. This result is illustrated in Fig. 3.2, center. As one can see, the extent of the fluctuations increases linearly, as we move away from the “clamping height” z_c . At the clamping height itself, $z = z_c$, fluctuations are still present due to the finite cross-link strength $1/a^2$.

In the full system, the chains will be cross-linked to each other at *random heights*. Similarly, we may define a disorder average $[\cdot]$ for this simple system as the uniform distribution of all clamping heights z_c (resp. s_c), so that the disorder-averaged fluctuations may be interpreted as the average of many (non-interacting) chains clamped to random points on the z -axis:

$$\begin{aligned} [\Delta \mathbf{r}^2(s)] &:= \int_0^1 ds_c \Delta \mathbf{r}^2(s) = D \left(\frac{L}{\sigma} \int_0^1 ds_c |s_c - s| + a^2 \right) \\ &= D \left(\frac{L}{\sigma} \cdot \frac{s^2 + (1 - s)^2}{2} + a^2 \right). \end{aligned} \quad (3.13)$$

Using the radius of gyration $\ell^2 = L/(2\sigma)$ from the previous section 3.2.1 we get:

$$\boxed{\frac{[\Delta \mathbf{r}^2(s)]}{D} = \ell^2 (s^2 + (1 - s)^2) + a^2,} \quad (3.14)$$

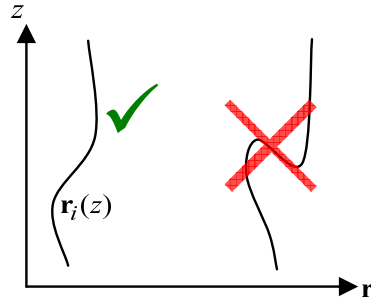


Figure 3.3: The polymer chains are modeled as functions $\mathbf{r}_i(z)$. Thus, at every height z , a chain can only appear once (as on the left) and cannot turn around in z direction (as on the right).

which is illustrated in Fig. 3.2 on the right side. It might be surprising that in spite of the homogeneous distribution of clamping points, the fluctuations are not uniform along the z -axis. This behavior arises from the fact that the chain has open ends, which can move around freely. We will see a similar behavior also for the cross-linked polymers and discuss it more extensively in Sec. 3.7.2.

3.3 Full Model: Cross-Linked Directed Polymers

In the rest of this chapter, we consider a system of N identical chains. As in the previous section, the positions of these chains are modeled as D -dimensional functions $\mathbf{r}_1(z), \dots, \mathbf{r}_N(z)$ which depend on the “height” z in the system, see Fig. 3.3, left chain. Thus, the dimensionality of the system is $D+1$. This way of modeling distinguishes the z -direction such that the chains cannot “turn around” and go back in that direction as the chain on the right hand side in Fig. 3.3.

The chains reside in a volume $V = L \cdot A$, where L is the total height in z -direction and A is the area (or hyper-area) of the system in the other D spatial dimensions. Most important is, of course, the case $D = 2$ for a three-dimensional system. The thermodynamic limit is taken such that $N, A \rightarrow \infty$ with a constant chain density $n_0 = N/A$. However, we keep the L -dependence, since the free ends of the chain may cause a z -dependence of the structural and mechanical properties that provides interesting physics. It is interesting to note about this model that the limit $L \rightarrow 0$ (a thin D -dimensional “slice”) reproduces the D -dimensional RLP model.

3.3.1 Interactions

The chains are subject to the following interactions:

The **cross-link interaction** permanently connects a pair of chains (i_e and j_e) at a fixed height ($z_e \in [0, L]$), which is the same for both constituents (see left side of Fig. 3.4). As in the randomly cross-linked particle model, the M cross-links are modeled as Hookian springs. Thus, for a given cross-link configuration

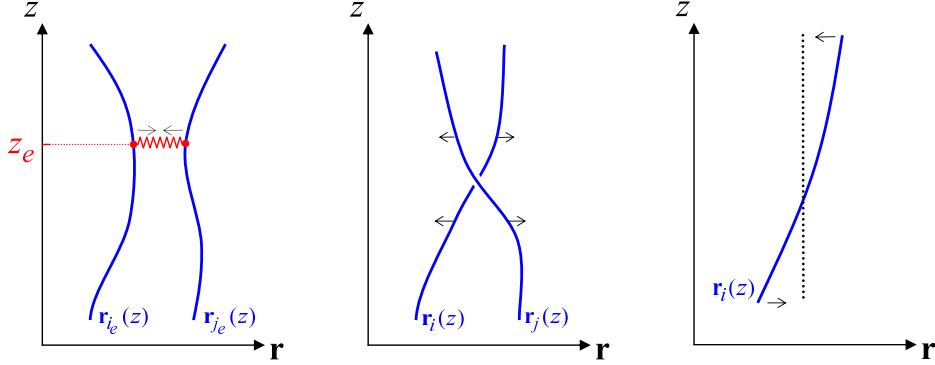


Figure 3.4: Illustration of the interactions between the directed polymers (λ). Left: A cross-link ($\color{red}{\text{wavy line}}$) permanently connects chains i_e and j_e at height z_e . Center: Through the excluded volume interaction, chains which come too close are repelled from each other. Right: The alignment interaction straightens the chains along the z axis by penalizing high derivatives $\dot{\mathbf{r}}_i(z)$.

$\mathcal{C} := \{(i_e, j_e, z_e)\}_{e=1}^M$, the cross-link Hamiltonian is:

$$H_{\text{Xlink}} = \frac{1}{2a^2} \sum_{e=1}^M (\mathbf{r}_{i_e}(z_e) - \mathbf{r}_{j_e}(z_e))^2, \quad (3.15)$$

where a is the typical cross-linking length.

As in the RLP model, a **repulsive excluded-volume interaction** is introduced, to prevent the system from collapsing. Here, it acts between all pairs of chains and along the whole length of them (see Fig. 3.4, center):

$$H_{\text{ev}} = \frac{\lambda}{2} \sum_{i,j=1}^N \int_0^L dz U(\mathbf{r}_i(z) - \mathbf{r}_j(z)), \quad (3.16)$$

with a short ranged $U(\mathbf{x})$ subject to the restriction $\int_A U(\mathbf{x}) d\mathbf{x} = 1$.

Furthermore, we add an **alignment interaction**, which straightens the chains along the z -axis by penalizing high values of the derivatives $\dot{\mathbf{r}}_i(z)$ (Fig. 3.4, right):

$$H_{\text{align}} = \frac{\sigma}{2} \sum_{i=1}^N \int_0^L dz \dot{\mathbf{r}}_i^2(z) \quad (3.17)$$

In Sec. 3.2, we have already seen the effect of this interaction on a *single* chain.

From a physical point of view, this interaction can be interpreted as a means to penalize stretching of the chains: While the unperturbed length, *i.e.* the length of a

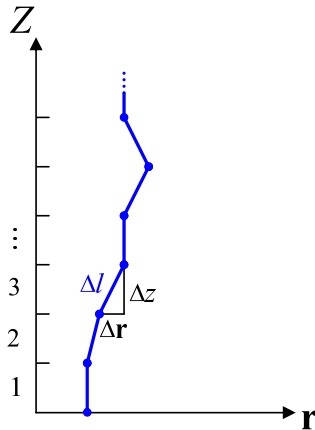


Figure 3.5: Discretization of a polymer chain into segments $Z = 1, 2, \dots, Z_L$. The length of one chain segment Δl depends on the length of the segment in z -direction (Δz) and in transverse direction ($\Delta \mathbf{r}$).

chain with $\mathbf{r}(z) \equiv \mathbf{0}$, equals the system height L , any fluctuation $\mathbf{r}(z) \neq \mathbf{0}$ stretches the chain such that the arc length L_{arc} becomes larger than L : Discretizing the chain into Z_L segments of height $\Delta z = L/Z_L$ (see Fig. 3.5), we get for its arc length:

$$L_{\text{arc}} = \sum_{Z=1}^{Z_L} \Delta l_Z = \sum_{Z=1}^{Z_L} \sqrt{\Delta z^2 + \Delta \mathbf{r}_Z^2} = \sum_{Z=1}^{Z_L} \Delta z \sqrt{1 + \frac{\Delta \mathbf{r}_Z^2}{\Delta z^2}}$$

$$\stackrel{\Delta z \rightarrow 0}{=} \int_0^L \sqrt{1 + \dot{\mathbf{r}}^2(z)} dz = L + \frac{1}{2} \int_0^L \dot{\mathbf{r}}^2(z) dz + \mathcal{O}(\dot{\mathbf{r}}^4). \quad (3.18)$$

Thus, to lowest order in $\dot{\mathbf{r}}(z)$, the alignment Hamiltonian (3.17) comprises the excess length $L_{\text{arc}} - L = \frac{1}{2} \int_0^L \dot{\mathbf{r}}^2(z) dz$, of all chains. This discretization of a chain into small segments $Z = 1, 2, \dots, Z_L$ will appear frequently throughout the investigation of this model and is a useful way to visualize and calculate functional integrals.

A more sophisticated approach [Marko & Siggia, 1995] shows that this Hamiltonian is also obtained for a **worm-like chain** (a model of semiflexible and inextensible chains), if it is **strongly stretched along the z -axis** by applying an external force f to the ends of the chain. With our notation the energy of this stretched worm-like chain can be written as $E = \frac{1}{2} \int_0^{L_{\text{arc}}} dl (A \dot{\mathbf{r}}^2(l) + f \mathbf{r}^2(l))$, where A is the persistence length. If the force on the chain ends f is strong enough, the bending contribution can be neglected. Changing the integral over the arc length $\int_0^{L_{\text{arc}}} dl$ to an integral over the stretching direction $\int_0^L dz$ we get a transformation factor $dl/dz = \sqrt{1 + \dot{\mathbf{r}}^2(z)} = 1 + \mathcal{O}(\dot{\mathbf{r}}^2)$. Thus this strongly stretched worm-like chain reproduces to lowest order the alignment Hamiltonian (3.17) with σ playing the role of f .

Another physical realization of Eq. (3.17) is that of **worm-like chains interacting with a strong nematic field**, in the *limit of vanishing persistence length* of the chains [Kamien *et al.*, 1992]. In this context, σ controls the coupling of a

(homogeneous) nematic matrix to the local polymer direction. Different from the interpretation of stretched chains, polymers in a nematic field would in general be allowed to make hairpin turns, since the nematic field is typically symmetric under flipping the direction of the director field $\mathbf{n} \rightarrow -\mathbf{n}$; the energy of a hairpin is estimated to be $E_h \approx \sqrt{\sigma\kappa}$ (with the bending stiffness κ) [de Gennes, 1982; Nelson, 2002]. In this context it is important to note that our model is an approximation, since such hairpins cannot be modeled by a function $\mathbf{r}(z)$. However, since we are interested in the limit in which coupling strength σ is dominating further terms, hairpins are energetically unfavorable and can safely be neglected if $\sigma \gg \kappa^{-1}$ is ensured.

The alignment interaction (3.17) is also necessary from a mathematical point of view, because there is no other interaction that makes sure that the chains are continuous in z -direction. Without the alignment interaction, every slice in z -direction could be treated independently and system would decouple.

The alignment interaction has the convenient form of a **sum over all individual chains**,

$$H_{\text{align}} = \sum_{j=1}^N H_{\text{align},1}\{\mathbf{r}_j\}, \quad (3.19)$$

which, as we will see, is easy to handle in the calculation. It is even possible, without impeding the calculation, to add further interactions of that form. Beyond the alignment interaction, further possible terms proposed in Appendix C.2.

The total Hamiltonian is $H_C = H_{\text{Xlink}} + H_{\text{ev}} + H_{\text{align}}$. For the partition function, we have to use functional integrals over all possible conformations of all chains

$$Z_C = \int \mathcal{D}\mathbf{r}_1(z) \cdots \mathcal{D}\mathbf{r}_N(z) e^{-H_C}. \quad (3.20)$$

The functional integral $\int \mathcal{D}\mathbf{r}(z)$ was already briefly mentioned in Sec. 3.2. For a quantity $\mathcal{A}(\{\mathbf{r}(z)\})$ that depends on the chain conformation, it is represented by a discretization as in Fig. 3.5:

$$\int \mathcal{D}\mathbf{r}(z) \mathcal{A}(\{\mathbf{r}(z)\}) = \frac{\int d\mathbf{r}_1 \cdots d\mathbf{r}_{Z_L} \mathcal{A}(\{\mathbf{r}_Z\}_{Z=1}^{Z_L})}{\int d\mathbf{r}_1 \cdots d\mathbf{r}_{Z_L} \exp\left(-\frac{\sigma}{2} \sum_{Z=1}^{Z_L-1} \frac{(\mathbf{r}_{Z+1} - \mathbf{r}_Z)^2}{\Delta z^2}\right)}. \quad (3.21)$$

The normalization constant in the denominator prevents divergences for $Z_L \rightarrow \infty$ and is such that $\int \mathcal{D}\mathbf{r}(z) e^{-H_{\text{align},1}\{\mathbf{r}(z)\}} = 1$.

3.3.2 Disorder Average

As in the RLP model, we use the Deam-Edwards distribution as cross-link distribution (see Sec. 2.3). In addition to taking into account all possible cross-links by

summing over all possible pairs of chains, we also have to integrate over all possible cross-link heights z_e :

$$[\mathcal{A}_C] = \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{1}{L^M} \int_0^L dz_1 \cdots dz_M \mathcal{A}_C \cdot P_{\text{DE}}(\mathcal{C}) \quad (3.22a)$$

$$\text{with } P_{\text{DE}}(\mathcal{C}) = \frac{1}{\mathcal{Z}_1} \cdot \frac{(\tilde{\mu}^2)^M}{M!} Z_{\mathcal{C}} \quad (3.22b)$$

$$\text{where } \tilde{\mu}^2 = \frac{\mu^2}{2N\phi} \quad \text{and} \quad \phi = \frac{(2\pi a^2)^{D/2}}{A}. \quad (3.22c)$$

And the normalization constant \mathcal{Z}_1 is such that $[1] = 1$.

The average cross-link density is given by $[M]/N = \mu^2/2$, as in the RLP model; the calculation of $[M]$ is deferred to Appendix C.7, as it is very similar to the RLP model (Secs. 2.9.1 and B.6.1).

3.4 Replica Calculation of the Free Energy

3.4.1 Disorder-Averaged Free Energy: First Steps

We are interested in the *disorder-averaged free energy* and, again, use the replica trick to handle the average over the logarithm:

$$[F_C] = -[\ln Z_C] = -\left[\lim_{n \rightarrow 0} \frac{Z_C^n - 1}{n} \right] = -\lim_{n \rightarrow 0} \frac{[Z_C^n] - 1}{n}. \quad (3.23)$$

The disorder-averaged Z_C^n is calculated by applying Eq. (3.22), which leads to the introduction of $n+1$ replicas:

$$\begin{aligned} [Z_C^n] &= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2)^M}{M!} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{1}{L^M} \int_0^L dz_1 \cdots dz_M Z_C \cdot Z_C^n \quad (3.24) \\ &= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2/L)^M}{M!} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \int_0^L dz_1 \cdots dz_M \\ &\quad \left(\int \mathcal{D}\mathbf{r}_1(z) \cdots \mathcal{D}\mathbf{r}_N(z) e^{-H_{\text{ev}}\{\mathbf{r}_j\} - H_{\text{align}}\{\mathbf{r}_j\}} \exp\left(-\sum_{e=1}^M \frac{(\mathbf{r}_{i_e}(z_e) - \mathbf{r}_{j_e}(z_e))^2}{2a^2}\right) \right)^{n+1} \\ &= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2/L)^M}{M!} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \int_0^L dz_1 \cdots dz_M \\ &\quad \int \mathcal{D}\hat{\mathbf{r}}_1(z) \cdots \mathcal{D}\hat{\mathbf{r}}_N(z) e^{-H_{\text{ev}}^{(n+1)}\{\hat{\mathbf{r}}_j\} - H_{\text{align}}^{(n+1)}\{\hat{\mathbf{r}}_j\}} \prod_{e=1}^M \exp\left(-\frac{(\hat{\mathbf{r}}_{i_e}(z_e) - \hat{\mathbf{r}}_{j_e}(z_e))^2}{2a^2}\right). \end{aligned}$$

Here we implemented the $(n+1)$ -times replicated hatted vectors $\hat{r}_j(z) = (\mathbf{r}_j^{(0)}(z), \dots, \mathbf{r}_j^{(n)}(z))$, which is the “launch” of the replica theory. Different from the calculation of the RLP model, these vectors are functions of z . The replicated Hamiltonians are:

$$H_{\text{ev}}^{(n+1)}\{\hat{r}_j\} = \sum_{\alpha=0}^n H_{\text{ev}}\{\mathbf{r}_j^{(\alpha)}\} \quad (3.25)$$

$$H_{\text{align}}^{(n+1)}\{\hat{r}_j\} = \sum_{\alpha=0}^n H_{\text{align}}\{\mathbf{r}_j^{(\alpha)}\} \quad (3.26)$$

The sum over M in Eq. (3.24) can be performed, as shown in Appendix C.3.1. We obtain a result similar to Eqs. (2.14) for the RLP model:

$$[Z_{\mathcal{C}}^n] = \frac{\mathcal{Z}_{n+1}}{\mathcal{Z}_1}, \quad (3.27a)$$

$$\text{with } \mathcal{Z}_{n+1} = \int \left(\prod_{j=1}^N \mathcal{D}\hat{r}_j(z) \right) \exp(-N\tilde{f}_{n+1}\{\hat{r}_j\}) \quad (3.27b)$$

$$\text{and } \tilde{f}_{n+1}\{\hat{r}_j\} = -\frac{\tilde{\mu}^2}{LN} \sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{(\hat{r}_i(z) - \hat{r}_j(z))^2}{2a^2}\right) + \frac{H_{\text{ev}}^{(n+1)}}{N} + \frac{H_{\text{align}}^{(n+1)}}{N} \quad (3.27c)$$

where \mathcal{Z}_{n+1} is the **replica partition function** and $\tilde{f}_{n+1}\{\hat{r}_j\}$ is the **replica free energy**. At this point, it is interesting to note that, while the chains $\hat{r}_j(z)$ are $n+1$ replicated vectors, z is not replicated. This is because the integration over z was originally part of the disorder average (like the sum $\sum_{i,j}$), see Eq. (3.24), while the integration over the chain positions $\int \mathcal{D}\hat{r}_j(z)$ comes from the (replicated) partition function.

As in Eq. (2.57) for the RLP model, \mathcal{Z}_{n+1} is related to the disorder-averaged free energy by:

$$-[F_{\mathcal{C}}] = [\ln Z_{\mathcal{C}}] = \lim_{n \rightarrow 0} \frac{\partial}{\partial n} \ln \mathcal{Z}_{n+1}. \quad (3.28)$$

3.4.2 Introduction of the Replicated Density Field

As for the RLP model, it is useful to define a replicated density field, which can characterize the state of the system and easily distinguish between sol and gel phase. Now, it additionally depends on the height z :

$$\text{real space: } O(\hat{x}, z) := \frac{1}{N} \sum_{j=1}^N \delta(\hat{x} - \hat{r}_j(z)), \quad (3.29a)$$

$$\text{Fourier space: } O(\hat{q}, z) := \frac{1}{N} \sum_{j=1}^N \exp(i\hat{q}\hat{r}_j(z)). \quad (3.29b)$$

For a *slice at a given system height* z , the replicated density $O(\hat{x}, z)$ gives the probability to find a particle at $\mathbf{x}^{(0)}$ in replica (0), at $\mathbf{x}^{(1)}$ in replica (1),... and at $\mathbf{x}^{(n)}$ in replica (n). Hence it can answer the critical question, whether $|\mathbf{r}_j^{(\alpha)}(z) - \mathbf{r}_j^{(\beta)}(z)|$ remains finite for different replicas α and β , i.e. whether or not the chain segment of polymer j at height z is localized; and if so, it can give information about the extent ξ of the localization.

This localization of the polymer chains occurs when the cross-link structure gives rise to a macroscopic cluster: if a chain is part of that macroscopic cluster, it will be hindered from traversing the whole system¹; hence the replicated density (3.29) is capable of detecting the emergence of a macroscopic cluster and therefore distinguish between sol and gel phase.

As for the RLP model, we distinguish between the **one replica sector** (1RS), the set of \hat{q} -vectors with exactly one of the $n+1$ replica components non-zero, and the **higher replica sector** (HRS), the set of \hat{q} -vectors with at least two non-zero components; see Sec. 2.4.3. For $\hat{q} \in 1RS$, Eq. (3.29b) is the simple Fourier density of the respective replica, and therefore describes density fluctuations at height z :

$$O(\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}^{(\alpha)}, \mathbf{0}, \dots, \mathbf{0}), z) = \frac{1}{N} \sum_{j=1}^N \exp(i\mathbf{q}^{(\alpha)} \mathbf{r}_j^{(\alpha)}(z)). \quad (3.30)$$

In Appendix C.3.2, we see how we can rewrite the replica free energy (3.27c) in terms of the density field (3.29b):

$$\begin{aligned} \tilde{f}_{n+1}\{\hat{r}_j\} = & -\phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |O(\hat{q}, z)|^2 \Delta(\hat{q}) + \frac{1}{2} \int_0^L dz \sum_{\hat{q} \in 1RS} |O(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) \\ & + f_0 + \frac{H_{\text{align}}^{(n+1)}\{\hat{r}_j\}}{N} \end{aligned} \quad (3.31a)$$

where we defined the mean chain density $n_0 = N/A$ as the number of chains per (hyper-)area and:

$$f_0 = -\phi^n \frac{\mu^2}{2} + (n+1) \frac{\lambda n_0 L}{2}, \quad (3.31b)$$

$$\Delta(\hat{q}) = \exp\left(-\frac{\hat{q}^2 a^2}{2}\right), \quad (3.31c)$$

$$\tilde{\lambda}(\hat{q}) = \lambda n_0 U(\hat{q}) - \phi^n \frac{\mu^2}{L} \Delta(\hat{q}). \quad (3.31d)$$

As we can see in Eq. (3.31a), $\tilde{\lambda}(\hat{q})$ controls the fluctuations of the unreplicated density in the 1RS, and as in the RLP model, it has to be positive to make sure the

¹If chain i_e is clamped to the macroscopic cluster at height z_e , one might ask the question, if its chain segment at a height z_0 far from the “clamping point” is still localized. However, as long as the transverse *radius of gyration* of the chain is much less than the transverse *extent of the system*, i.e. $\ell = \sqrt{L/(2\sigma)} \ll A^{1/D}$ (*), a chain localized at height z_e should be localized at any other height z_0 . If the thermodynamic limit is taken such that $L \propto A^{1/D}$ or $L < A^{1/D}$, (*) is fulfilled.

system does not collapse. In Eq. (3.31d) it is interesting to see that the strength of the excluded volume interaction is counteracted by the cross-link density *per height*, μ^2/L , not by the cross-link density μ^2 alone as in the RLP model. That means that the cross-link density μ^2 has to scale linearly with the system height L , if we want to keep the system properties the same, in particular to compensate a fixed excluded volume interaction. This makes sense intuitively, since the excluded volume interaction acts on the polymers along the *whole height of the system*, see Eq. (3.16), whereas the cross-link interaction only acts at the given *points* z_e , see Eq. (3.15).

3.4.3 Introduction of a Field Theory and Decoupling

The expression (3.31a) has the disadvantage that the density $O(\hat{q}, z)$ in the upper line appears in a quadratic form. As we did for the RLP model (in Sec. 2.4.4/Appendix B.2), we want to linearize those two terms with the help of the **Hubbard-Stratonovich transformation**. Since $O(\hat{q}, z) \propto \sum_j e^{i\hat{q}r_j(z)}$, this has the great advantage that those two terms can be written as a *single* sum over all particles; then in the partition function (3.27b) the integral $\int \mathcal{D}\hat{r}_1(z) \cdots \mathcal{D}\hat{r}_N(z)$ decouples with respect to the particles $j = 1, \dots, N$, which drastically simplifies the calculation. The drawback of the linearization is that we have to introduce the integration over a new field $\Omega(\hat{q}, z)$, whose expectation value, however, is the same as $O(\hat{q}, z)$.

As we see in Appendix C.3.3, we obtain for the **replica partition function**:

$$\mathcal{Z}_{n+1} = \exp(-Nf_0) \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\}) \quad (3.32a)$$

with a **replica free energy**, dependent on the field $\Omega(\hat{q}, z)$:

$$f_{n+1}\{\Omega\} = \phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |\Omega(\hat{q}, z)|^2 \Delta(\hat{q}) + \frac{1}{2} \int_0^L dz \sum_{\hat{q} \in \text{IRS}} |\Omega(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) - \ln \mathfrak{z}, \quad (3.32b)$$

with an **effective one-particle partition function**:

$$\mathfrak{z} = \int \mathcal{D}\hat{r}(z) \exp \left(\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} + i \int_0^L dz \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} - H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\} \right), \quad (3.32c)$$

and the constant contribution f_0 as before (see Eq. (3.31b)).

This form can be used as a starting point for answering many questions arising from this system, like the density-density correlation function, the response of the system to a tilt field, the dependence of the bulk modulus on λ , the shear modulus, *etc.* It is important to note that **no approximation has been made** so far.

In this thesis, we will focus on the gelation transition of this system and the localization length of the chains. We take the limit of infinitely strong excluded

volume interaction by taking the limit $\lambda \rightarrow \infty$ and hence $\tilde{\lambda}(\hat{q}) \rightarrow \infty$. This prevents fluctuations of the density in the one replica sector and simplifies the equation:

$$\begin{aligned} f_{n+1}\{\Omega\} &= \phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |\Omega(\hat{q}, z)|^2 \Delta(\hat{q}) - \ln \mathfrak{z} , \\ \mathfrak{z} &= \int \mathcal{D}\hat{r}(z) \exp \left(\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} - H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\} \right) \end{aligned} \quad (3.33)$$

3.4.4 Saddle Point Equation

In the RLP model, we plugged in an Ansatz for the order parameter into the replica free energy, in order to calculate quantities like the energy cost of shear deformations. Here, we are interested in the localization of the chains. Therefore, we set up the saddle point equation for the order parameter and plug in a physically motivated Ansatz similar to the one employed in Sec. 2.6. We will see that the Ansatz consistently solves the saddle point equation, yielding an equation for the distribution of localization lengths.

The saddle point equation can be retrieved from deriving $f_{n+1}\{\Omega\}$, Eq. (3.33), with respect to Ω :²

$$0 \stackrel{!}{=} \frac{\delta f_{n+1}\{\Omega\}}{\delta \Omega(\hat{q}_0, z_0)} = \phi^n \frac{\mu^2}{L} \Omega(-\hat{q}_0, z_0) \Delta(\hat{q}_0) - \left\langle \phi^n \frac{\mu^2}{L} \Delta(\hat{q}_0) e^{-i\hat{q}_0 \hat{r}(z_0)} \right\rangle_{\mathfrak{z}} , \quad (3.34)$$

which simplifies to a very plausible form:

$$\Omega(-\hat{q}_0, z_0) = \left\langle e^{-i\hat{q}_0 \hat{r}(z_0)} \right\rangle_{\mathfrak{z}} =: \frac{I(\hat{q}_0, z_0)}{I(\hat{0}, 0)} \quad (3.35)$$

Here $\langle \rangle_{\mathfrak{z}}$ represents averaging with the statistical weight of the one-particle partition function \mathfrak{z} (see Eq. (3.33)), and $I(\hat{q}_0, z_0)$ is defined accordingly:

$$\begin{aligned} I(\hat{q}_0, z_0) &= \\ & \int \mathcal{D}\hat{r}(z) e^{-i\hat{q}_0 \hat{r}(z_0)} \exp \left(\phi^n \frac{\mu^2}{L} \int_0^L dz \sum'_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} - H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\} \right) . \end{aligned} \quad (3.36)$$

²Strictly speaking, one has to take into account that Ω is complex and take *both* derivatives $\delta f_{n+1}\{\Omega\}/\delta \text{Re} \Omega = \delta f_{n+1}\{\Omega\}/\delta \text{Im} \Omega = 0$. By setting $\Omega(\hat{q}, z) = \text{Re} \Omega(\hat{q}, z) + i \text{Im} \Omega(\hat{q}, z)$ in Eq. (3.33), one can easily see that this yields the same result.

The primed sum is defined by:

$$\sum'_{\hat{q} \in \text{HRS}} (\dots) := Q + \sum_{\hat{q} \in \text{HRS}} (\dots) \quad (3.37)$$

The additional Q seems unnecessary, since it does not depend on $\hat{r}(z)$ and hence cancels out in numerator and denominator of Eq. (3.35); however it will simplify further calculations.

As next step, we want to develop an Ansatz for the order parameter $\Omega(\hat{q}, z)$ and then show that it solves the saddle-point equation (3.35).

3.5 Ansatz for the Order Parameter

Here we try to develop an Ansatz for the order parameter

$$\langle \Omega(\hat{q}, z) \rangle_{f_{n+1}} = \frac{1}{N} \left\langle \sum_{j=1}^N \exp(i\hat{q}\hat{r}_j(z)) \right\rangle_{f_{n+1}} \quad (3.38)$$

The assumptions we make are similar to the assumptions that [Castillo *et al.* \[1994\]](#) made to solve the saddle-point equations for an isotropic system, which were already mentioned in Sec. 2.6. For our system, the assumptions are:

1. A fraction Q of chains is **localized** (“loc.”). If chain j is part of that fraction, its chain segment $\mathbf{r}_j^{(\alpha)}(z)$ at height z in replica α performs Gaussian fluctuations around a fixed mean position $\mathbf{R}_j(z)$, which is *independent of the replica* (see Fig. 3.6). The width of this distribution is the localization length ξ_j , which follows a distribution $\mathcal{P}(\xi^2, z)$ that may depend on the height z .
2. To account for the disorder average, the mean positions $\mathbf{R}_j(z)$ are randomly placed with equal probability throughout the system.
3. The remaining fraction $1 - Q$ is **delocalized** (“deloc.”) and therefore has equal probability to be found anywhere in the system.

It is important to note that these assumptions will not give rise to correlations in z -direction. This is sufficient for our case, because we only need these assumptions for calculating the expectation value $\langle \Omega(\hat{q}, z) \rangle$, for which every slice is treated independently. If, however, we needed correlators like $\langle \Omega(\hat{q}, z) \Omega(-\hat{q}, z') \rangle \propto \sum_{i,j=1}^N \langle \exp(i\hat{q}(\hat{r}_i(z) - \hat{r}_j(z'))) \rangle$, we would need to include further assumptions, e.g. the joint probability distribution for mean positions $\mathbf{R}_j(z)$ and $\mathbf{R}_j(z')$ with at different heights z and z' .

To set up the Ansatz for the order parameter, we first incorporate the Gaussian fluctuations of a chain segment around a mean position: The joint probability density to find chain segment j at position $\mathbf{x}^{(0)}$ in replica 0, at $\mathbf{x}^{(1)}$ in replica 1, *etc.*

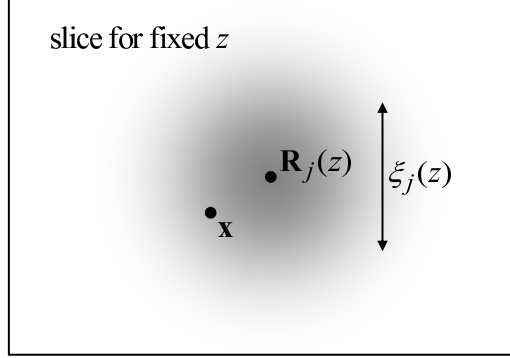


Figure 3.6: Transverse cut through the system, showing one polymer chain j . The probability distribution that the chain segment is at position \mathbf{x} (depicted as grayscale values) is a Gaussian distribution of width $\xi_j(z)$ and mean position $\mathbf{R}_j(z)$.

is

$$\langle \delta(\hat{x} - \hat{r}_j(z)) \rangle = \begin{cases} \frac{1}{(2\pi\xi_j^2(z))^{D(n+1)/2}} \exp\left(-\frac{(\hat{x} - \hat{R}_j(z))^2}{2\xi_j^2(z)}\right) & \text{if } j \text{ loc.} \\ A^{-(n+1)} & \text{if } j \text{ deloc.} \end{cases} \quad (3.39)$$

where the mean position is the same in all replicas: $\hat{R}_j = (\mathbf{R}_j, \dots, \mathbf{R}_j)$. Its Fourier transform is given by:

$$\begin{aligned} \langle \exp(i\hat{q}\hat{r}_j(z)) \rangle &= \quad (3.40) \\ &= \int d\hat{x} \exp(i\hat{q}\hat{x}) \langle \delta(\hat{x} - \hat{r}_j(z)) \rangle = \begin{cases} \exp\left(-\frac{\hat{q}^2\xi_j^2(z)}{2} + i\hat{q}\hat{R}_j(z)\right) & \text{if } j \text{ loc.} \\ \delta_{\hat{q},\hat{0}} & \text{if } j \text{ deloc.} \end{cases} \end{aligned}$$

Hence, we get for the order parameter:

$$\begin{aligned} \langle \Omega(\hat{q}, z) \rangle &= \frac{1}{N} \sum_{j=1}^N \langle \exp(i\hat{q}\hat{r}_j(z)) \rangle \\ &= \frac{1}{N} \sum_{j \text{ loc.}} \exp\left(-\frac{\hat{q}^2\xi_j^2(z)}{2} + i\hat{q}\hat{R}_j(z)\right) + \frac{1}{\mathcal{N}} \cdot \mathcal{N}(1-Q)\delta_{\hat{q},\hat{0}} \quad (3.41) \end{aligned}$$

Now, to account for the disorder average, we assume an even distribution of the mean positions $\mathbf{R}_j(z)$. We also embed the distribution for the localization lengths, which may depend on the height z :

$$\mathcal{P}(\xi^2, z) = \frac{1}{QN} \sum_{j \text{ loc.}} \delta(\xi^2 - \xi_j^2(z)). \quad (3.42)$$

So our Ansatz for the order parameter, denoted as $\bar{\Omega}$, becomes:

$$\begin{aligned} \bar{\Omega}(\hat{q}, z) &= (1 - Q)\delta_{\hat{q}, \hat{0}} \\ &= \frac{1}{N} \sum_{j \text{ loc.}} \exp\left(-\frac{\hat{q}^2 \xi_j^2(z)}{2}\right) \int \frac{d\mathbf{R}_j(z)}{A} \exp\left(i \sum_{\alpha=0}^n \mathbf{q}^{(\alpha)} \mathbf{R}_j(z)\right) \\ &= Q\delta_{\mathbf{q}_{\parallel}, \mathbf{0}} \int_0^\infty d\xi^2 \mathcal{P}(\xi^2, z) \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right) \end{aligned} \quad (3.43)$$

and hence:

$$\boxed{\bar{\Omega}(\hat{q}, z) = (1 - Q)\delta_{\hat{q}, \hat{0}} + Q\delta_{\mathbf{q}_{\parallel}, \mathbf{0}} \int_{\xi^2, z} \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right)} \quad (3.44)$$

where we defined:

$$\int_{\xi^2, z} := \int_0^\infty d\xi^2 \mathcal{P}(\xi^2, z) \quad (3.45)$$

3.6 The Saddle-Point Equation with Ansatz for Ω

Now we plug in the Ansatz (3.44) into the saddle point equation (3.35) and get:

$$\boxed{(1 - Q)\delta_{\hat{q}_0, \hat{0}} + Q\delta_{\mathbf{q}_{0\parallel}, \mathbf{0}} \int_{\xi^2, z_0} \exp\left(-\frac{\hat{q}_0^2 \xi^2}{2}\right) = \frac{I(\hat{q}_0, z_0)}{I(\hat{0}, 0)}} \quad (3.46)$$

When plugging the Ansatz into $I(\hat{q}_0, z_0)$, Eq. (3.36), we note that the primed sum can be turned into a sum over all \hat{q} :

$$\sum'_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \bar{\Omega}(\hat{q}, z) e^{-i\hat{q}r(z)} = Q \sum_{\hat{q}} \Delta(\hat{q}) e^{-i\hat{q}r(z)} \delta_{\mathbf{q}_{\parallel}, \mathbf{0}} \int_{\xi^2, z} \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right), \quad (3.47)$$

because

1. for $\hat{q} \in \text{IRS}$, the Ansatz $\bar{\Omega}(\hat{q}, z) = 0$ due to the constraint $\delta_{\mathbf{q}_{\parallel}, \mathbf{0}}$, and
2. the additional Q of the primed sum (see Def. (3.37)) on the left hand side and the term for $\hat{q} = \hat{0}$ on the right hand side cancel out.

Furthermore, looking at the definition of $\Delta(\hat{q})$, Eq. (3.31c), we can simplify the calculation by absorbing the cross-linking length a into the localization length ξ :

$$\Delta(\hat{q}) \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right) = \exp\left(-\frac{\hat{q}^2 a^2}{2}\right) \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right) = \exp\left(-\frac{\hat{q}^2 \tilde{\xi}^2}{2}\right) \quad (3.48)$$

$$\text{with } \tilde{\xi}^2 \equiv \xi^2 + a^2 \quad (3.49)$$

Therefore:

$$\begin{aligned}
 I(\hat{q}_0, z_0) &= \int \mathcal{D}\hat{r}(z) e^{-i\hat{q}_0\hat{r}(z_0)} \\
 &\times \exp\left(\phi^n \frac{\mu^2 Q}{L} \int_0^L dz \sum_{\hat{q}} e^{-i\hat{q}\hat{r}(z)} \delta_{\mathbf{q}_{\parallel}, \mathbf{0}} \int_{\xi^2, z} \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right) - H_{\text{align}, 1}^{(n+1)}\{\hat{r}(z)\}\right)
 \end{aligned}
 \tag{3.50}$$

3.6.1 Expansion to second order in Q

The functional integral $I(\hat{q}_0, z_0)$, Eq. (3.50), can be calculated to second order in Q , as seen in Appendix C.4. Plugging the result for $I(\hat{q}_0, z_0)$ back into the saddle point equation (3.46) yields:

$$\begin{aligned}
 &Q (1 + \mu^2 Q) \int_{\xi^2, s_0} \exp\left(-\frac{q_{0\perp}^2 \xi^2}{2}\right) \\
 &= \mu^2 Q \int_0^1 ds_1 \int_{\xi_1^2, s_1} \exp\left(-\frac{q_{0\perp}^2}{2} (2\ell^2 |s_0 - s_1| + a^2 + \xi_1^2)\right) \\
 &\quad + \frac{(\mu^2 Q)^2}{2} \int_0^1 ds_1 ds_2 \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} \\
 &\quad \exp\left(\frac{q_{0\perp}^2}{2} \left\{ \frac{(\ell^2 g(\mathbf{s}) - a^2 - \xi_2^2)^2}{\xi_1^2 + \xi_2^2 + 2a^2 + 2\ell^2 |s_1 - s_2|} - 2\ell^2 |s_0 - s_2| - a^2 - \xi_2^2 \right\}\right) \\
 &\quad + \mathcal{O}(Q^3).
 \end{aligned}
 \tag{3.51}$$

Here, we introduced a few simplifying definitions: The dimensionless system height is defined as

$$s_\gamma := \frac{z_\gamma}{L} \in [0, 1] \quad \text{for } \gamma = 0, 1, 2. \tag{3.52}$$

Accordingly the integral

$$\int_{\xi^2, s} := \int_0^\infty d\xi^2 \mathcal{P}(\xi^2, s), \tag{3.53}$$

where $\mathcal{P}(\xi^2, s)$ depends on the dimensionless height s , but apart from that it is the same as $\mathcal{P}(\xi^2, z)$. Moreover:

$$g(\mathbf{s}) \equiv g(s_0, s_1, s_2) := |s_0 - s_1| - |s_0 - s_2| - |s_1 - s_2| \tag{3.54}$$

And in addition, the dependence on L and σ can be absorbed into

$$\ell^2 := \frac{L}{2\sigma}, \tag{3.55}$$

where ℓ is the transverse radius of gyration of an uncrosslinked polymer chain (see Sec. 3.2.1).

Eq. (3.51) is correct for arbitrary $q_{\perp 0}$ and s_0 . However the case $q_{\perp 0} = \hat{0}$ was excluded, which trivially yields $1 = 1$, as can already be seen in the basic saddle point equation (3.35).

3.6.2 Sol-Gel Transition

Eq. (3.51) in the limit $q_{0\perp}^2 \rightarrow 0$ yields the well known relation for the gel-fraction Q . (See Sec. 2.9.2, and [Castillo *et al.*, 1994; Goldbart *et al.*, 1996; Broderix *et al.*, 2002].) Here we restrict ourselves to the neighborhood of the sol-gel transition:

$$Q(1 + \mu^2 Q) = \mu^2 Q + \frac{\mu^4 Q^2}{2} + \mathcal{O}(Q^3) \quad (3.56)$$

This equation has a trivial solution $Q = 0$, the sol phase, known to be stable for $\mu^2 < 1$. For $Q > 0$, we can divide by Q to get:

$$Q = 2(\mu^2 - 1) + \mathcal{O}(Q^2), \quad (3.57)$$

which goes to zero when approaching the sol-gel transition, $\mu^2 \downarrow 1$. Defining $\varepsilon = \mu^2 - 1$, the distance from the transition point, and noting that $Q = \mathcal{O}(\varepsilon)$, as seen in the above equation, we can write:

$$Q = 2\varepsilon + \mathcal{O}(\varepsilon^2), \quad (3.58)$$

which is the same form as for the RLP model (see Eq. (2.69)).

3.7 The Equation for the Localization Length

In this section we transform the saddle point equation (3.51) into an integro-differential equation for the distribution of localization lengths. The basic steps are presented here, while the actual calculation, being a bit intricate and draggy, is deferred to Appendix C.5.

3.7.1 Normalization of Length Scales

As we can see in Eq. (3.51), there are three length scales in our system: ξ , ℓ and a . We can simplify the equations by **measuring all lengths in units of ℓ** , *i.e.* dividing all lengths by ℓ and multiplying all \hat{q} -vectors with ℓ . Hence, we define:

$$\xi_\ell^2 := \xi^2 / \ell^2 \quad (3.59a)$$

$$a_\ell^2 := a^2 / \ell^2 \quad (3.59b)$$

$$q_{0\ell\perp}^2 := \ell^2 q_{0\perp}^2 \quad (3.59c)$$

With this scaling there is no explicit dependence on ℓ in the saddle point equation anymore. See *e.g.* Eq. (C.62).

As studies of different randomly cross-linked systems have shown before, the localization length diverges as we approach the sol-gel transition from the gel side [Broderix *et al.*, 2002; Goldbart *et al.*, 1996; Peng *et al.*, 1998; Castillo *et al.*, 1994]. Hence, as a next step, we *normalize the localization length with ε* , the distance from

the critical point, which was introduced in Sec. 3.6.2. Thereby, we introduce the new variable, the **normalized inverse and squared localization length**

$$\theta := \frac{f}{\varepsilon \xi_\ell^2} = \frac{f \ell^2}{(\mu^2 - 1) \xi^2}, \quad (3.60a)$$

$$\text{with } f = \frac{2}{3} + \frac{a^2}{\ell^2} \quad (3.60b)$$

and define the corresponding distribution $\pi(\theta, s)$ for θ such that:

$$d\xi^2 \mathcal{P}(\xi^2, s) = d\theta \pi(\theta, s) \quad (3.60c)$$

As we will see, θ remains of order $\mathcal{O}(1)$, when we approach the transition point.³ Similar approaches have already been used successfully by the four above-mentioned studies to account for the diverging localization length. In our system, however, we have two relevant length scales besides the localization length, which is accounted for by the (dimensionless) factor f . The particular choice (3.60b) for f seems arbitrary, but will yield a parameter free equation for the mean distribution of θ .

3.7.2 Result for the Distribution of Localization Lengths

With the definitions and concepts of the previous section, we transform the saddle point equation (3.51) into an integro-differential equation for the distribution $\pi(\theta, s)$. The calculation, done in Appendix C.5, yields

$$\begin{aligned} & (1 + 2\varepsilon)\pi(\theta, s) \\ &= (1 + \varepsilon) \int_0^1 ds_1 \pi(\theta, s_1) + \frac{\varepsilon}{f} \int_0^1 ds_1 \partial_\theta \left(\theta^2 \pi(\theta, s_1) \right) \left\{ 2|s - s_1| + a_\ell^2 \right\} \\ & \quad + \varepsilon \int_0^1 ds_1 ds_2 \int_0^\theta d\theta_1 \pi(\theta_1, s_1) \pi(\theta - \theta_1, s_2) + \mathcal{O}(\varepsilon^2). \end{aligned} \quad (3.61)$$

This equation for $\pi(\theta, s)$ is still difficult to handle, since it involves two variables, the normalized inverse localization length θ and the normalized height s . To get a better understanding of the equation, it is useful to define

$$\bar{\pi}(\theta) = \int_0^1 ds \pi(\theta, s), \quad (3.62)$$

³Alternatively we could define $\hat{\xi}^2 \sim \varepsilon \xi^2$, which is also of the order $\mathcal{O}(1)$. However, defining the reciprocal, Eq. (3.60a), turns out to yield a simpler differential equation in the end.

the *mean value* of the distribution with respect to s . Using that definition and by sorting Eq. (3.61) in orders of ε , we get:

$$\begin{aligned} \pi(\theta, s) - \bar{\pi}(\theta) &= \varepsilon \left(-2\pi(\theta, s) + \bar{\pi}(\theta) + \frac{1}{f} \int_0^1 ds_1 \partial_\theta \left(\theta^2 \pi(\theta, s_1) \right) \left\{ 2|s - s_1| + a_\ell^2 \right\} \right. \\ &\quad \left. + \int_0^\theta d\theta_1 \bar{\pi}(\theta_1) \bar{\pi}(\theta - \theta_1) \right) + \mathcal{O}(\varepsilon^2). \end{aligned} \quad (3.63)$$

This equation states that the deviation of the distribution from its mean value is of the order ε . Hence it is suggestive to define $\delta\pi(\theta, s)$ with:

$$\pi(\theta, s) \equiv \bar{\pi}(\theta) + \varepsilon \delta\pi(\theta, s), \quad (3.64a)$$

$$\text{with } \int_0^1 ds \delta\pi(\theta, s) = 0, \quad (3.64b)$$

and we are able to calculate both parts, $\bar{\pi}(\theta)$ and $\delta\pi(\theta, s)$, separately to lowest order in ε :

Therefore we first plug this definition into the right hand side of Eq. (3.63). Since these expressions are already of order ε , we can just replace $\pi(\theta, s)$ by $\bar{\pi}(\theta)$, because the corrections are of the order $\mathcal{O}(\varepsilon^2)$:

$$\begin{aligned} \varepsilon \delta\pi(\theta, s) &= \varepsilon \left(-\bar{\pi}(\theta) + \frac{1}{f} \int_0^1 ds_1 \partial_\theta \left(\theta^2 \bar{\pi}(\theta) \right) \left\{ 2|s - s_1| + a_\ell^2 \right\} + \int_0^\theta d\theta_1 \bar{\pi}(\theta_1) \bar{\pi}(\theta - \theta_1) \right) \\ &\quad + \mathcal{O}(\varepsilon^2). \end{aligned} \quad (3.65)$$

Now it is possible to perform the integration over s_1 in the second term on the right hand side

$$\int_0^1 ds_1 |s - s_1| = \frac{s^2 + (1-s)^2}{2}. \quad (3.66)$$

Furthermore we plug in $f = \frac{2}{3} + a_\ell^2$, see Eq. (3.60b), and divide by ε :

$$\begin{aligned} \delta\pi(\theta, s) &= \\ &= -\bar{\pi}(\theta) + \frac{s^2 + (1-s)^2 + a_\ell^2}{2/3 + a_\ell^2} \partial_\theta \left(\theta^2 \bar{\pi}(\theta) \right) + \int_0^\theta d\theta_1 \bar{\pi}(\theta_1) \bar{\pi}(\theta - \theta_1) + \mathcal{O}(\varepsilon) \end{aligned}$$

(3.67)

This equation can be used to obtain the form of the deviation $\delta\pi(\theta, s)$ from the mean distribution, once we know the mean distribution $\bar{\pi}(\theta)$. In order to get the equation for $\bar{\pi}(\theta)$, we integrate both sides over s , use

$$\int_0^1 ds \left(s^2 + (1-s)^2 \right) = \frac{2}{3} \quad (3.68)$$

for the second term and get:

$$0 = -\bar{\pi}(\theta) + \partial_{\theta}(\theta^2 \bar{\pi}(\theta)) + \int_0^{\theta} d\theta_1 \bar{\pi}(\theta_1) \bar{\pi}(\theta - \theta_1) + \mathcal{O}(\varepsilon), \quad (3.69)$$

and hence:

$$(1 - 2\theta)\bar{\pi}(\theta) = \theta^2 \bar{\pi}'(\theta) + \int_0^{\theta} d\theta_1 \bar{\pi}(\theta_1) \bar{\pi}(\theta - \theta_1) + \mathcal{O}(\varepsilon). \quad (3.70)$$

Comparing Eqs. (3.67) and (3.69), we can simplify the equation for the deviation from the mean value:

$$\delta\pi(\theta, s) = \left(\frac{s^2 + (1-s)^2 + a_{\ell}^2}{2/3 + a_{\ell}^2} - 1 \right) \partial_{\theta}(\theta^2 \bar{\pi}(\theta)) + \mathcal{O}(\varepsilon), \quad (3.71)$$

and hence:

$$\begin{aligned} \delta\pi(\theta, s) &= w(s) \partial_{\theta}(\theta^2 \bar{\pi}(\theta)) + \mathcal{O}(\varepsilon), \\ \text{with } w(s) &= \frac{s^2 + (1-s)^2 - 2/3}{a^2/\ell^2 + 2/3}. \end{aligned} \quad (3.72)$$

Here we have to keep in mind that the actual localization lengths are related to θ by

$$\xi_{\ell}^2 = \frac{f}{\varepsilon\theta} = \frac{\frac{2}{3} + a_{\ell}^2}{(\mu^2 - 1)\theta} \quad (3.73)$$

as defined in Eqs. (3.60), and with Eq. (3.59a):

$$\xi^2 = \frac{\frac{2}{3}\ell^2 + a^2}{(\mu^2 - 1)\theta} \quad (3.74)$$

The result for $\bar{\pi}(\theta)$ from Eq. (3.70) can be computed numerically: This is done by discretizing $\bar{\pi}(\theta)$ into points $\bar{\pi}_j = \bar{\pi}(j \cdot \Delta\theta)$ with $j = 0, \dots, j_{\max}$ and solving the resulting set of (polynomial) equations. The constraint $\sum_{j=0}^{j_{\max}} \Delta\theta \cdot \bar{\pi}_j = 1$ has to be taken into account. The result for $\bar{\pi}(\theta)$ obtained by this procedure is plotted in Fig. 3.7.

Indeed Eq. (3.70) for the (normalized and inverse) localization length is well known⁴: It is also found and investigated for isotropic polymer systems [Broderix *et al.*, 2002; Goldbart *et al.*, 1996].

Here, however, the system is not isotropic, but has the preferred z -direction. Since the ends of the chains are loose, the behavior at the top and bottom boundaries

⁴Different ways of normalization of the localization lengths may yield different prefactors for the terms of Eq. (3.70), however, they can always be reconciled with Eq. (3.70)

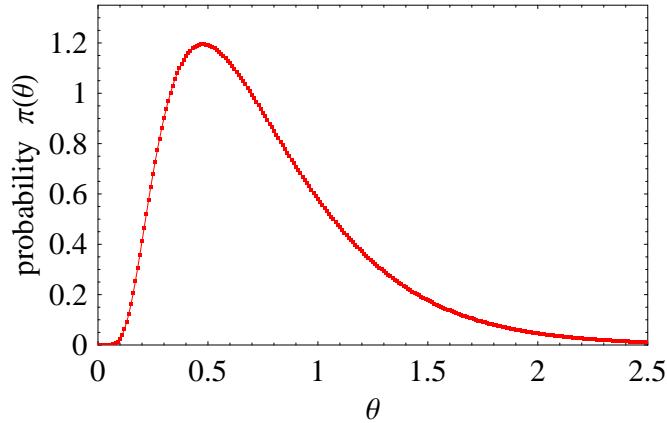


Figure 3.7: Mean probability distribution $\bar{\pi}(\theta)$ of the normalized inverse localization length θ . The curve was obtained by solving the integro-differential equation (3.70) numerically.

is different from the bulk. This difference can be understood in the following way: A chain segment far from the boundaries has a certain probability to have a cross-link above or below. However, if this segment is close to, let's say, the top boundary, the probability for a cross-link to be above is lower. Consequently, since close to the boundary the chains have fewer nearby cross-links, one would expect an increased localization length. This behavior is indeed seen in Fig. 3.8, where the deviation $\delta\pi(\theta, s)$, obtained from Eq. (3.72), is plotted vs. θ and the system height s . One can see that, indeed, close to the boundaries $s \approx 0$ and $s \approx 1$, smaller θ and hence larger localization lengths ξ are favored. On the other hand, inside the bulk $s \approx 1/2$, the opposite is true.

The strength of this inhomogeneity is controlled by the ratio of the two *internal* length scales $a_\ell^2 = a^2/\ell^2$. When taking the limit $\ell^2 = L/(2\sigma) \rightarrow 0$, i.e. considering thin slices ($L \rightarrow 0$) or well aligned parallel chains ($\sigma \rightarrow \infty$), the presented $(D+1)$ -dimensional system becomes effectively D -dimensional and isotropic. This can also be seen in Eq. (3.72): For $\ell^2 \rightarrow 0$, the prefactor $w(s) \rightarrow 0$ and hence the deviations of the localization length from its mean value vanish. In this case, the relevant length scale for the localization length is a , see Eq. (3.74):

$$\xi^2 = \frac{a^2}{(\mu^2 - 1)\theta} \quad \Rightarrow \quad \xi \propto \frac{a}{\sqrt{\mu^2 - 1}}. \quad (3.75)$$

The other and more relevant case is the limit of large system heights $L \rightarrow \infty$. In this case the relevant length scale for the localization length is the radius of gyration $\ell^2 = L/(2\sigma)$ of the chains (perpendicular to the z -axis); the cross-linking length a becomes unimportant:

$$\xi^2 = \frac{2}{3} \frac{\ell^2}{(\mu^2 - 1)\theta} \quad \Rightarrow \quad \xi \propto \frac{\sqrt{L}}{\sqrt{\mu^2 - 1}}. \quad (3.76)$$

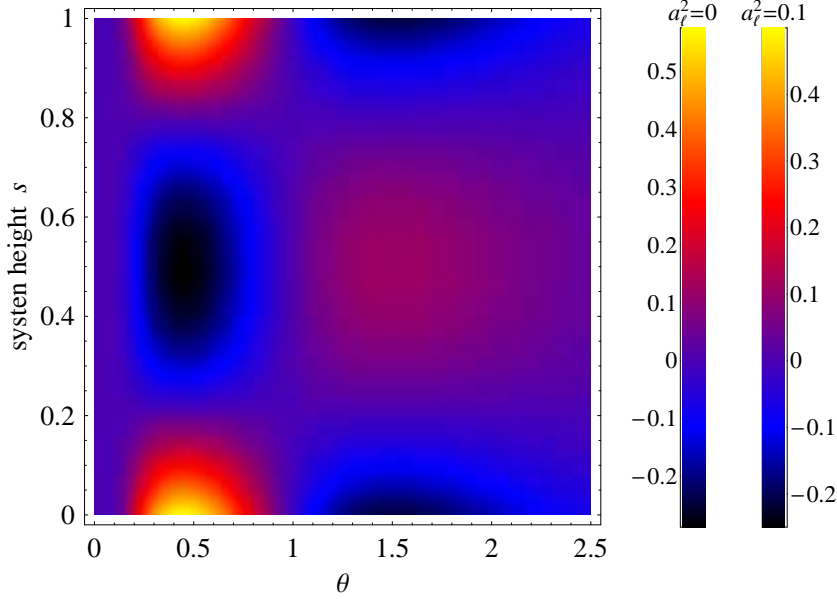


Figure 3.8: Deviation $\delta\pi(\theta, s)$ of the probability distribution of θ from its mean value, as defined in Eq. (3.64a). The plot was obtained by applying the data from Fig. 3.7 to Eq. (3.72). For different ratios $a_l^2 = a^2/\ell^2$, $\delta\pi(\theta, s)$ can be rescaled simply resulting in different color bars.

One can also see that larger system heights L yield larger localization lengths, since, due to a larger radius of gyration $\ell \propto \sqrt{L}$, the chains can span a larger region in the sample. Also the deviation $\delta\pi(\theta, s)$ is most dominant in this large- L regime (see Eq. (3.72) and compare color bars of Fig. 3.8).

As a last thing, it is instructive to see how the *typical* localization length is affected by the deviation $\delta\pi(\theta, s)$. For that we will need the first two moments of the mean distribution $\bar{\pi}(\theta)$:

$$\bar{\theta} = \int_0^\infty d\theta \theta \bar{\pi}(\theta) \quad \text{and} \quad \bar{\theta}^2 = \int_0^\infty d\theta \theta^2 \bar{\pi}(\theta). \quad (3.77)$$

With that we can easily calculate the average inverse localization length:

$$\begin{aligned} \theta_{\text{typ}}(s) &:= \int_0^\infty d\theta \theta \pi(\theta, s) = \int_0^\infty d\theta \theta \bar{\pi}(\theta) + \varepsilon \int_0^\infty d\theta \theta \delta\pi(\theta, s) = \\ &\stackrel{\text{Eq. (3.72)}}{=} \bar{\theta} + \varepsilon w(s) \int_0^\infty d\theta \theta \partial_\theta (\theta^2 \bar{\pi}(\theta)) + \mathcal{O}(\varepsilon^2) = \\ &= \bar{\theta} + \varepsilon w(s) \left(\theta \cdot \theta^2 \bar{\pi}(\theta) \Big|_0^\infty - \int_0^\infty d\theta \theta^2 \bar{\pi}(\theta) \right) + \mathcal{O}(\varepsilon^2). \end{aligned} \quad (3.78)$$

As stated by [Castillo *et al.* \[1994\]](#), the mean distribution $\bar{\pi}(\theta)$, coming from Eq. (3.70), should fall off exponentially for $\theta \rightarrow \infty$. Hence we can omit the boundary term $\theta^3 \bar{\pi}(\theta) \Big|_0^\infty$ and get:

$$\theta_{\text{typ}}(s) = \bar{\theta} - \varepsilon \bar{\theta}^2 w(s) + \mathcal{O}(\varepsilon^2). \quad (3.79)$$

And hence, the typical localization length becomes:

$$\xi_{\text{typ}}^2(s) := \frac{f\ell^2}{\varepsilon\theta_{\text{typ}}(s)} = \frac{\frac{2}{3}\ell^2 + a^2}{\varepsilon\bar{\theta}} \left(1 + \varepsilon w(s) \frac{\bar{\theta}^2}{\theta} \right) + \mathcal{O}(\varepsilon). \quad (3.80)$$

On the right hand side of this equation, the prefactor $(\frac{2}{3}\ell^2 + a^2)/\varepsilon$ sets the length scale for the localization length, which diverges in the limit $\varepsilon \rightarrow 0$. ($\bar{\theta}$ is just a dimensionless number of order 1.) In the parentheses, the dominant term “1” does not incorporate spatial dependencies, yielding similar behavior as for isotropic systems. Relative to this term, the s -dependent second term provides corrections of order ε . Hence these anisotropic corrections are of the same order as $\frac{2}{3}\ell^2 + a^2$ and do not diverge or vanish in the limit $\varepsilon \rightarrow 0$.

Furthermore we have to note that the definition of the typical localization length $\xi_{\text{typ}}^2(s)$ is, for simplification, just the inverse of $\theta_{\text{typ}}(s)$, and thus it is not exactly the mean of the distribution $\mathcal{P}(\xi^2, s)$.

3.8 Conclusions

In this chapter we investigated a system of cross-linked directed polymer chains in a $(D+1)$ -dimensional volume $L \cdot A$, where L is the height in z -direction, along which the chains are preferably aligned, and A is the base area.

The chains are subject to an alignment interaction, which penalizes tilting of the chains with respect to the z -axis. The strength σ of this interaction may be interpreted as the force f with which a worm-like chain is stretched along the z -axis, or as the coupling strength of the polymers to a spatially uniform nematic matrix. This interaction constitutes the typical extension of an uncrosslinked chain perpendicular to the z -axis, $\ell = \sqrt{L/(2\sigma)}$.

The cross-linking interaction permanently connects random pairs of chains at a fixed height, with a cross-link density determined by the parameter μ^2 . This interaction introduces a second length scale, the typical cross-link length a .

The system exhibits a continuous phase transition – the sol-gel transition – as the cross-link concentration is increased beyond the critical value $\mu_c^2 = 1$. This transition is characterized by the gel-fraction Q , which vanishes like $Q \propto (\mu^2 - \mu_c^2)$ as the transition point is approached from the gel phase $\mu^2 \downarrow \mu_c^2$. Thereby the typical localization length diverges as $\xi_{\text{typ}} \propto (\mu^2 - \mu_c^2)^{-1/2}$. The relevant length scale is the transverse extension $\ell = \sqrt{L/(2\sigma)}$ of the chains, given that $\ell \gg a$.

The distribution of localization lengths varies throughout the height z of the system: Close to the boundaries $z \approx 0$ and $z \approx L$, larger localization lengths are favored. This is due to the fact the ends of the chains are loose and hence, the probability that a given chain segment has nearby cross-links is lower if this segment is close to a boundary.

As an outlook, it would be interesting to calculate *mechanical* properties of the system like it was done for the randomly cross-linked particle model. Since

the system is not isotropic, the *shear modulus*, describing the energy cost of a density conserving deformation of the type $\nabla_{\mathbf{r}}\mathbf{u}(\mathbf{r}, z)$, is expected to differ from the *tilt modulus*, accounting for deformations like $\partial_z\mathbf{u}(\mathbf{r}, z)$. The latter is expected to remain non-zero in the liquid phase (even in the totally uncrosslinked phase), because of the energy cost of tilting due to the alignment interaction. Also the bulk modulus, describing the energy cost of density fluctuations, would be interesting to calculate; this could be done by incorporating a pressure field to the Hamiltonian H_{align} (as proposed in Eq. (C.14)) and investigate the response of the system to pressure fluctuations.

Another matter would be to evaluate the saddle-point equation (3.46) to higher or even infinite order, as it was possible for the RLP model in Chapter 2. It would be interesting to see how stronger cross-linking affects the inhomogeneity of the localization lengths close to the boundaries, and if higher orders of the mean distribution $\bar{\pi}(\theta)$ are in accordance with corresponding *isotropic* models.

An extension to the model would be to incorporate cross-link at different heights z_e and z'_e . This would be relevant, since recent developments suggest that the tilt modulus is not influenced by cross-links which are fixed at the same height on both chains, as in the presented model. [Ulrich *et al.*, 2010]

Another modification is the introduction of so called “slip-links”: Thereby the quenched cross-link configuration *only* specifies which pairs of chains are connected. The height of the cross-link z_e , however, is not quenched but may move along the chains, as they fluctuate around. These slip-links can be seen as a means to account for fixed entanglements of the chains. In this respect, an entanglement can move along a pair of chains, but these chains cannot be disentangled.

Structure of Spider Silk Modeled as a Random Network

Contents

| | | |
|------------|--|-----------|
| 4.1 | Introduction | 57 |
| 4.2 | Model | 60 |
| 4.2.1 | Unit Cell | 60 |
| 4.2.2 | Crystallite | 60 |
| 4.2.3 | Ensemble of Crystallites | 63 |
| 4.2.4 | Continuous Background | 64 |
| 4.3 | Scattering Function | 65 |
| 4.3.1 | Incoherent Part $G_1(\mathbf{q})$ | 66 |
| 4.3.2 | Coherent Part $G_2(\mathbf{q})$ | 67 |
| 4.3.3 | Scattering Amplitude of a Single Crystallite | 68 |
| 4.4 | Atomic Configuration of the Unit Cell | 69 |
| 4.4.1 | Unshifted Unit Cells | 69 |
| 4.4.2 | Possible Shifts inside the Unit Cell | 72 |
| 4.4.3 | Variations between Crystallites | 72 |
| 4.5 | Results | 73 |
| 4.5.1 | Experimental Scattering Function | 73 |
| 4.5.2 | Scattering Function from the Model | 74 |
| 4.6 | Conclusions | 79 |

4.1 Introduction

Spider silk has long been widely known, but has only recently received great appreciation from the scientific community for its outstanding material properties [Grubb & Jelinski, 1997; Fossey & Kaplan, 1999]. In general, a spider is capable of producing a variety of different types of silk fibers; scientific interest here has focused on the so-called **dragline fibers**, fibers known for their high strength. Orb web spiders produce them essentially from only two proteins and use them to build their net's frame and radii, and also to support their own body weight after an intentional fall during escape from attack. Evolution has optimized dragline fibers for **tensile**

strength (i.e. the stress at which the material breaks), **extensibility** and **tenacity** (i.e. the energy per volume dissipated before the material breaks, a.k.a. *toughness*). Dragline silk can support large strains and has a tensile strength comparable to steel or Kevlar. Regarding tenacity, values of 160 MJ/m^3 have been reported [Kaplan *et al.*, 1994; Gosline *et al.*, 1999], e.g. for different *Nephila* species, on which most studies have been carried out.

An understanding of the structural origins of these mechanical properties is of fundamental interest, and may at the same time serve the development of biomimetic material design [Huemmerich *et al.*, 2004; Scheibel, 2004] using recombinant and synthetic approaches [Huemmerich *et al.*, 2006; Foo *et al.*, 2006; Rammensee *et al.*, 2006]. As for other biomaterials, the correlation between structure and the mechanical properties can only be clarified by advanced structural characterization accompanied by numerical modeling. To this end, not only the mechanical properties [Vollrath & Porter, 2006; Zbilut *et al.*, 2006] resulting from the structure, but also the structure itself has to be modeled to interpret the experimental data. Such efforts have in the past led to a quantitative understanding of many biomaterials like bone, tendons and wood [Roschger *et al.*, 2001; Puxkandl *et al.*, 2002].

As deduced from X-ray scattering [Warwicker, 1960; Kaplan *et al.*, 1994; Grubb & Jelinski, 1997] and NMR experiments [van Beek *et al.*, 2002], spider silks are characterized by a seemingly simple design (see Fig. 4.1): The alanine-rich segments of the fibroin polypeptide chains fold into β -sheet nano-crystallites (similar to poly-L-alanine crystals), which are embedded in an **amorphous network of chains**, containing predominately glycine. In this polymer network, the crystalline component makes up an estimated 20% - 30% of the total volume, and may be considered as **cross-links**, interconnecting several different chains.

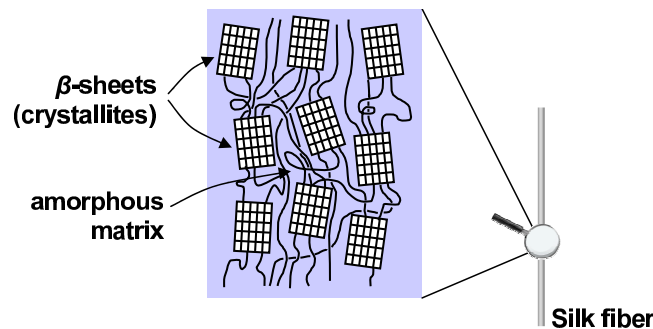


Figure 4.1: Schematic illustration of dragline silk. β -sheets, mainly composed of polyalanine, are embedded in the *amorphous matrix*, an amorphous network of chains.

The detailed investigation of the structure is complicated, at least on the **single fiber** level, by the relatively small diameters in the range of 1 - 10 μm , depending on the species. Using highly brilliant microfocused synchrotron radiation, diffraction patterns can be obtained not only on thick samples of fiber bundles, but also on a single fiber [Riekel *et al.*, 1999a,b, 2000; Riekel & Vollrath, 2001; Riekel *et al.*, 2004;

[Sapede *et al.*, 2005](#)]. Single fiber diffraction has been used under simultaneous controlled mechanical load, in order to investigate changes of the molecular structure with increasing strain up to failure [[Glišović *et al.*, 2008](#)]. Single fiber diffraction, where possible, is much better suited to correlate the structure to controlled mechanical load, since the strain distribution in bundles is intrinsically inhomogeneous, and the majority of load may be taken up by a small minority of fibers.

While progress of the experimental diffraction studies has been evident, the analysis of the data still relies on the classical classification and indexing scheme introduced by Warwicker. According to Warwicker, the β -sheet crystallites of the dragline of *Nephila* fall into the so-called system 3 of a nearly orthorhombic unit cell [[Marsh *et al.*, 1955a](#); [Warwicker, 1960](#)] with lattice constants $10.6 \text{ \AA} \times 9.44 \text{ \AA} \times 6.95 \text{ \AA}$ [[Warwicker, 1960](#)]. To fix the coordinate system, they define the x -axis to be in the direction of the amino acid side chains connecting different β -sheets, while the y -axis denotes the direction along the hydrogen bonds of the β -sheets. Finally, the z -axis corresponds to the axis along the covalent peptide bonds (main chain). The main chain is preferably aligned along the fiber axis (for an illustration, see top panel of Fig. 4.8 on page 71). Note that while we follow this common convention, other notations and choices of axes are also used in literature.

While helpful, the indexing scheme does not give information regarding the **exact structure of the unit cell**, whether the β -pleated sheets are composed of **parallel or antiparallel strands**, or how the two-dimensional **sheets are arranged** to stacks. To this end, not only peak positions but the entire rather broad intensity distribution has to be analyzed. To interpret the scattering image, it is essential to know whether correlations between different crystallites are important, or whether the measured data can be accounted for by the scattering of single crystallites averaged over fluctuating orientations. It is also not clear, if correlations between translational and rotational degrees of freedom are important. Finally, the powder averaging taking into account the fiber symmetry experimental mosaicity (orientational distribution) must be quantitatively taken into account.

In this chapter a **scattering model based on kinematic scattering theory** is developed and the numerically calculated scattering intensity is compared with the **experimental wide angle scattering distribution measured from aligned silk fibers**. The numerical calculations allow for a quantitative comparison to the experimental data and yield both **structural** and **statistical** parameters.

Note that the small size of crystallites, leading to correspondingly broad reflections and a generally rather low number of external peaks, exclude a standard crystallographic approach. The presented model is based on a quite general approach, independently allowing for correlations between center-of-mass positions (translations) and varying crystallite orientations (rotations). The structural parameters concern the crystal structure, in particular the atomic positions in the unit cell, and the crystallite size. The statistical parameters relate to the orientational distribution of the crystallite symmetry axis with respect to the fiber axis and the correlations between crystallites.

This chapter is organized as follows: In Sec. 4.2 the basic model is introduced

with parameters for the crystallite size, lattice constants and statistical parameters for the crystallites' position and orientation. Subsequently, in Sec. 4.3 we compute the scattering function for this model. Sec. 4.4 specifies the different atomic configurations which are conceivable for polyalanine. The main results and the comparison of calculated and measured intensities are presented in Sec. 4.5, before conclusion in Sec. 4.6. Most of the presented results can also be found in [Ulrich *et al.*, 2008].

4.2 Model

In this section we develop a simple model of spider silk which allows us to compute the scattering function

$$G(\mathbf{q}) = \left\langle \left| \sum_j f_j \exp(i\mathbf{q}\mathbf{r}_j) \right|^2 \right\rangle \quad (4.1)$$

as measured in X-ray scattering experiments. The atomic positions are denoted by \mathbf{r}_j and the atomic form factors by f_j . The modeling proceeds on three different levels:

- 1) On the largest length scales, spider silk is modeled as an **ensemble of crystallites** embedded in an amorphous matrix and preferentially oriented along the fiber axis.
- 2) Each crystallite is composed of **parallel or antiparallel β -sheets**.
- 3) Each unit cell contains a given number of **amino acids**, whose arrangements have been classified by Warwicker [1960]. In Fig. 4.2 (right) we see an illustration of Bombyx mori by Geis [Zubay, 1998].

In the following we shall build up a model, starting with the smallest scales and working up to the whole system. Subsequently, we will compare the calculated scattering functions with experimental data. Thereby, we are able to determine the arrangement of atoms in the unit cell which *optimizes the agreement between model and experiment*.

4.2.1 Unit Cell

One unit cell of a crystallite is described as a set of atoms at *positions* \mathbf{r}_k relative to the center of the unit cell, where $k = 1, 2, \dots, K$ runs through the atoms of the unit cell (see Fig. 4.2, left panel, for a schematic drawing). Each atom is assigned a *form factor* f_k specifying the scattering strength of the respective atom type.

4.2.2 Crystallite

A crystallite is composed of $M = M_x M_y M_z$ unit cells, replicated M_x , M_y and M_z times along the *primitive vectors* \mathbf{a}_x , \mathbf{a}_y and \mathbf{a}_z respectively. The unit cells in a crystallite are numbered by a vector index $\mathbf{m} = (m_x, m_y, m_z)$, where $m_\nu =$

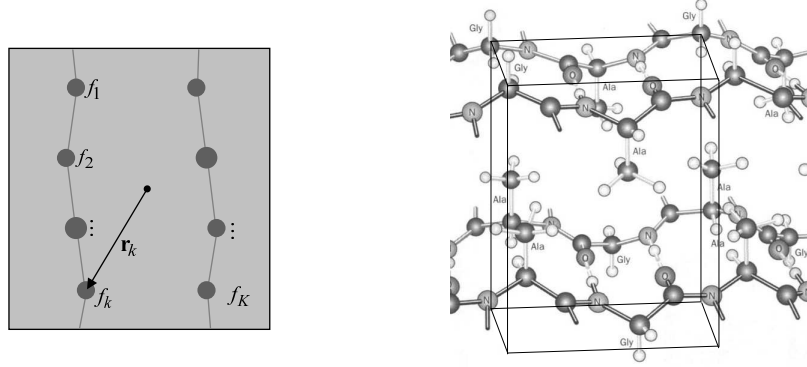


Figure 4.2: Left: Schematic view of the unit cell. Atom k is located at position \mathbf{r}_k and the atom type is specified by the form factor f_k of the atom. For simplification schematic illustrations are in 2D, if possible, even though the model refers to three space dimensions, $D = 3$. Right: A possible configuration inside the unit cell (illustration adapted from [Zubay, 1998]).

$1, 2, \dots, M_\nu$ for $\nu = x, y, z$. Hence the center of mass of unit cell \mathbf{m} has position vector

$$\tilde{\mathbf{s}}_{\mathbf{m}} = m_x \mathbf{a}_x + m_y \mathbf{a}_y + m_z \mathbf{a}_z . \quad (4.2)$$

In fact, it is more convenient to measure all distances with respect to the center of the whole crystallite

$$\mathbf{s}_{\text{cm}} = \frac{(M_x + 1)\mathbf{a}_x + (M_y + 1)\mathbf{a}_y + (M_z + 1)\mathbf{a}_z}{2} , \quad (4.3)$$

so that $\mathbf{s}_{\mathbf{m}} = \tilde{\mathbf{s}}_{\mathbf{m}} - \mathbf{s}_{\text{cm}}$ denotes the position of unit cell \mathbf{m} relative to the center of the respective crystallite (see Fig. 4.3). The position of atom k in unit cell \mathbf{m} relative to the center of the crystallite is

$$\mathbf{r}_{\mathbf{m},k} = \mathbf{s}_{\mathbf{m}} + \mathbf{r}_k . \quad (4.4)$$

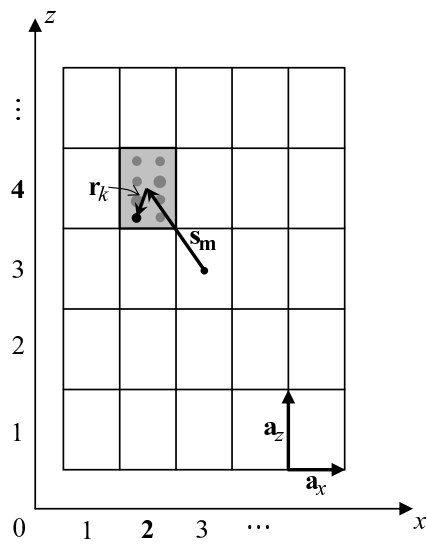


Figure 4.3: Schematic view of the crystallite. The vector \mathbf{s}_m points from the center of the crystallite to the unit cell \mathbf{m} . From there, \mathbf{r}_k points to atom k .

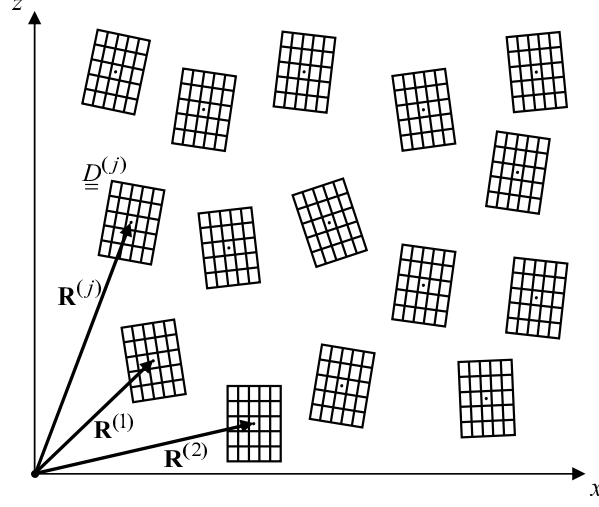


Figure 4.4: The whole system is composed of crystallites at positions $\mathbf{R}^{(j)}$. They can be rotated by rotation matrices $\underline{\underline{D}}^{(j)}$.

4.2.3 Ensemble of Crystallites

The whole system is composed of N such crystallites at positions $\mathbf{R}^{(j)}$ with $j = 1, 2, \dots, N$ (see Fig. 4.4). The crystallites are not perfectly aligned with the fiber axis, but their orientations fluctuate. The orientation of a single crystallite is specified by three Euler angles $\phi^{(j)}, \theta^{(j)}, \psi^{(j)}$ (see right panel of Fig. 4.5). Here the z -axis is the fiber axis and θ denotes the angle between the z -direction of the crystallites (direction of the covalent bonds) and the fiber axis. The atomic positions of the rotated crystallite are obtained from the configuration which is perfectly aligned with the z -axis, by applying a rotation matrix $\underline{\underline{D}}^{(j)}$ (see left panel of Fig. 4.5):

$$\mathbf{r}_{\mathbf{m},k}^{(j)} = \mathbf{R}^{(j)} + \underline{\underline{D}}^{(j)} \mathbf{r}_{\mathbf{m},k}. \quad (4.5)$$

In the experiment, the scattering intensity is obtained for a large system consisting of many crystallites. Hence, it is reasonable to assume that the scattering function is self-averaging and therefore can be averaged over the positions and orientations of the crystallites. Angular brackets $\langle \rangle$ denote the average of an observable \mathcal{A} over crystallite positions $\mathbf{R}^{(j)}$ and orientations $\underline{\underline{D}}^{(j)}$:

$$\langle \mathcal{A} \rangle = \int \prod_{j=1}^N \left(d\mathbf{R}^{(j)} \mathcal{D}\underline{\underline{D}}^{(j)} \right) \mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)}) \mathcal{A}. \quad (4.6)$$

Here, the **crystallite positions** follow the distribution function $\mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)})$, which in general includes correlations. In contrast, the

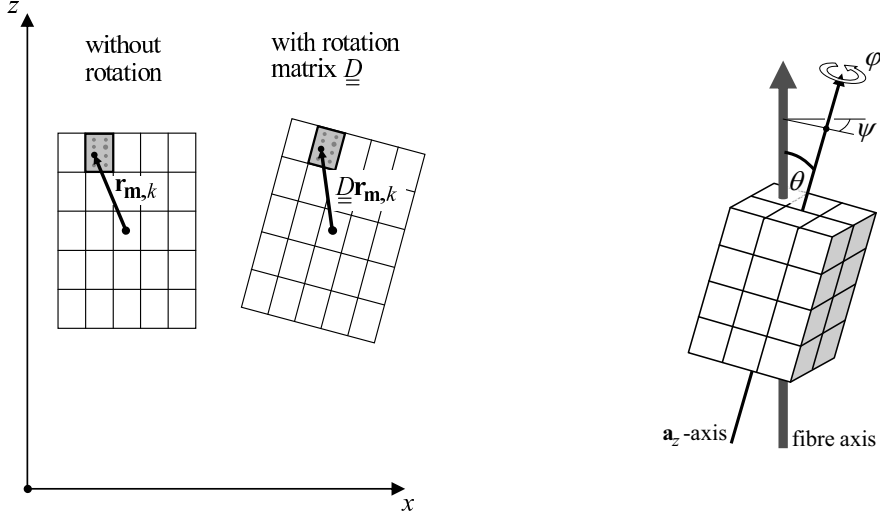


Figure 4.5: Left: The position of atom k in unit cell \mathbf{m} of the rotated crystallite is obtained by applying the rotation matrix $\underline{\underline{D}}\mathbf{r}_{\mathbf{m},k}$ to the position vector in the aligned configuration. Right: Illustration of Euler angles in 3D. The angle θ specifies the deviation of the crystallite's \mathbf{a}_z -axis from the fiber axis of the strand. ϕ is the rotation of the crystallite about its own \mathbf{a}_z -axis, and ψ is the rotation of the \mathbf{a}_z -axis about the fiber axis (after the θ -rotation).

orientation of each crystallite is assumed to be independent of the others. The average over all orientations

$$\underline{\underline{D}}\underline{\underline{D}}^{(j)} = d\phi^{(j)}d\theta^{(j)}d\psi^{(j)} \sin\theta^{(j)} \mathcal{P}_{\text{angle}}(\phi^{(j)}, \theta^{(j)}, \psi^{(j)}) \quad (4.7)$$

involves the **angular distribution function** $\mathcal{P}_{\text{angle}}(\phi, \theta, \psi)$, which is the same for each crystallite. In the simplest model we assume a Gaussian distribution for the deviations of the crystallite axis from the fiber axis $\mathcal{P}_{\text{angle}}(\phi, \theta, \psi) \propto \exp(-\theta^2/\theta_0^2)$, while all values of ϕ and ψ between 0 and 2π are equally likely.

4.2.4 Continuous Background

The space between the crystallites is filled with water molecules and strands connecting the crystallites, *i.e.* the amorphous matrix. In the scope of this work, we are not interested in the details of its structure and thus we model it as a continuous background density ϱ_0 , chosen to match the average scattering density of the crystallite (see Fig. 4.6): $\varrho_0 = \sum_{k=1}^K f_k/V_{\text{uc}}$. Here V_{uc} is the volume of the unit cell. There is no background intensity inside the crystallite, achieved in our model by cutting out a spherical cavity $V(\mathbf{r})$ around each atom $\mathbf{r} = \mathbf{r}_{\mathbf{m},k}^{(j)}$. For simplicity we assume a Gaussian cavity

$$V(\mathbf{r}) = \frac{\bar{f}}{(2\pi\xi^2)^{3/2}} \exp\left(-\frac{\mathbf{r}^2}{2\xi^2}\right), \quad (4.8)$$

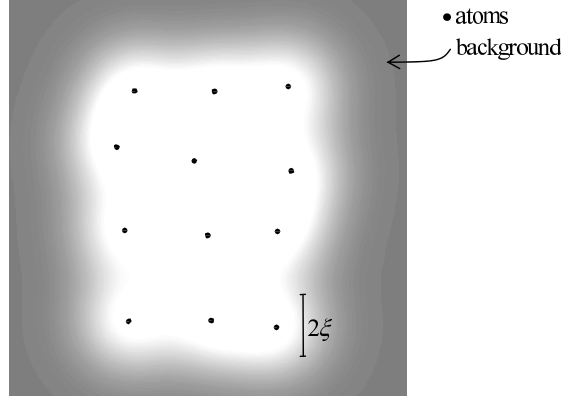


Figure 4.6: Illustration of the continuous background, modeling the amorphous matrix. Outside the crystallites it has a homogeneous scattering density ϱ_0 , matching the mean scattering density of the crystallites. In this illustration ξ is 0.45 of the mean nearest-neighbor distance.

and choose the amplitude such that the average density inside the crystallites is zero:

$$0 = \int_{\text{crystallite}} d\mathbf{r} \left(\varrho_0 - \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^K V(\mathbf{r} - \mathbf{r}_{\mathbf{m},k}) \right), \quad (4.9)$$

where the sum over the vector index $\sum_{\mathbf{m}=1}^{\mathbf{M}}$ means $\sum_{m_x=1}^{M_x} \sum_{m_y=1}^{M_y} \sum_{m_z=1}^{M_z}$.

Using the assumption (4.9), \bar{f} becomes simply the *average* form factor,

$$\bar{f} = \frac{1}{K} \sum_{k=1}^K f_k, \quad (4.10)$$

as can be easily verified. The typical size of the cavity ξ has to be comparable to the nearest-neighbor distance to make sure that there is no “background” inside the crystallites. Models both with and without continuous background are compared in Appendix E.1.

This completes the specification of our model. We now proceed to compute the scattering function as predicted by the model.

4.3 Scattering Function

Given the atomic positions $\mathbf{r}_{\mathbf{m},k}^{(j)}$, the background density $\varrho(\mathbf{r})$ and the statistics of the crystallites’ orientations and positions, we calculate the scattering function:

$$G(\mathbf{q}) = \left\langle \left| \sum_{j=1}^N \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^K f_k \exp(i\mathbf{q}\mathbf{r}_{\mathbf{m},k}^{(j)}) + \int d\mathbf{r} \varrho(\mathbf{r}) \exp(i\mathbf{q}\mathbf{r}) \right|^2 \right\rangle. \quad (4.11)$$

Here, $\varrho(\mathbf{r})$ is the background intensity whose Fourier transform reads as:

$$\begin{aligned} \int d\mathbf{r} \varrho(\mathbf{r}) \exp(i\mathbf{q}\mathbf{r}) &= \int d\mathbf{r} \exp(i\mathbf{q}\mathbf{r}) \left(\varrho_0 - \bar{f} \sum_{j=1}^N \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^K V(\mathbf{r} - \mathbf{r}_{\mathbf{m},k}^{(j)}) \right) \\ &= \varrho_0 V \delta_{\mathbf{q},\mathbf{0}} - \tilde{V}(\mathbf{q}) \sum_{j=1}^N \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^K \exp(i\mathbf{q}\mathbf{r}_{\mathbf{m},k}^{(j)}), \end{aligned} \quad (4.12)$$

with $\tilde{V}(\mathbf{q})$ being the Fourier transform of the cavity $V(\mathbf{r})$. The uniform density ϱ_0 , giving rise to a contribution proportional to $\delta_{\mathbf{q},\mathbf{0}}$, does not contain information about the structure of the system. Furthermore, the central beam has to be gated out in the analysis of the experimental data. Hence, we neglect the uniform contribution and obtain for the scattering intensity (4.11)

$$G(\mathbf{q}) = \left\langle \left| \sum_{j=1}^N \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^K \underbrace{(f_k - \tilde{V}(\mathbf{q}))}_{F_k(\mathbf{q})} \exp(i\mathbf{q} \cdot \mathbf{r}_{\mathbf{m},k}^{(j)}) \right|^2 \right\rangle. \quad (4.13)$$

The cavities give rise to effective form factors $F_k(\mathbf{q}) = f_k - \tilde{V}(\mathbf{q})$ accounting for the scattering of the atoms themselves, f_k , and the cavities surrounding them, $\tilde{V}(\mathbf{q})$. Note that, if we want to switch off the background $\bar{f} \rightarrow 0$, we return to the original form factors $F_k(\mathbf{q}) \rightarrow f_k$.

Inserting the average $\langle \rangle$ and multiplying out the magnitude squared in (4.13) yields:

$$\begin{aligned} G(\mathbf{q}) &= \int \prod_{j=1}^N \left(d\mathbf{R}^{(j)} \mathcal{D}\underline{\underline{D}}^{(j)} \right) \mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)}) \\ &\quad \times \sum_{j,j'=1}^N \exp\left(i\mathbf{q}(\mathbf{R}^{(j)} - \mathbf{R}^{(j')})\right) \\ &\quad \times \sum_{\mathbf{m},\mathbf{m}'=1}^{\mathbf{M}} \sum_{k,k'=1}^K F_k(\mathbf{q}) F_{k'}^*(\mathbf{q}) \exp\left(i\mathbf{q}(\underline{\underline{D}}^{(j)} \mathbf{r}_{\mathbf{m},k} - \underline{\underline{D}}^{(j')} \mathbf{r}_{\mathbf{m}',k'})\right). \end{aligned} \quad (4.14)$$

Note that the double sum over the crystallites j and j' also applies to the rotation matrices $\underline{\underline{D}}^{(j)}$ and $\underline{\underline{D}}^{(j')}$ in the last line. We now split the scattering function into two terms $G(\mathbf{q}) = G_1(\mathbf{q}) + G_2(\mathbf{q})$: the first, $G_1(\mathbf{q})$, includes only the terms $j = j'$ of that sum and thus incorporates scattering of the same crystallite, but not scattering from different crystallites. The second, $G_2(\mathbf{q})$, includes only the terms with $j \neq j'$, taking into account coherent scattering of two different crystallites.

4.3.1 Incoherent Part $G_1(\mathbf{q})$

We first consider the case $j = j'$ of the sum, *i.e.* the contribution to the scattering function which is incoherent with respect to different crystallites. Here, the term

$\exp(i\mathbf{q}(\mathbf{R}^{(j)} - \mathbf{R}^{(j')}))$ gives 1, and therefore the integral over the crystallite positions $\mathbf{R}^{(j)}$ can be performed and trivially gives 1 due to the normalization of the spatial distribution $\mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)})$. Furthermore, for each summand j all integrations over the orientations $\underline{\underline{D}}^{(1)}, \dots, \underline{\underline{D}}^{(N)}$, except $\underline{\underline{D}}^{(j)}$, can be performed and also yield 1. Therefore, the N terms with $j = j'$ simplify to:

$$\begin{aligned} G_1(\mathbf{q}) &= N \int \mathcal{D}\underline{\underline{D}} \sum_{\mathbf{m}, \mathbf{m}'=1}^{\mathbf{M}} \sum_{k, k'=1}^{\mathbf{K}} F_k(\mathbf{q}) F_{k'}^*(\mathbf{q}) \exp(i\mathbf{q}(\underline{\underline{D}}\mathbf{r}_{\mathbf{m},k} - \underline{\underline{D}}\mathbf{r}_{\mathbf{m}',k'})) \\ &= N \int \mathcal{D}\underline{\underline{D}} \left| \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{p=1}^{\mathbf{P}} F_k(\mathbf{q}) \exp(i(\underline{\underline{D}}^T \mathbf{q}) \cdot \mathbf{r}_{\mathbf{m},k}) \right|^2 \\ &= N \int \mathcal{D}\underline{\underline{D}} |A(\underline{\underline{D}}^T \mathbf{q})|^2, \end{aligned} \quad (4.15)$$

where $\underline{\underline{D}}^T$ is the transpose of the matrix $\underline{\underline{D}}$ and

$$A(\mathbf{q}) := \sum_{\mathbf{m}}^{\mathbf{M}} \sum_{k=1}^{\mathbf{K}} F_k(\mathbf{q}) \exp(i\mathbf{q} \cdot \mathbf{r}_{\mathbf{m},k}) \quad (4.16)$$

is the unaveraged scattering amplitude of a single unrotated crystallite. Hence, the interpretation of the incoherent part $G_1(\mathbf{q})$ is straightforward: each crystallite contributes independently and each with a given orientation. The orientation can be absorbed into the scattering vector $\mathbf{q} \rightarrow \underline{\underline{D}}^T \mathbf{q}$, so that the contributions of two crystallites with different orientations are related by a rotation of the scattering vector. In the macroscopic limit, we are allowed to average over all possible orientations and get a sum of N identical terms.

4.3.2 Coherent Part $G_2(\mathbf{q})$

We now consider the contribution to the scattering intensity from *different* crystallites, i.e. the case $j \neq j'$ in Eq. (4.14). In analogy to the above calculation, all integrations over the orientations $\underline{\underline{D}}^{(1)}, \dots, \underline{\underline{D}}^{(N)}$ can be performed except for $\underline{\underline{D}}^{(j)}$ and $\underline{\underline{D}}^{(j')}$:

$$\begin{aligned} G_2(\mathbf{q}) &= \int d\mathbf{R}^{(1)} \dots d\mathbf{R}^{(N)} \mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)}) \\ &\quad \times \sum_{j \neq j'} \exp(i\mathbf{q}(\mathbf{R}^{(j)} - \mathbf{R}^{(j')})) \\ &\quad \times \int \mathcal{D}\underline{\underline{D}} \mathcal{D}\underline{\underline{D}}' \sum_{\mathbf{m}, \mathbf{m}'=1}^{\mathbf{M}} \sum_{k, k'=1}^{\mathbf{K}} F_k(\mathbf{q}) F_{k'}^*(\mathbf{q}) \exp(i\mathbf{q}(\underline{\underline{D}}\mathbf{r}_{\mathbf{m},k} - \underline{\underline{D}}'\mathbf{r}_{\mathbf{m}',k'})) , \end{aligned} \quad (4.17)$$

where we have used that the angular distribution is the same for all crystallites. We introduce the structure factor of the crystallite positions:

$$\begin{aligned} S(\mathbf{q}) &= \frac{1}{N} \left\langle \sum_{j,j'=1}^N \exp(i\mathbf{q}(\mathbf{R}^{(j)} - \mathbf{R}^{(j')})) \right\rangle \\ &= 1 + \frac{1}{N} \int d\mathbf{R}^{(1)} \dots d\mathbf{R}^{(N)} \mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)}) \\ &\quad \times \sum_{j \neq j'} \exp(i\mathbf{q}(\mathbf{R}^{(j)} - \mathbf{R}^{(j')})) \end{aligned} \quad (4.18)$$

and observe that the two upper lines in (4.17) are just $N(S(\mathbf{q}) - 1)$. This simplifies $G_2(\mathbf{q})$ to:

$$\begin{aligned} G_2(\mathbf{q}) &= N(S(\mathbf{q}) - 1) \left| \int \mathcal{D}\underline{D} \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^{\mathbf{K}} F_k(\mathbf{q}) \exp(i(\underline{D}^T \mathbf{q}) \cdot \mathbf{r}_{\mathbf{m},k}) \right|^2 \\ &= N(S(\mathbf{q}) - 1) \left| \int \mathcal{D}\underline{D} A(\underline{D}^T \mathbf{q}) \right|^2 \end{aligned} \quad (4.19)$$

Again the interpretation is straightforward: for coherent scattering the *amplitudes* of individual crystallites with different orientations add up, as expressed by $\int \mathcal{D}\underline{D} A(\underline{D}^T \mathbf{q})$. Spatial correlations of the *centers of the crystallites* are accounted for by the structure function $S(\mathbf{q})$.

The total scattering function

$$\boxed{\frac{G(\mathbf{q})}{N} = \int \mathcal{D}\underline{D} |A(\underline{D}^T \mathbf{q})|^2 + (S(\mathbf{q}) - 1) \left| \int \mathcal{D}\underline{D} A(\underline{D}^T \mathbf{q}) \right|^2} \quad (4.20)$$

is reduced to the scattering amplitude of a *single crystallite* $A(\mathbf{q})$, which we compute next. Note that if the angular spread of the crystallites can be neglected, in other words all crystallites are approximately aligned, then the above expression reduces to $G(\mathbf{q}) = NS(\mathbf{q}) |A(\mathbf{q})|^2$, as would be expected.

If there is thermal motion of the atoms around their equilibrium positions due to finite temperature, the intensity of the scattering function $G(\mathbf{q})$ is weakened for larger q -values [e.g. Willis & Pryor, 1975]. The resulting scattering function has to be multiplied with the Debye-Waller factor

$$G_{\text{DW}}(\mathbf{q}) = G(\mathbf{q}) \cdot \exp(-\mathbf{q}^2 \langle u^2 \rangle), \quad (4.21)$$

where $\langle u^2 \rangle$ is the mean square displacement of the atoms in any direction.

4.3.3 Scattering Amplitude of a Single Crystallite

The calculation of the scattering amplitude of a single crystallite $A(\mathbf{q})$ (defined in Eq. (4.16)) follows standard procedures. We substitute the atomic positions of

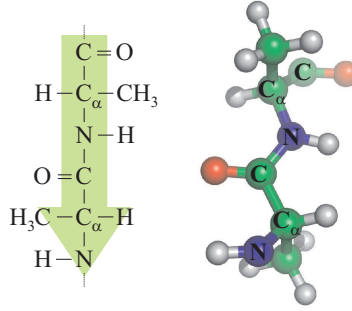


Figure 4.7: Chemical structure (left) and conformation (right) of a polyaniline strand. Two alanines are shown. The CH_3 group is characteristic for the alanine amino acid and is bound to the so called C_α atom. The arrow indicates the direction $\text{C} \rightarrow \text{C}_\alpha \rightarrow \text{N}$ along the backbone.

Sec. 4.2.2 and note that the sums over m_x , m_y and m_z are geometric progressions which can easily be performed. As seen in Appendix E.3, the scattering amplitude becomes:

$$A(\mathbf{q}) = L_{M_x}(\mathbf{q}\mathbf{a}_x)L_{M_y}(\mathbf{q}\mathbf{a}_y)L_{M_z}(\mathbf{q}\mathbf{a}_z) \sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q}\mathbf{r}_k). \quad (4.22)$$

Here $L_{M_\nu}(\mathbf{q}\mathbf{a}_\nu) = \frac{\sin(\mathbf{q}\mathbf{a}_\nu M_\nu/2)}{\sin(\mathbf{q}\mathbf{a}_\nu/2)}$ is the well-known Laue function, which has an extreme value when its argument $\mathbf{q}\mathbf{a}_\nu$ is a multiple of 2π . If M_ν is finite, the peak has a finite width and gets wider with a decreasing number of repetitions M_ν .

It is noteworthy that the extremum of the magnitude of the *scattering amplitude* $A(\mathbf{q})$ may be shifted to a slightly different position, if the form factor of the unit cell $\sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q}\mathbf{r}_k)$ has a non-vanishing gradient at that position. In this case, the resultant peak may be shifted by a value of the order of its peak width. Consequently, care has to be taken when determining lattice constants from experimental peak positions.

4.4 Atomic Configuration of the Unit Cell

The computation of the scattering function $G(\mathbf{q})$ requires the atomic configuration $\{\mathbf{r}_k\}_{k=1}^K$ of the unit cell, which we discuss next.

4.4.1 Unshifted Unit Cells

It is known that the crystallites are composed of polyaniline strands [see [Arnott et al., 1967](#); [Grubb & Jelinski, 1997](#)]. In Fig. 4.7 two alanine amino acids of this strand are shown. Its conformation, shown on the right, is well established and was modeled with Yasara [[Meling, 2006](#)]. There are two constraints for the strand: firstly, the subsequent alanines in the strand must have the same orientation so that the strand does not have a “twist” and can produce periodic structures. And

secondly, the distance between adjacent alanines has to match the size of the unit cell. These two constraints allow for a unique choice of the two degrees of freedom, namely the Ramachandran angles [Ramakrishnan & Ramachandran, 1965] Φ and Ψ , which are the dihedral angles for the bonds C_α -N and C_α -C, respectively.

Many of the described polyalanine strands placed side by side form a stable crystalline configuration, the *β -pleated-sheet*. We assume an orthorhombic unit cell¹, consisting of four alanine strands, as illustrated in Fig. 4.8. Thus one unit cell contains 8 alanine amino acids. Furthermore, we define the spatial directions in the usual way [e.g. Warwicker, 1960]:

- x : Direction of the CH_3 -groups and Van-der-Waals interactions between sheets lying upon each other.
- y : Direction of the hydrogen bonds between the O-atom of one strand and the H-atom of the neighboring strand.
- z : Direction of the covalent bonds along the backbone.

Accordingly, \mathbf{a}_x , \mathbf{a}_y and \mathbf{a}_z are the principal vectors pointing in these directions and a_x , a_y , a_z their magnitudes. Note furthermore that due to symmetry the distance between the strands has to be $a_x/2$ in the x -direction and $a_y/2$ in the y -direction.

In general, one has to distinguish between the **parallel** and **antiparallel** structure. In the parallel structure, the direction of the atom sequence $\text{C} \rightarrow \text{C}_\alpha \rightarrow \text{N}$ in the strand's backbone is the same for all strands (left side of Fig. 4.8). For the antiparallel structure this direction alternates along the y -axis (right side of Fig. 4.8). Both will be considered in the following analysis.

¹Note that the assumption of an orthorhombic unit cell is restrictive. In a more general approach one can give up this assumption and use a smaller unit cell, allowing for different shifts. However, the best fit to the data is obtained for a shift which can also be achieved with an orthorhombic unit cell. For the sake of clarity, we stick to the established unit cell notations and indexing of the reflexes here.

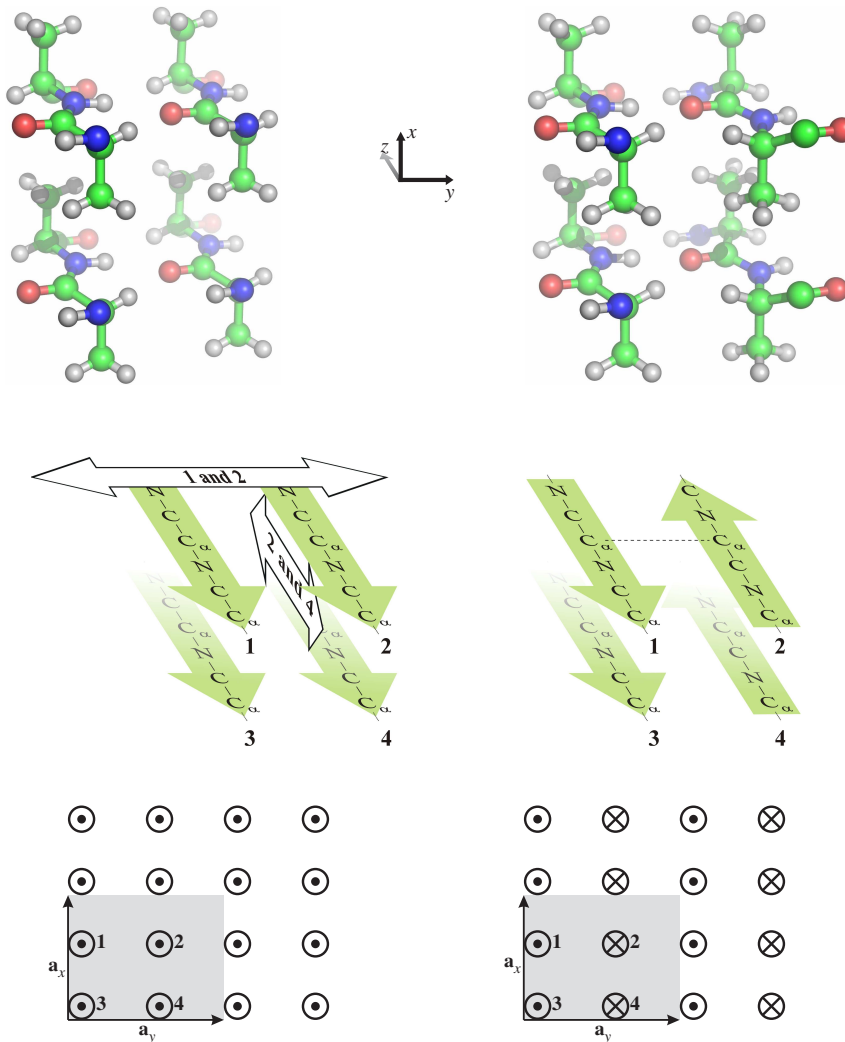


Figure 4.8: Illustration of one unit cell, containing four neighboring polyaniline strands. Top: Configuration of the atoms. Center: Schematic view displaying only the backbone atoms. The white arrows illustrate two possible adjustments of the structure, which are to be optimized to match the experimental scattering image. Bottom: Simplified schematic view along the backbone axis \mathbf{a}_z , where \odot indicates an arrow pointing towards the reader and \otimes pointing away from the reader. Here, the unit cell is shown as a gray rectangle. In each case, a parallel structure is shown on the left side, and an antiparallel structure on the right side. The illustration shows an *unshifted* configuration, which means: the C_α atoms of neighboring strands are aligned and have no shift in z -direction (indicated by the dashed line in the middle right image). Additionally, the strands are exactly aligned in the x - and y -direction.

4.4.2 Possible Shifts inside the Unit Cell

The scattering intensity is not only sensitive to the conformation of the polyalanine strands, but also to the distance and orientation of different strands relative to each other. Besides the question of parallel or antiparallel structure, the four strands can also be shifted with respect to each other. In principle there are four ways to displace them (see Fig. 4.8):

- Shifting strands 1 and 2 in the y -direction by a value Δy_{12} . This displacement is performed in Fig. 4.9.
- Shifting strands 1 and 2 in the z -direction by a value Δz_{12} . Because of the CH_3 -groups extending into the layers above and below (as seen in Fig. 4.8, top panel), a displacement like this is only possible if those two strands have a shift in the y -direction of $\Delta y_{12} \approx \pm a_y/4$ as well. In this case, the CH_3 -groups can pass each other without overlapping.
- Shifting strands 2 and 4 in the z -direction by a value Δz_{24} . This shift was originally suggested by *Arnott et al.* [1967].
- Shifting strands 2 and 4 in the x -direction. However this displacement would have a high energy cost, because it would break the hydrogen bonds between the H- and O-atoms of neighboring strands. Since, moreover, no reasonable result could be achieved performing such shifts, it will not be included in the discussion any further.

Note that shifting strands 1 and 2 in the x -direction, or strands 2 and 4 in the y -direction, is associated with resizing the unit cell in the x - or y -direction, respectively.

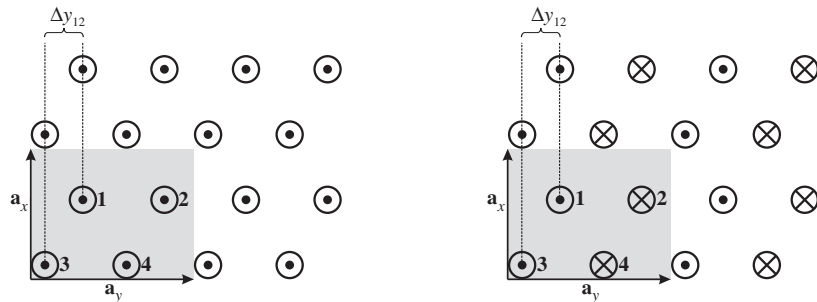


Figure 4.9: Illustration of the unit cell, as at the bottom of Fig. 4.8, but with the conducted shift $\Delta y_{12} = a_y/4$ of strands 1 and 2 the in y -direction.

4.4.3 Variations between Crystallites

For real systems, the composition of the crystallite is certainly not fixed but may vary from crystallite to crystallite. These variations can easily be included into

our model by introducing a probability distribution $P(\{r_{\mathbf{m},k}\})$ for the crystallite configuration $\{r_{\mathbf{m},k}\}$, with the normalization $\sum_{\{r_{\mathbf{m},k}\}} P(\{r_{\mathbf{m},k}\}) = 1$. Note that the probability distribution may be chosen such that the *crystallite size* (M_x, M_y, M_z) is variable and also that the unit cells may have *different types* and *different numbers* of atoms. The calculation is performed completely analogously to the orientational distribution. The scattering function becomes

$$\begin{aligned} \frac{G(\mathbf{q})}{N} &= \sum_{\{r_{\mathbf{m},k}\}} P(\{r_{\mathbf{m},k}\}) \int \mathcal{D}\underline{D} |A(\underline{D}^T \mathbf{q})|^2 \\ &+ (S(\mathbf{q}) - 1) \left| \sum_{\{r_{\mathbf{m},k}\}} P(\{r_{\mathbf{m},k}\}) \int \mathcal{D}\underline{D} A(\underline{D}^T \mathbf{q}) \right|^2. \end{aligned} \quad (4.23)$$

Later (Sec. 4.5.2), we investigate the effect of **variable crystallite sizes** as well as the influence of **small fractions of glycine** inside the crystallites. The effect of the **coherent scattering term** on the bottom line of Eq. (4.23) is analyzed in Appendix E.2.

4.5 Results

4.5.1 Experimental Scattering Function

Two types of samples have been investigated: fiber bundles and single fiber preparations. Single fibers demand a highly collimated and brilliant beam, but are better defined in orientation and are amenable to simultaneous strain-stress measurements.

An oriented bundle of major ampullate silk (MAS) of *Nephila Clavipes* was measured at the D4 bending magnet of HASYLAB/DESY in Hamburg. The fibers were reeled on a steel holder and oriented horizontally in the beam. The number of threads was estimated at 400-600. Photon energy was set to $E = 10.9$ keV by a Ge(111) crystal monochromator, located behind a mirror to suppress higher harmonics. Data was collected by a CCD X-ray camera (SMART Apax, AXS Bruker). Illumination was triggered by a fast shutter. The momentum transfer was calibrated by a standard (corrundum) and raw data was corrected by an empty image (background subtracted).

The single fiber experiments were performed at the microfocus beamline ID13 at ESRF, Grenoble [Riekkel & Davies, 2005]. A 12.7 keV X-ray beam was focused with a pair of short focal length Kirkpatrick Baez mirrors [Kirkpatrick & Baez, 1948] to a 7 μm spot on the sample. This focusing scheme provides a sufficient flux density ($6.8 \cdot 10^{15}$ cps/mm²) to obtain diffraction patterns from single dragline fibers. The single fiber diffraction patterns were recorded with a CCD detector positioned 131 mm behind the sample (Mar 165 detector, Mar USA, Evanston, IL). One of the beamline's custom-made lead beamstops (approx. 300 μm diameter) was used to block the intense primary beam. The raw data was treated as follows: (i) both the image and the background (empty beam) were corrected by dark current, and (ii)

the background was subtracted from the image. The peaks which are significantly broadened by the small crystallite size (see below) can then be indexed by the orthorhombic lattice described above.

Typical scattering distributions for both types of sample preparations are shown in Fig. 4.10, as a function of parallel and vertical momentum transfer. More details on experimental procedures and on the sample preparation by forced silking can be found in [Glišović & Salditt, 2007; Glišović *et al.*, 2008].

4.5.2 Scattering Function from the Model

It is our aim to determine those crystallites' parameters which best match the experimental result. The free parameters of our model are the three shifts Δy_{12} , Δz_{12} and Δz_{24} , the unit cell dimensions a_x , a_y and a_z , the crystallite size in the three directions M_x , M_y and M_z , as well as θ_0 , the tilting angle of the crystallites away from the fiber axis.

These parameters affect the scattering intensity in different ways, allowing us to at least partially separate the effects of different parameters. The crystallite size (M_x, M_y, M_z) determines the peak widths, whereas the length of the principal vectors \mathbf{a}_x , \mathbf{a}_y and \mathbf{a}_z determine the peak position. (We have to keep in mind, however, that the peak position can differ from the extremal values of the Laue functions, as explained in Sec. 4.3.3.) The shifts Δy_{12} , Δz_{12} and Δz_{24} , as described in Sec. (4.4.2), affect the relative peak intensities *via* the form factors of the unit cell $\sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q}\mathbf{r}_k)$. Finally, the parameter θ_0 is responsible for the peak widths in the *azimuthal* direction on the scattering image.

From Eq. (4.22) it is clear that the z -components of the atom positions $\{\mathbf{r}_k\}_{k=1}^K$ are irrelevant for the scattering amplitude $A(\mathbf{q})$ in the xy -plane, *i.e.* for a zero z -component of \mathbf{q} . Therefore, parameters affecting only the z -components – especially the mentioned shifts in the z -direction – will not influence the intensity profile of $G(\mathbf{q})$ in the xy -plane.² Analogously, the scattering profile in the z -direction is independent of parameters influencing the x - and y -directions. Consequently, the sections of the scattering profile *along* and *perpendicular* to the fiber axis can separately be matched to subsets of the parameters. The intensity profile off the z - and xy -axes, taking into account all dimensions of the crystallite, can be seen as a consistency check for the found parameters.

The experimental scattering data clearly reveal a (002) peak, Fig. 4.10. This peak is allowed by symmetry; however, it is extremely weak in the antiparallel structure suggested by Marsh *et al.* [1955b] and shown in Fig. 4.11. The reason is that the electron density within the unit cell projected along the z -axis is almost uniform, varying by approximately 10%. We therefore consider two alternative

²In principle there can be an influence because of the θ -tilt of the crystallites with respect to the fiber axis (see Sec. 4.2.3 and Fig. 4.5). However, the scattering amplitude $A(\mathbf{q})$ shows a *discrete* peak structure and, for small θ -rotations, the out-of-plane reflections (which *are* influenced by the z -components) are too far away from the xy -plane to have an impact on the in-plane intensity profile.

mechanisms generalizing the classical model (Marsh *et al.*) of the antiparallel unit cell. By both mechanisms, the intensity of the (002) peak will increase in agreement with the experiment:

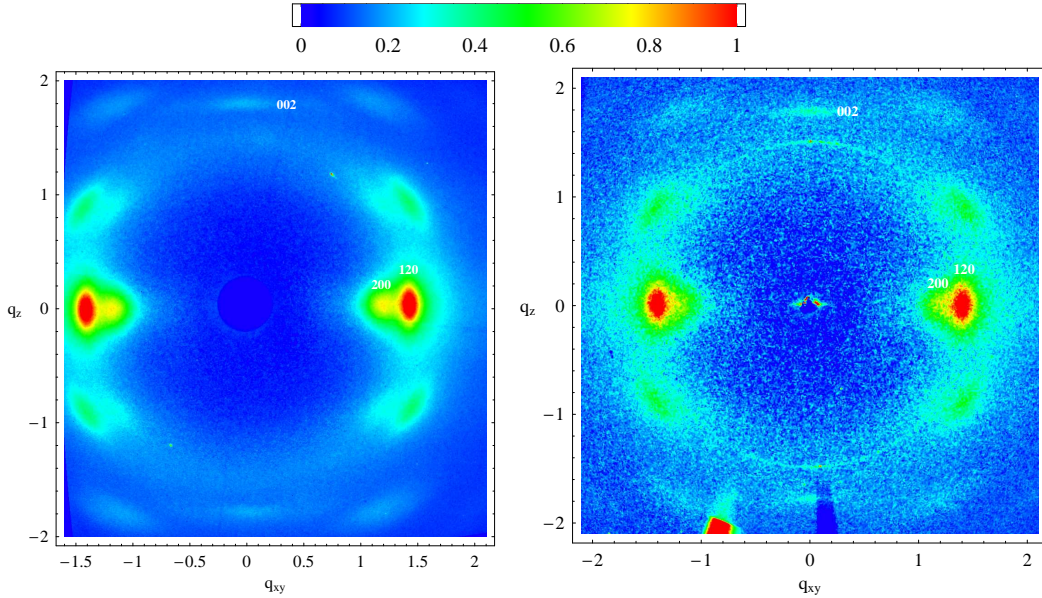


Figure 4.10: Scattering images of spider silk. The experimental scattering image of spider silk from *Nephila clavipes* is shown, both for bundle measurements (left) and single fiber diffraction (right). The fiber axis runs vertically. At the top of the figure the color-bar shows scattering intensities, which are normalized by the intensity of the (120) peak. By courtesy of Tim Salditt.

- a) the **shift of strands 2 and 4** in the z -direction, *i.e.* a non-zero Δz_{24} -shift or
- b) **structural disorder** affecting the almost uniform electron density.

We first discuss case a). The uniform electron density is disturbed by a shift $\Delta z_{24} \neq 0$. The intensity of the (002) reflection grows accordingly with an increasing shift Δz_{24} . Adjusting the Δz_{24} -shift yields results consistent with experiments. Table 4.1 shows the results for the parameters of the model, obtained from optimizing the agreement between the calculated scattering function and the experimental one. For comparison the set of parameters for *both* the parallel and the antiparallel structure are presented. On the basis of the experimental data, one can not discriminate between the parallel and the antiparallel structure.

The scattering intensities, as calculated with these values, are shown in Fig. 4.12. The crystallites are randomly rotated about the fiber axis so that on average the system is invariant under rotations around the fiber axis. Consequently, the scattering image also has rotational symmetry about the z -axis and the q_x - and q_y -axis are indistinguishable and denoted by q_{xy} . A section along the q_{xy} -axis is shown in Fig. 4.13, top panel. The mismatch for q -values slightly larger than the (120) peak

| structure of | presented calculation | | Warwicker [1954] | Marsh <i>et al.</i> [1955 <i>b</i>] | Arnott <i>et al.</i> [1967] |
|-----------------------|-----------------------|--------------------|---------------------|---|--------------------------------|
| | Nephila clavipes | Bombyx mori | Tussah Silk | poly-L-alanine | |
| alignment | parallel | anti-parallel | anti-parallel | anti-parallel | anti-parallel |
| a_x | 10.0 Å | 10.0 Å | 10.6 Å | 10.6 Å | 10.535 Å |
| a_y | 9.3 Å | 9.3 Å | 9.44 Å | 9.44 Å | 9.468 Å |
| a_z | 6.95 Å | 6.95 Å | 6.95 Å | 6.95 Å | 6.89 Å |
| M_x | 1.5 (*) | 1.5 (*) | - | - | - |
| M_y | 5 | 5 | - | - | - |
| M_z | 9 | 9 | - | - | - |
| Δy_{12} | $a_y/4$ | $a_y/4$ | 0 | $a_y/4$ | $\pm a_y/4$ (**) |
| Δz_{12} | 0 | 0 | 0 | 0 | 0 |
| Δz_{24} | 0 | $-a_z/6$ | 0 | 0 | $-a_z/10$ |
| θ_0 | 7.5° | 7.5° | - | - | - |
| $\langle u^2 \rangle$ | 0.1 Å ² | 0.1 Å ² | - | - | - |

Table 4.1: Summary of parameters. The left two columns show the best match between experimental and calculated scattering functions. For $a_z = 6.95$ Å, the resulting Ramachandran angles are $\Phi = -139.0^\circ$ and $\Psi = 136.9^\circ$. $\langle u^2 \rangle$ was used for the Debye-Waller factor in Eq. (4.21). The three right columns compare our obtained parameters with the literature.

(*) Note that each unit cell contains two layers of alanine-strands in x -direction. Therefore $M_x = 1.5$ corresponds to three layers of β -sheets in a single crystallite.

(**) Statistical model: A layer is shifted by a value $+a_y/4$ or $-a_y/4$ with respect to the previous layer, where $+$ and $-$ are equally likely.

is plausible, because in this region the amorphous matrix contributes noticeably to the experimental scattering intensity, but has been neglected in the model. The oscillations of the calculated scattering image for low q -values are side maxima, which are suppressed by fluctuations in the crystallite sizes (see Sec. 4.4.3). The corresponding scattering intensities are shown in Fig. 4.13, and the full scattering image is in Fig. E.2 (left) in Appendix E.2 (page 189). Clearly, the side maxima have been flattened out.

We now discuss an alternative mechanism to generate a stronger (002) peak by introduction of disorder into the amino acid composition of the unit cell (case b). Polyalanine as a model for the crystallites in spider silk is an over-simplification, since the amino acid sequence hardly allows for a pure polyalanine crystallite. Instead, we expect that other residues must be incorporated into the crystallite even if energetically less favorable to compromise the given sequence. In particular, it is highly likely that glycine amino acids are also embedded in the crystallites [Marsh *et al.*, 1955*a*]. This can be easily implemented by replacing randomly selected alanine amino acids of the crystallites with glycine (see Sec. 4.4.3). It is found that the intensity of the (002) peak increases with the fraction of substituted alanines. In Fig. 4.11 we compare the original Marsh-structure (without glycine) to a structure with the same parameters, but with alanine randomly replaced with glycine with a

probability $p_{\text{gl}} = 0.375$. The random substitution has clearly produced an intensity of the (002) peak comparable to the experiment.

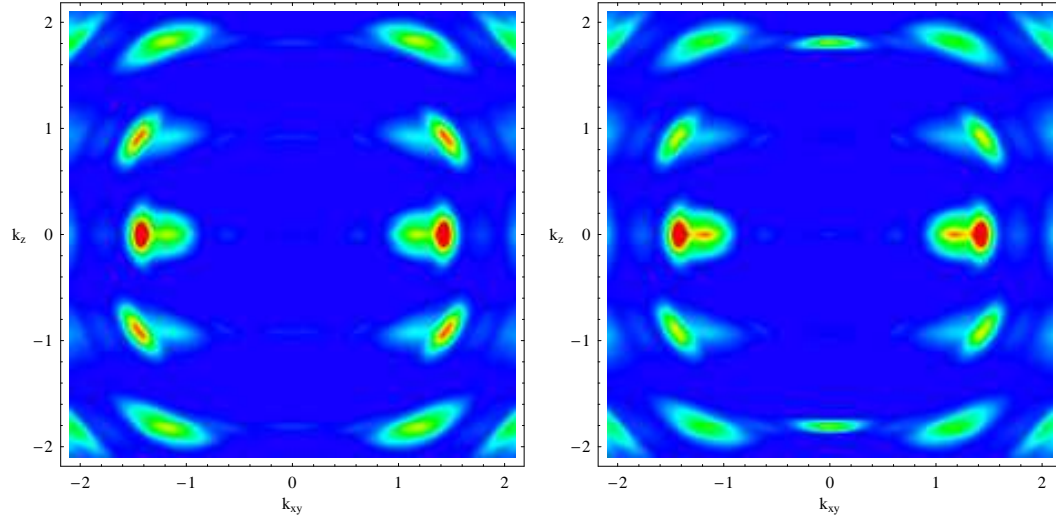


Figure 4.11: Calculated scattering images of the structure proposed by *Marsh et al.* [1955*b*]. The unit cell size is $(a_x, a_y, a_z) = (10.6, 9.44, 6.95)\text{\AA}$, and we used the crystallite size $(M_x, M_y, M_z) = (1.5, 6, 9)$. Left: The crystallites are made purely of alanine amino acids. Right: The crystallites' alanine amino acids are replaced with glycine with a probability $p_{\text{gl}} = 0.375$.

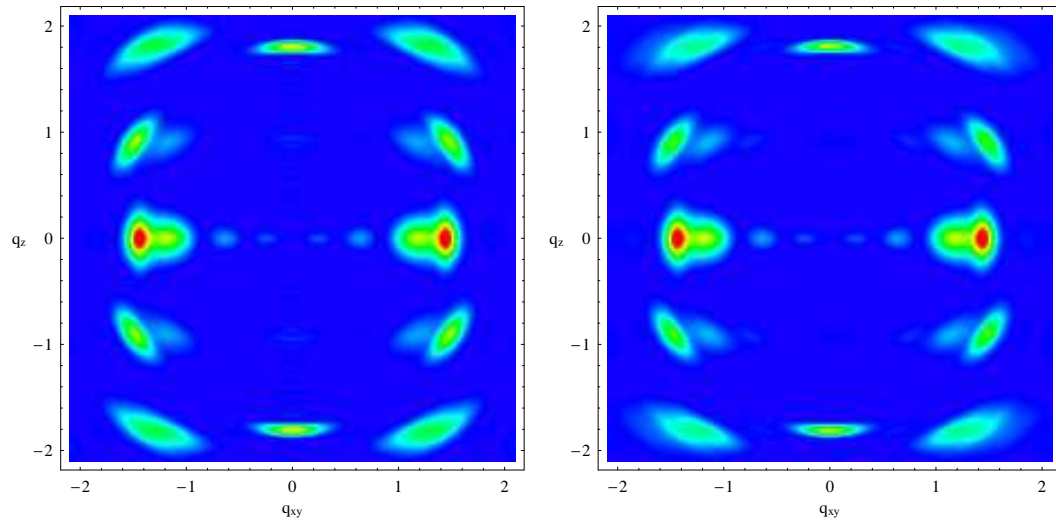


Figure 4.12: Scattering images, as calculated from Eq. (4.20), for the parallel structure on the left side and the antiparallel structure on the right side.

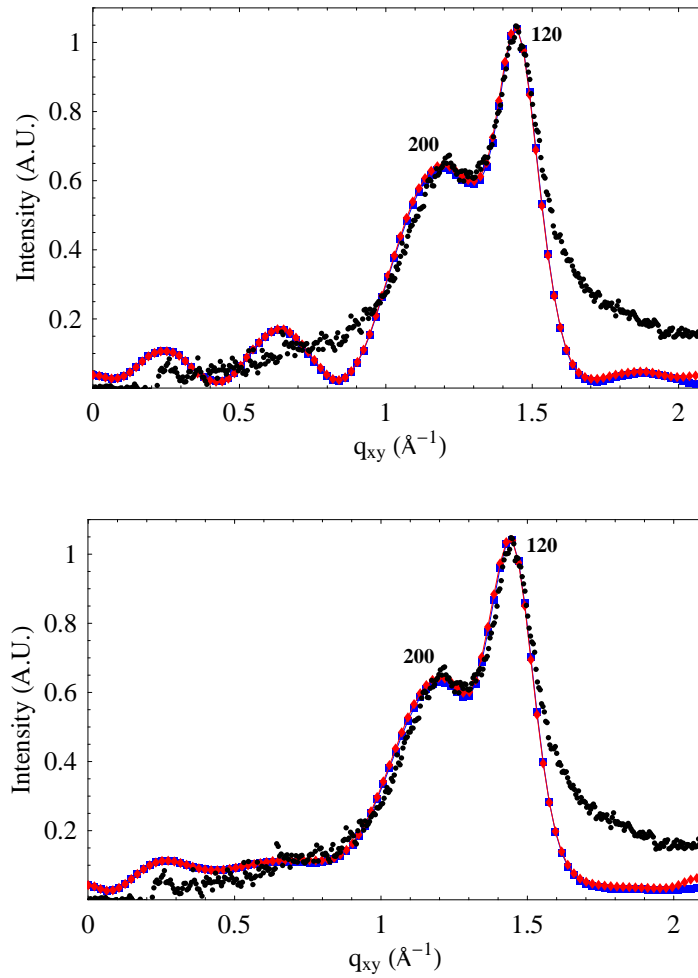


Figure 4.13: Top: Comparison of a section of the experimental (\bullet) and the calculated scattering intensity (\blacksquare parallel, \blacklozenge antiparallel). Sections of the profiles in Fig. 4.12 along the q_{xy} -axis, *i.e.* the scattering profile perpendicular to the fiber axis, are shown. Bottom: As top, but with a Gaussian distribution (rounded to integers) of the crystallite sizes M_x , M_y and M_z . The widths are $\Delta M_x = 2$, $\Delta M_y = 0.75$ and $\Delta M_z = 3$ respectively.

4.6 Conclusions

We have developed a **microscopic model of the structure of spider silk**. The main ingredients of the model are the following:

- a) many small crystallites are distributed randomly in an amorphous matrix,
- b) the orientation of the crystallites fluctuate with a preferential alignment along the fiber axis,
- c) each crystallite is composed typically of $5 \times 2 \times 9$ unit cells,
- d) each unit cell contains four alanine strands shifted with respect to each other. Disorder can be generated by randomly replacing alanine with glycine.

We have computed the scattering intensity of our model and compared it to **wide angle X-ray scattering data of spider silk** from *Nephila clavipes*. Possible **inter-crystallite correlations are unimportant** given the measured orientational distribution. In other words, even if significant center-of-mass correlations between crystallites were present, the orientational distribution would suppress interference effects, with the exception of the (002) peak, which is least sensitive to orientational disorder. The contribution of coherent scattering is discussed in detail in Appendix E.2.

A **homogeneous electron density background** is a necessary feature of the scattering model. Calculation of the crystal structure factor in vacuum does not only lead to an incorrect overall scaling prefactor (which is important if absolute scattering intensities are measured), but also leads to a scattering intensity distribution with artifacts at small and intermediate momentum transfer.

The comparison between model and data fixes the parameters of the unit cell and the crystallite for the two possible cases, the parallel and the antiparallel structure, respectively, as shown in Table 4.1. The two models with parallel and antiparallel alignment of the alanine strands yield comparable agreement with the experimental data. Furthermore a more refined model, where alanine is randomly replaced with glycine, gives reasonable results. Hence we cannot rule out one of these structures.

Our model is similar to the model of the poly-L-alanine of [Arnott *et al.* \[1967\]](#). Their model does incorporate a Δz_{24} -shift. However, our structure shows a better agreement with the experimentally measured scattering function using a value of $\Delta z_{24} = -a_z/6$.

While we have concentrated here on the wide angle scattering reflecting the crystalline structure on the molecular scale, the same model can be used for small angle scattering to analyze the short range order between crystallites in the presence of orientational and positional fluctuations. In particular, the model can describe the entire range of momentum transfer and the transition from wide angle scattering (WAXS) to small angle scattering (SAXS). Note that WAXS is usually described

only in the single object approximation, neglecting inter-particle correlations. Contrarily, SAXS is mostly described in continuum models without crystalline parameters. Here, both are treated by the same approach, which is a significant advantage for systems where the length scales are not decoupled.

Aggregation and Gelation in Wet Granular Materials

Contents

| | | |
|------------|--|------------|
| 5.1 | Introduction | 81 |
| 5.2 | Models | 83 |
| 5.3 | Cooling Dynamics | 86 |
| 5.3.1 | Early Stage of Cooling | 87 |
| 5.3.2 | Late Stage of Cooling | 90 |
| 5.3.3 | Velocity Distribution | 95 |
| 5.3.4 | Partitioning of the Energy into Translational, Rotational and Internal Degrees of Freedom | 98 |
| 5.4 | Aggregation | 101 |
| 5.4.1 | Self-Similar Growth | 103 |
| 5.4.1.1 | Fractal Dimension of the Aggregates | 103 |
| 5.4.1.2 | Cluster Mass Distribution | 105 |
| 5.4.1.3 | Number of Clusters and Mean Cluster Mass | 106 |
| 5.4.2 | Properties of the Asymptotic Cluster | 107 |
| 5.4.2.1 | Fractal Dimension from Radius of Gyration | 108 |
| 5.4.2.2 | Fractal Dimension from Box Counting Algorithm | 110 |
| 5.4.2.3 | Pair Correlation Function and Correlation Dimension | 114 |
| 5.4.2.4 | Coordination Number | 118 |
| 5.4.2.5 | Inertia Tensor and Spatial Extension | 122 |
| 5.5 | Conclusions | 123 |

5.1 Introduction

Granular materials are systems of macroscopic particles, which are typically characterized by two properties: They interact *only on contact* and their interactions are *dissipative*. In spite of their simple interaction, they are of high importance for both science and for industry: On the one hand, because granular systems yield a broad range of interesting nonequilibrium phenomena [Brilliantov & Pöschel, 2004;

Kudrolli, 2004; Umbanhowar *et al.*, 1996] and have been considered as model systems for structure formation on various length scales, including the formation of planetesimals from interstellar dust and the formation of planets and stars from accretion disks [Blum *et al.*, 2000; Bridges *et al.*, 1984]. And on the other hand, storage and handling of bulk solids is among the most significant tasks in industrial technology and still poses a large number of unsolved problems [Jaeger *et al.*, 1996; Duran, 2000]; in addition granular materials are the second-most manipulated material by mankind (the first one being water) [de Gennes, 1999].

Most studies discuss **dry granular systems**, where the dissipative contact interaction consists in the loss of a certain fraction of the kinetic energy in every impact [Duran, 2000; Brilliantov & Pöschel, 2004, and references therein]. Adding a small amount of liquid to the granular system changes its properties dramatically: while dry sand can flow through one's fingers similarly to a liquid, wet sand has properties of a plastic solid, even suitable to build structures like sand castles. This difference in the macroscopic behavior is reflected in the corresponding difference in particle interactions [Herminghaus, 2005]:

Wet granular particles are covered by a **thin liquid film**. When two wet particles come into contact, the films merge and a capillary bridge is formed, exerting an attractive force on the particles. As the particles separate from each other again, the bridge stays intact up to a critical distance d_c . At this point the bridge ruptures [Willett *et al.*, 2000]. Integrating this attractive force over the separation of the particles up to d_c yields the amount of energy which is dissipated in each bond rupture process.

This dissipative mechanism is an essential difference between wet and dry granular systems: The collisions of dry granulates are purely *repulsive* and characterized by the coefficient of restitution ε , which specifies which *fraction of the kinetic energy* is dissipated. Wet granular particles, however, are characterized by a hysteretic *attractive* interaction giving rise to a *fixed amount of energy*, which is dissipated in each rupture.

This existence of a **well defined energy scale**, which is absent in dry materials, is the essential microscopic ingredient not only of wet granulates but also of cohesive gases. In fact the liquid bridge can be thought of as a particular realization of a more general cohesive force. A particularly important aspect of free cooling in cohesive gases is the **aggregation process** which sets in, when the kinetic energy falls below the bond breaking energy. Wet granular systems may provide a realization of various aggregation models and so-called sticky gases [Carnevale *et al.*, 1990], where particles move diffusively or ballistically until they collide and get stuck to an aggregate which is thereby growing. Such models have attracted a lot of interest [Liang & Kadanoff, 1985; Jiang & Leyvraz, 1993, 1994; Carnevale *et al.*, 1990; Alves & Ferreira, 2006; van Dongen & Ernst, 1985; Westbrook *et al.*, 2004; Jullien & Kolb, 1984; Trizac & Krapivsky, 2003; Trizac & Hansen, 1995], due to a wide range of applications ranging from the formation of dust filaments, snowflakes and clouds to the size distribution and impact probability of planetesimals in accretion disks.

Kinetic properties of granular gases have been discussed mainly for dry materials.

In particular, free cooling has been studied extensively [Ben-Naim *et al.*, 1999; Nie *et al.*, 2002; Baldassarri *et al.*, 2002; Krapivsky & Ben-Naim, 2002], and it was shown that the dissipative interactions are responsible for many novel phenomena, unexpected from the kinetic theory of molecular gases: The particles' velocities are not distributed according to a Maxwell-Boltzmann distribution [Goldshtein & Shapiro, 1995], equipartition does not hold [Huthmann & Zippelius, 1997; Garzo & Dufty, 1999; Losert *et al.*, 1999], a spatially homogeneous state is generically unstable [Goldhirsch & Zanetti, 1993; Goldhirsch *et al.*, 1993; McNamara, 1993], and linear and angular motion are correlated [Brilliantov *et al.*, 2007].

Much less is known about *wet* granular media, which have been addressed only recently [Thornton *et al.*, 1996; Lian *et al.*, 1998; Huang *et al.*, 2005; Herminghaus, 2005; Zaburdaev *et al.*, 2006; Fingerle & Herminghaus, 2006, 2008; Fingerle *et al.*, 2008], focussing on nonequilibrium phase transitions [Fingerle *et al.*, 2008], the equation of state [Fingerle & Herminghaus, 2008], agglomeration [Ennis *et al.*, 1991; Thornton *et al.*, 1996; Lian *et al.*, 1998], shear flow [Huang *et al.*, 2005], and cooling in one dimension [Zaburdaev *et al.*, 2006; Fingerle & Herminghaus, 2006].

Structure formation in wet granulates during free cooling is the focus of this chapter, which is organized as follows: In Sec. 5.2 the model is introduced, and the decay of the average kinetic energy is discussed in Sec. 5.3. Aggregation is investigated in Sec. 5.4, before we conclude in Sec. 5.5. A short summary of the presented results has appeared in [Ulrich *et al.*, 2009b] and an even shorter one in [Ulrich *et al.*, 2009a].

5.2 Models

We are interested in the zero-gravity free cooling dynamics of wet granular gases. We assume the particles to be covered by a thin liquid film, as it is the case if the liquid completely wets the particle material [Israelachvili, 1992]. The particles move freely, until these surface films come into contact. The liquid then rapidly accumulates around the contact due to the interfacial forces. A capillary bridge forms at the contact, exerting an attractive force on the grains due to its negative Laplace pressure. This liquid bridge is stretched but stays intact (or even continues to grow) as the particles move apart. The attractive force thus remains until a certain critical separation s_c is reached, where the liquid neck becomes unstable and ruptures. As mentioned above, the hysteretic formation and rupture of the bridge gives rise to a characteristic loss of energy, ΔE , which depends upon the thickness of the liquid film wetting the grains.

In order to design a suitable model, we briefly discuss the nature of this process. The formation of capillary bridges is quite fast in real systems. Between typical grains of one millimeter diameter it takes less than a millisecond [Herminghaus, 2005, Sec. II D]. It is clear, however, that this formation cannot in general be considered instantaneous if the velocity of the impacting grains v_i is large. If the time scale of the impact process, which may be written as s_c/v_i , is of the same order

or even smaller than the time of capillary bridge formation, the accumulated liquid volume of the bridge, and hence ΔE , will be smaller than for slow impacts. However, this will not greatly affect the main features of the wet system, in particular as to its characteristic difference from the dry granulate. In order to see that, we compare the effective restitution coefficient of the dry and of the wet system. This is shown in Fig. 5.1, where the restitution coefficient for the dry system is shown as the dotted curve. It tends to be mildly depending on impact energy [Brilliantov & Pöschel, 2004], E_i , with a negative slope throughout. The effective restitution coefficient of the wet system, $\varepsilon_{\text{eff}} = \sqrt{1 - \Delta E/E_i}$, is shown as the solid curve, assuming constant ΔE . In strong contrast to the dry system, it has a zero at $\Delta E/E_i = 1$ and a markedly positive slope. This illustrates the dramatic difference between these two systems. The dashed line qualitatively accounts for the effect of finite formation time of the capillary bridge. Since ε_{eff} must stay below one, the qualitative picture concerning the comparison of dry and wet granular gases remains unchanged.

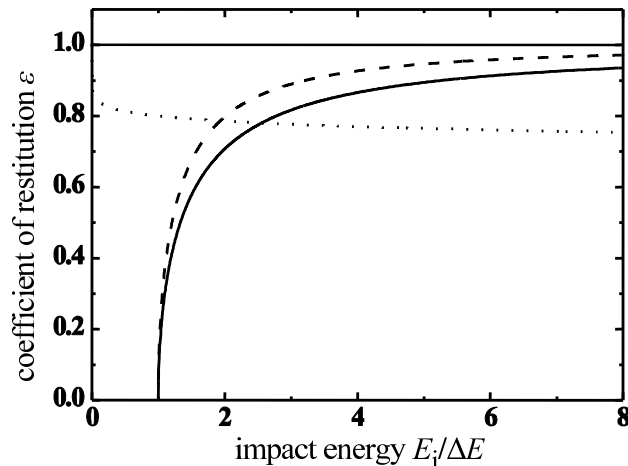


Figure 5.1: Restitution coefficients for dry (dotted) and wet (solid and dashed) granular systems, plotted vs. the impact energy in units of the wet energy loss, ΔE . The main feature in the wet case is the zero at $E_i = \Delta E$, which is unchanged if the finite formation time for capillary bridges is taken into account (dashed curve). The horizontal line $\varepsilon = 1$ corresponds to the limit of fully elastic spheres.

Our system consists of N identical and spherical particles with diameter d and mass m in a three-dimensional cubic volume $V = L^3$. They are at positions $\mathbf{r}_1, \dots, \mathbf{r}_N$ and have the velocities $\mathbf{v}_1, \dots, \mathbf{v}_N$. The particles have a hard-core interaction, such that two particles are *reflected elastically*, if their centers of mass reach the hard-core distance, which is the particle diameter d .

To account for the liquid film, a liquid bridge is allowed to form between a pair of particles if they come close enough (“close enough” is specified later). When these particles are moving apart and their distance exceeds the bond breaking distance d_c , the liquid bridge will break and a fixed amount of kinetic energy ΔE is dissipated; in that process, momentum is conserved and the relative velocity v_{rel} changes to v'_{rel}

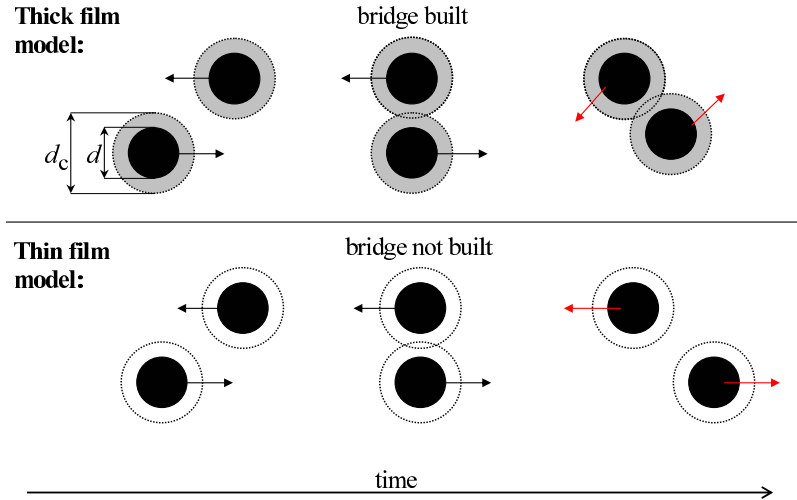


Figure 5.2: Illustration of the thin film model and thick film model. In the thick film model, the liquid bridge forms, as soon as the bond breaking distances d_c overlap. The same initial configuration in the thin film model does not create a liquid bridge, since the hard cores of the particles do not touch. Thus, the particles just pass by.

according to

$$\frac{\mu}{2}v_{\text{rel}}^2 = \frac{\mu}{2}v_{\text{rel}}^2 - \Delta E \quad (5.1)$$

with the reduced mass $\mu = m/2$. If, however, the relative kinetic energy is smaller than ΔE , the particles are elastically reflected towards each other. In this case the two particles oscillate between d and d_c , until a collision with a third particle or a wall supplies enough energy to break the liquid bridge.

The effect of the capillary force, which is present in reality for distances up to d_c , is thus solely modeled by the energy loss which occurs at the distance d_c . This has been shown before to be a very good approximation [Fingerle *et al.*, 2008], and enables event-driven simulations as discussed below. For the formation of the liquid bridge, we distinguish between two models:

In the *thick film model*, a liquid bridge forms as soon as particles come closer than the critical bond breaking distance d_c . This model assumes that the outer diameter of the liquid film is d_c and its shape stays spherical and is not deformed by the particles.

In the *thin film model*, the liquid bridge forms when the particles touch, *i.e.* the distance of their centers is equal to d . This model assumes that the liquid film covering the particles is infinitesimally thin and the capillary bridges form a thin liquid neck, which breaks off at the critical distance d_c . Raw data for this model was provided by courtesy of the Max Planck Institute for Dynamics and Self-Organization.

As it will turn out, the differences in most of the results are only minute. The two models are illustrated in Fig. 5.2. Note that in the limit $d_c \rightarrow d$ the difference between the two models vanishes, since in a collision the bond breaking distance

and hard core distance are always passed concurrently. For the sake of clarity, in all graphs throughout this chapter data points displayed as *filled symbols* (like \bullet) represent the *thick film model* and *open symbols* (like \circ) the *thin film model*.

In general there is some energy being transferred to the atomic degrees of freedom of wet grains as well. We are going to neglect this dissipation mechanism for our simple model, because it is usually small as compared to the energy loss due to the breaking of capillary bridges, especially if the granular temperature is small. It is evident, however, that such a dissipation mechanism can easily be incorporated in the simulations, replacing the elastic reflection by incomplete normal restitution. We restrict ourselves here to perfectly smooth particles, such that translational and rotational motion are decoupled. Furthermore, we investigate free cooling only, so no gravity is present, and no energy is injected into the system.

The particular way of accounting for the liquid film used in these models makes it possible to use an event-driven simulation scheme. The possible events are the reflection of the particles at the hard core distance d and the crossing of the bond-breaking distance d_c , at which the bond-breaking energy ΔE is dissipated. [Fingerle et al. \[2008\]](#) have previously compared event-driven simulations of the wet system with full molecular dynamics simulations, integrating the equation of motion; they found good quantitative agreement in the results of both methods, justifying the event-driven approach chosen exclusively for the present study.

We use dimensionless units such that $\Delta E = 1$, particle mass $m = 1$ and particle diameter $d = 4$. The bond-breaking distance is chosen as $d_c = 1.07d$, unless noted otherwise, and volume fraction, $\phi = \pi d^3/6 \cdot N/V$, is varied from $\phi \approx 0.06\%$ up to 15.6% . We use periodic boundary conditions in the x - and y -direction and hard walls in z -direction. The hard walls are completely elastic and do not exert capillary interactions with the particles.

5.3 Cooling Dynamics

The central quantity in this section is the granular temperature T , defined *via*

$$\frac{3}{2} T = \frac{1}{N} \sum_{i=1}^N \frac{m \mathbf{v}_i^2}{2}. \quad (5.2)$$

We will investigate its decay in time from a given initial value $T_0 \gg \Delta E$; in all the simulations $T_0 = 45\Delta E$ was chosen.

Simple arguments can be used to derive an analytical form of the temperature decay. Therefore we first need the collision frequency f_{coll} , which is defined as the average number of collisions per time for a given particle. Since each collision involves *two* particles, $f_{\text{coll}} \cdot N \cdot \Delta t$ adds up to *twice* the number of collisions in the whole system during the short time interval Δt . In each collision a capillary bridge ruptures with probability P_{bb} , giving rise to dissipation of a fixed amount of energy, the bond-breaking energy ΔE . Hence, during this time the total kinetic

energy decreases by

$$\Delta E_{\text{kin}} = \Delta E \cdot \frac{N f_{\text{coll}} \Delta t}{2} \cdot P_{\text{bb}} . \quad (5.3)$$

Dividing by N and Δt yields, in the limit $\Delta t \rightarrow 0$, a rate equation for the temperature:

$$\frac{3}{2} \frac{dT}{dt} = -\frac{1}{2} \cdot f_{\text{coll}} \cdot \Delta E \cdot P_{\text{bb}} . \quad (5.4)$$

5.3.1 Early Stage of Cooling

In the early stage of cooling, the average kinetic energy per particle is much larger than the bond breaking energy, so that $P_{\text{bb}} \approx 1$ and almost every collision gives rise to dissipation by ΔE . For a dilute gas, the collision frequency

$$f_{\text{coll}} = 4g(d)\sigma n \sqrt{\frac{T}{\pi m}} \quad (5.5)$$

is well established [e.g. Brilliantov & Pöschel, 2004], with the particle density $n = N/V$ and the pair correlation function at contact $g(d)$, for which we use the Carnahan-Starling approximation [Carnahan & Starling, 1969], $g(d) = \frac{(2-\phi)}{2(1-\phi)^3}$. The two models differ only in the scattering cross section σ (see Fig. 5.2), which is given by $\sigma = d^2\pi$ in the thin film model and $\sigma = d_c^2\pi$ in the thick film model.

Applying Eq. (5.5) and $P_{\text{bb}} = 1$ to Eq. (5.4), gives rise to the following simple equation:

$$\frac{dT}{dt} = -\frac{4}{3}g(d)\sigma n \sqrt{\frac{T}{\pi m}} \Delta E \propto -\sqrt{T} , \quad (5.6)$$

which is solved by $T(t) \sim (t - t_0)^2$. Inserting the prefactors and setting the initial value $T(0) = T_0$, we obtain, similar to Haff's law [Haff, 1983], an analytical form of the decay of the temperature:

$$T(t) = \begin{cases} T_0 (1 - t/t_0)^2 & \text{for } t \leq t_0 \\ 0 & \text{for } t > t_0 \end{cases} \quad (5.7a)$$

with a characteristic time scale

$$t_0 = \frac{3\sqrt{\pi m T_0}}{2g(d)\sigma n \Delta E} . \quad (5.7b)$$

Note that, in this simplified model, the assumption that *every* collision causes an energy loss ΔE gives rise to a time-scale t_0 after which all energy is dissipated. Even though this assumption does not hold for all times in the simulation (since the bonds do not break anymore if the relative kinetic energy is too small), the timescale

t_0 has a clear physical relevance. It sets the time after which the temperature is comparable to the bond-breaking energy ΔE and after which persistent clusters will form.

Now, let us have a look at the evolution of the granular temperature T from the simulation and compare it to (5.7): In Fig. 5.3 the temperature of the thick film model is shown for different volume fractions. One can see that the agreement between theory and simulation is quite good up to t_0 , when the assumption contained in Eq. (5.4) that *every* collision causes an energy loss ΔE breaks down.

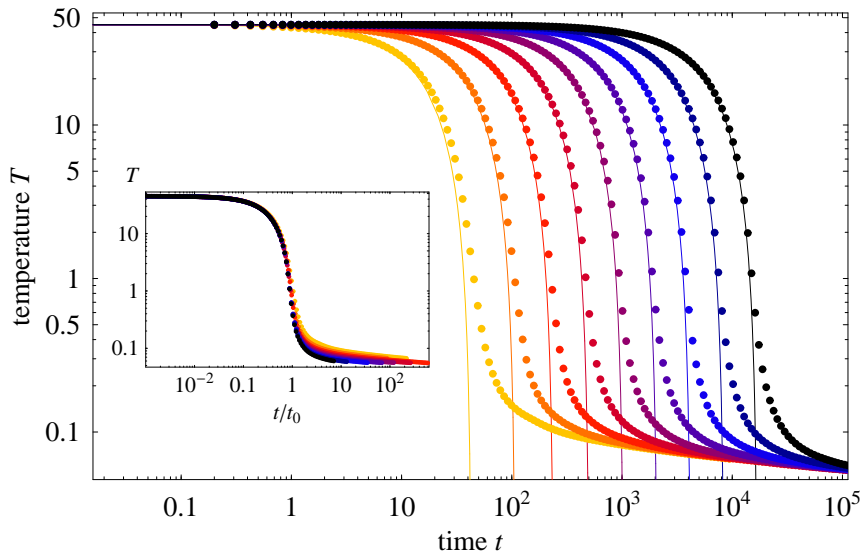


Figure 5.3: Decay of the granular temperature T for volume fractions (from left to right) $\phi = 15.6\%$, 7.81% , 3.90% , 1.95% , 0.98% , 0.49% , 0.24% , 0.12% , 0.061% ; the number of particles $N = 262144$ is fixed; comparison of theory (full line) and simulation (\bullet). Inset: T versus scaled time t/t_0 .

In the simplified cooling law (5.7a), the volume fraction only enters into t_0 . Hence we can try to superimpose the data by scaling time with t_0 . As can be seen in the inset of Fig. 5.3, the data obey the expected scaling well, except for the long time limit, which has different asymptotic behavior and is treated in the next section, Sec. 5.3.2.

In Fig. 5.4, the temperature evolution of the thick and thin film model is compared. The difference of the two models is solely due to different scattering cross-sections, entering in t_0 , Eq. (5.7b). Again, we observe good agreement of the simplified theory and the simulation up to the time t_0 . In the inset the time is rescaled with t_0 .

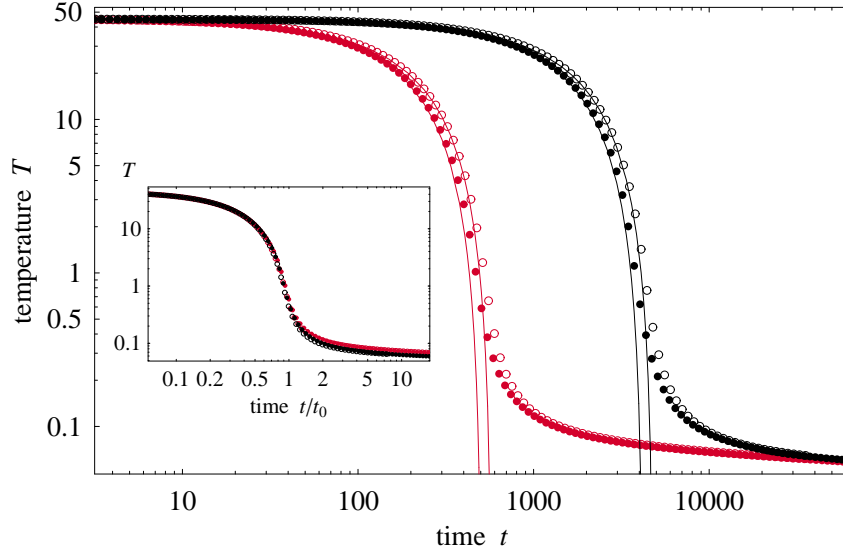


Figure 5.4: Decay of the granular temperature T for the thick film model (\bullet) and the thin film model (\circ) and for volume fractions $\phi = 1.95\%$ (red) and $\phi = 0.24\%$ (black). The inset shows T versus scaled time t/t_0 .

In Fig. 5.5, we investigate how the the bond breaking distance d_c influences the cooling behavior. As mentioned above, this will influence the scattering cross section $\sigma = d_c^2\pi$ for the thick film model and according to Eq. (5.7b), and t_0 should increase when decreasing d_c . This behavior can indeed be observed on the left side of Fig. 5.5, which shows the data for the thick film model.

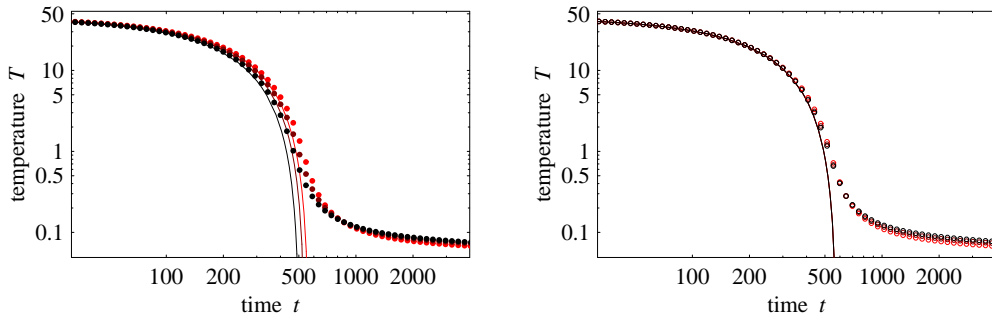


Figure 5.5: Decay of the granular temperature T for volume fraction $\phi = 1.95\%$ and $N = 262144$ particles. The bond breaking distances are $d_c = 1.07d$ (black), $d_c = 1.035d$ (brown) and $d_c = 1.01d$ (red). The left graph shows data for the thick film model, the right graph for the thin film model.

For the thin film model, the scattering cross section $\sigma = d^2\pi$ depends only on the hard core diameter d of the particles. Hence varying d_c should not influence the time t_0 . This can be seen on the right hand side of Fig. 5.5, where the different curves of

the thin film model coincide well, until the time t_0 is passed and aggregation starts.

Surprisingly one can see that in the aggregated state $t > t_0$ a smaller bond breaking distance d_c slightly expedites the cooling process, for both models. This is probably because a smaller d_c increases the collision frequency of the particles inside clusters.

5.3.2 Late Stage of Cooling

In the late stage of aggregation, when the system is strongly aggregated, it becomes very unlikely that a capillary bridge ruptures and hence we observe a very slow time evolution of our system. The slow decrease of the temperature can be understood with simple arguments. Therefore we start with Eq. (5.4),

$$\frac{3}{2} \frac{dT}{dt} = -\frac{1}{2} f_{\text{coll}} \Delta E P_{\text{bb}}, \quad (5.8)$$

as in the previous section, but omit the assumption $P_{\text{bb}} \approx 1$. The probability P_{bb} to break a bond is given by the probability to find a kinetic energy larger than ΔE :

$$P_{\text{bb}} = \int d\mathbf{v} \theta(m\mathbf{v}^2/2 - \Delta E) w(\mathbf{v}), \quad (5.9)$$

where $w(\mathbf{v})$ is the velocity distribution. As we will see in Sec. 5.3.3, in the clustered state it is reasonable to approximate it by a Maxwellian

$$w(\mathbf{v}) = \left(\frac{m}{2\pi T(t)} \right)^{3/2} \exp\left(-\frac{m\mathbf{v}^2}{2T(t)} \right). \quad (5.10)$$

As it is usually done for spherical symmetrical integrals like (5.9), we can integrate out the angular variables, yielding a factor of $4\pi v^2$, and then consider the dependence of the absolute value $v = |\mathbf{v}|$:

$$\begin{aligned} P_{\text{bb}} &= 4\pi \left(\frac{m}{2\pi T(t)} \right)^{3/2} \int_{\sqrt{2\Delta E/m}}^{\infty} dv v^2 \exp\left(-\frac{mv^2}{2T(t)} \right) \\ &= \sqrt{\frac{4\Delta E}{\pi T}} \exp\left(-\frac{\Delta E}{T} \right) + \text{erfc}\left(\sqrt{\frac{\Delta E}{T}} \right). \end{aligned} \quad (5.11)$$

We are interested in the asymptotic limit $T(t)/\Delta E \rightarrow 0$. Looking up the asymptotic expansion of the complementary error function [e.g. [Stegun & Abramowitz, 1954](#), Ch. 7],

$$\text{erfc}(x) = \frac{\exp(-x^2)}{x\sqrt{\pi}} (1 + \mathcal{O}(x^{-2})) \quad \text{for } x \rightarrow \infty, \quad (5.12)$$

we realize that the contribution of the $\text{erfc}(\dots)$ -term only makes a lower order contribution in the limit $T/\Delta E \rightarrow 0$ and hence:

$$P_{\text{bb}} = \sqrt{\frac{4\Delta E}{\pi T}} \exp\left(-\frac{\Delta E}{T} \right) \cdot \left(1 + \mathcal{O}\left(\frac{T}{\Delta E} \right) \right). \quad (5.13)$$

The collision frequency f_{coll} is not known for the clustered state, but it should be proportional to $T^{1/2}$: Increasing all particle velocities for a given system by a factor of two must also increase the collision frequency by a factor of two:

$$f_{\text{coll}} = \gamma T^{1/2} . \quad (5.14)$$

Roughly speaking, the prefactor γ contains information about how large the local density around a given particle is, *i.e.* how many close-by particles a given particles has to collide with. Hence, γ will clearly be higher in the clustered state than in the homogeneous state. By assuming that γ is not time dependent itself, we assume that the internal structure within the clusters does not change significantly as $t \rightarrow \infty$ (while the cluster is still allowed to change its overall shape). This seems to be a good assumption in the highly clustered state, however, it certainly can not correctly describe the transition state, where particles conglomerate around a given particle and the local density changes drastically¹.

With that, the rate equation (5.8) becomes:

$$\begin{aligned} \frac{dT}{dt} &= -\frac{\Delta E}{3} \gamma T^{1/2} \sqrt{\frac{4\Delta E}{\pi T}} \exp\left(-\frac{\Delta E}{T}\right) \cdot \left(1 + \mathcal{O}\left(\frac{T}{\Delta E}\right)\right) \\ \frac{1}{\Delta E} \frac{dT}{dt} &= -\nu \exp\left(-\frac{\Delta E}{T}\right) \cdot \left(1 + \mathcal{O}\left(\frac{T}{\Delta E}\right)\right) , \end{aligned} \quad (5.15)$$

where we defined $\nu = \frac{2}{3\sqrt{\pi}} \cdot \gamma \Delta E^{1/2}$, which has the dimension of a frequency². To solve this equation, it is useful to introduce the dimensionless inverse temperature:

$$x := \frac{\Delta E}{T} . \quad (5.16)$$

With that, Eq. (5.15) becomes:

$$\begin{aligned} \frac{d(x^{-1})}{dt} &= -\frac{1}{x^2} \frac{dx}{dt} = -\nu e^{-x} \cdot (1 + \mathcal{O}(x^{-1})) \\ \Rightarrow \frac{dx}{dt} &= \nu x^2 e^{-x} \cdot (1 + \mathcal{O}(x^{-1})) . \end{aligned} \quad (5.17)$$

This differential equation can be solved by separation of variables:

$$\int dx x^{-2} e^x \cdot (1 + \mathcal{O}(x^{-1})) = \nu \int dt . \quad (5.18)$$

The integration of the right hand side is trivial. For the left hand side we find by repeated partial integration that:

$$\int dx x^{-2} e^x = x^{-2} e^x + \mathcal{O}(x^{-3} e^x) , \quad (5.19)$$

$$\text{and } \int dx x^{-3} e^x = x^{-3} e^x + \mathcal{O}(x^{-4} e^x) . \quad (5.20)$$

¹Indeed, attempts to describe the temperature evolution in the transition state with this formalism failed.

²comparing it to Eq. (5.14) indicates, it is roughly the collision frequency, when $T = \Delta E$.

Hence Eq. (5.18) becomes:

$$x^{-2}e^x \cdot (1 + \mathcal{O}(x^{-1})) = \nu t + c, \quad (5.21)$$

where c is an integration constant, which has to be chosen to match the initial condition. Figuratively speaking, we can use c to shift the solution $x(t)$ along the time axis back and forth and thereby run through the set of solutions of the differential equation (5.17). However, we have to keep in mind that this equation is derived in the limit $x = \Delta E/T \gg 1$; therefore this initial condition has to be set in the low temperature regime, as well, and cannot be chosen as $T(0) = T_0 \gg \Delta E$ as in the previous section.

In order to see how the temperature evolves asymptotically in the limit $t \rightarrow \infty$, we need to invert Eq. (5.21), which is done in Appendix D.2:

$$x(t) = \ln \tilde{t} + 2 \ln \ln \tilde{t} + \frac{4 \ln \ln \tilde{t}}{\ln \tilde{t}} + \mathcal{O}\left(\frac{1}{\ln \tilde{t}}\right), \quad (5.22)$$

with $\tilde{t} = \nu t + c$. If we turn back to the actual temperature and resubstitute $T(t)/\Delta E = x^{-1}(t)$, we get in highest possible order from Eq. (5.22):

$$\frac{T(t)}{\Delta E} = \frac{1}{\ln \tilde{t} + 2 \ln \ln \tilde{t} + \frac{4 \ln \ln \tilde{t}}{\ln \tilde{t}}} + \mathcal{O}\left(\frac{1}{(\ln \tilde{t})^3}\right) \quad (5.23a)$$

and lower orders

$$\frac{T(t)}{\Delta E} = \frac{1}{\ln \tilde{t} + 2 \ln \ln \tilde{t}} + \mathcal{O}\left(\frac{\ln \ln \tilde{t}}{(\ln \tilde{t})^3}\right) \quad (5.23b)$$

$$= \frac{1}{\ln \tilde{t}} + \mathcal{O}\left(\frac{\ln \ln \tilde{t}}{(\ln \tilde{t})^2}\right). \quad (5.23c)$$

If furthermore $\nu t \gg c$ we can also neglect c . This is because the terms $\ln(\nu t + c)$ and $\ln \ln(\nu t + c)$ produce corrections of at least $\mathcal{O}(\frac{c}{\nu t})$, which one can easily confirm. Hence we find for the leading order in t :

$$\frac{T(t)}{\Delta E} = \frac{1}{\ln \nu t} + \mathcal{O}\left(\frac{\ln \ln \nu t}{\ln \nu t}\right). \quad (5.23d)$$

This result shows that the temperature exhibits a very slow, logarithmic time decay for $t \rightarrow \infty$, induced by the very low probability to break a bond, Eq. (5.13); this is in strong contrast to the algebraic time decay observed for granular systems with coefficient of restitution $\varepsilon < 1$ [Brilliantov & Pöschel, 2004].

The different orders of the approximations (5.23) as well as the numeric solution of the original differential equation (5.8) are shown in Fig. 5.6 and compared to the simulation data. The unknown prefactor γ , or equivalently ν , is a fit parameter. As one can see, in the limit $t \rightarrow \infty$, all curves show the same asymptotic behavior, as expected. Approaching the transition time $t \rightarrow t_0$, the lowest order approximation (5.23d) clearly deviates first, while already the second lowest order shows (5.23b) roughly the same conformance as the higher orders.

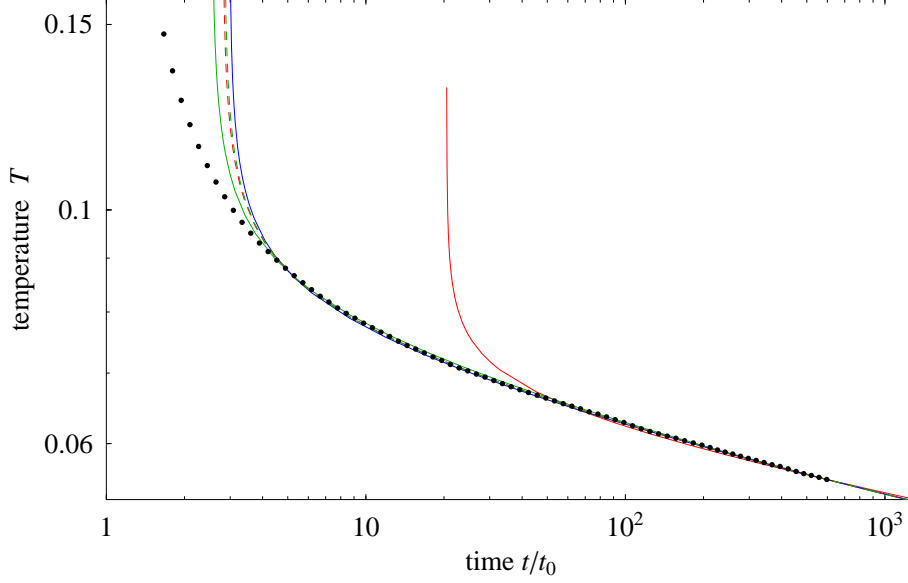


Figure 5.6: Asymptotic temperature evolution for a system with $\phi = 3.9\%$ and $N = 262144$. The data points show the simulation data, while the lines show analytical forms with different orders of precision: The red line is the **lowest order** approximation (5.23c), the green line the **second lowest** (5.23b) and the blue line the **third lowest** (5.23a). The dashed lines are numerical solutions of Eq. (5.8) with P_{bb} given by Eq. (5.11) (**red**) and its first order approximation Eq. (5.13) (**green**).

In Fig. 5.7, the second order solution (5.23b) is compared to the simulation data for different volume fractions, showing good agreement. One can see that for all densities, the temperature approaches a universal curve as $t \rightarrow \infty$. Considering the logarithmic decay Eq. (5.23c), this is not surprising: Even if two systems have different values for ν , let's say ν_1 and ν_2 , then in the limit $t \rightarrow \infty$

$$\begin{aligned} \frac{T_{\nu_2}(t)}{\Delta E} &= \frac{1}{\ln \nu_2 t} = \frac{1}{\ln \nu_1 t + \ln \nu_2 / \nu_1} = \frac{1}{\ln \nu_1 t} + \mathcal{O}((\ln \nu_1 t)^{-2}) \\ &= \frac{T_{\nu_1}(t)}{\Delta E} + \mathcal{O}((\ln \nu_1 t)^{-2}) . \end{aligned} \quad (5.24)$$

Hence, the difference between the curves vanishes.

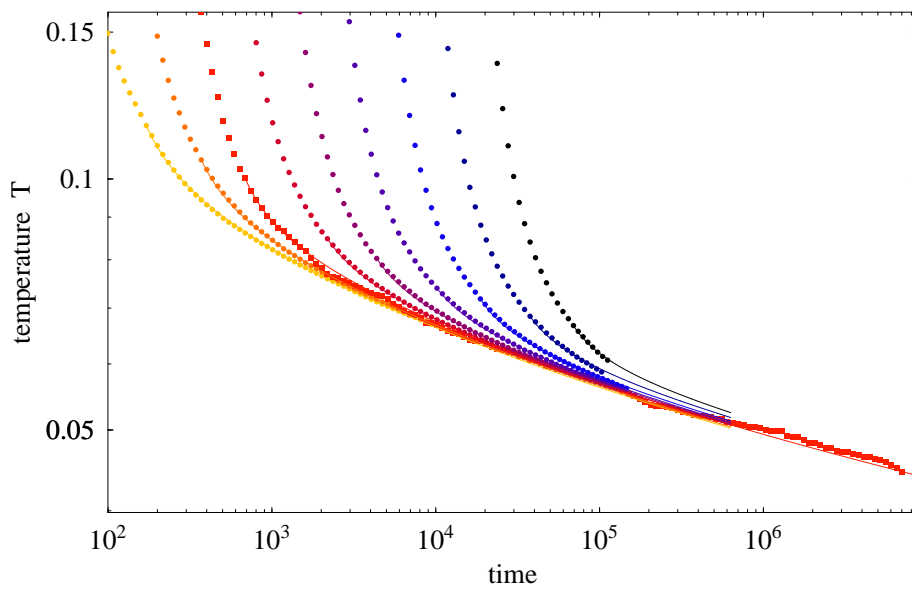


Figure 5.7: Asymptotic time dependence of the temperature for volume fractions (from left to right) $\phi = 15.6\%$, 7.81% , 3.90% , 1.95% , 0.98% , 0.49% , 0.24% , 0.12% , 0.061% . The number of particles is $N = 262144$, except for the third curve (\blacksquare), which has $N = 8192$ particles and is therefore somewhat noisy, but shows about two more decades in time; data (dots) in comparison to the analytical results, Eq. (5.23b) (lines)

5.3.3 Velocity Distribution

In this section we have a look at the velocity distribution $w(\mathbf{v})$, which is the probability for a particle to have the velocity \mathbf{v} . Fundamental for molecular gases is the Maxwell distribution; however, for dissipative systems, it is well known that deviations from the Maxwellian occur. This is found in experimental work as well as simulations and for free cooling as well as driven systems. However, the actual shape of the velocity distribution is still under discussion. In most studies the highly energetic particles are overpopulated, resulting in so called “high energy tails”.

Esipov & Pöschel [1997]; Brey *et al.* [1999]; Huthmann *et al.* [2000] find that the velocity distribution for undriven dry systems decays exponentially $w(\mathbf{v}) \propto \exp(-Av)$ in the high velocity limit. For a uniformly heated system, van Noije & Ernst [1998] find that a stretched exponential $w(\mathbf{v}) \propto \exp(-Av^{3/2})$ is a solution to the homogeneous Enskog-Boltzmann equation in arbitrary dimension in the high velocity limit. Ben-Naim & Machta [2005]; Ben-Naim *et al.* [2005] additionally find stationary states with algebraic tails $w(\mathbf{v}) \propto v^{-\sigma}$, with a homogeneous model using a very general collision kernel. In a two-dimensional system, Goldhirsch *et al.* [1993] find that the velocity distribution deviates from the classical Maxwell-Boltzmann distribution as the *clustering instability*³ sets in.

Zaburdaev *et al.* [2006] studied a one-dimensional *wet* granular system (meaning a model with well defined bond breaking energy) in the limit of vanishing bond breaking distance $d_c \rightarrow d$. It seems that the velocity distribution, initially set as Maxwellian, evolves towards an exponential profile, once the particles start to cluster. In that study, however, if the kinetic energy of a pair of particles is not sufficient to break a bond, all relative kinetic energy is dissipated and their relative motion is not considered anymore. Thus, clusters have no internal degrees of freedom and therefore, for late times, different results should be expected in our system.

In this regard, the question arises, how for the presented system the fixed bond-breaking energy and the clustering of the particles, starting at the transition time t_0 , influences the velocity distribution. Fig. 5.8 shows the typical behavior of the velocity distribution. Given the anisotropic boundary conditions, we compare the velocity distributions in the direction of the *periodic boundary conditions*, $w(v_x) = \int dv_y dv_z w(\mathbf{v})$ (left), and the *hard walls*, $w(v_z) = \int dv_x dv_y w(\mathbf{v})$ (right).

In the early stage of cooling, when $t \ll t_0$, the velocity distribution is well described by a Maxwellian. At the shown time in the top panel $t \approx 0.01t_0$, each particle has had on average three collisions. In the transition regime, where $t \approx t_0$ and $T \approx \Delta E$, deviations from the Maxwellian distribution occur, characterized by a pronounced peak in the center and by high energy tails, as in most other dissipative systems. Interestingly, in the clustering regime $t \gg t_0$, the velocity distribution returns to a Maxwellian, again. Note also that $w(v_x)$ and $w(v_z)$ coincide during the investigated simulation time, which means that $w(\mathbf{v})$ remains isotropic.

³This so called clustering instability is excessively studied and produces spatial inhomogeneities of typical size $L_0 \propto (1 - \varepsilon^2)^{1/2}$, given that the system size is large enough ($L > L_0$) and the system is given enough time to evolve.

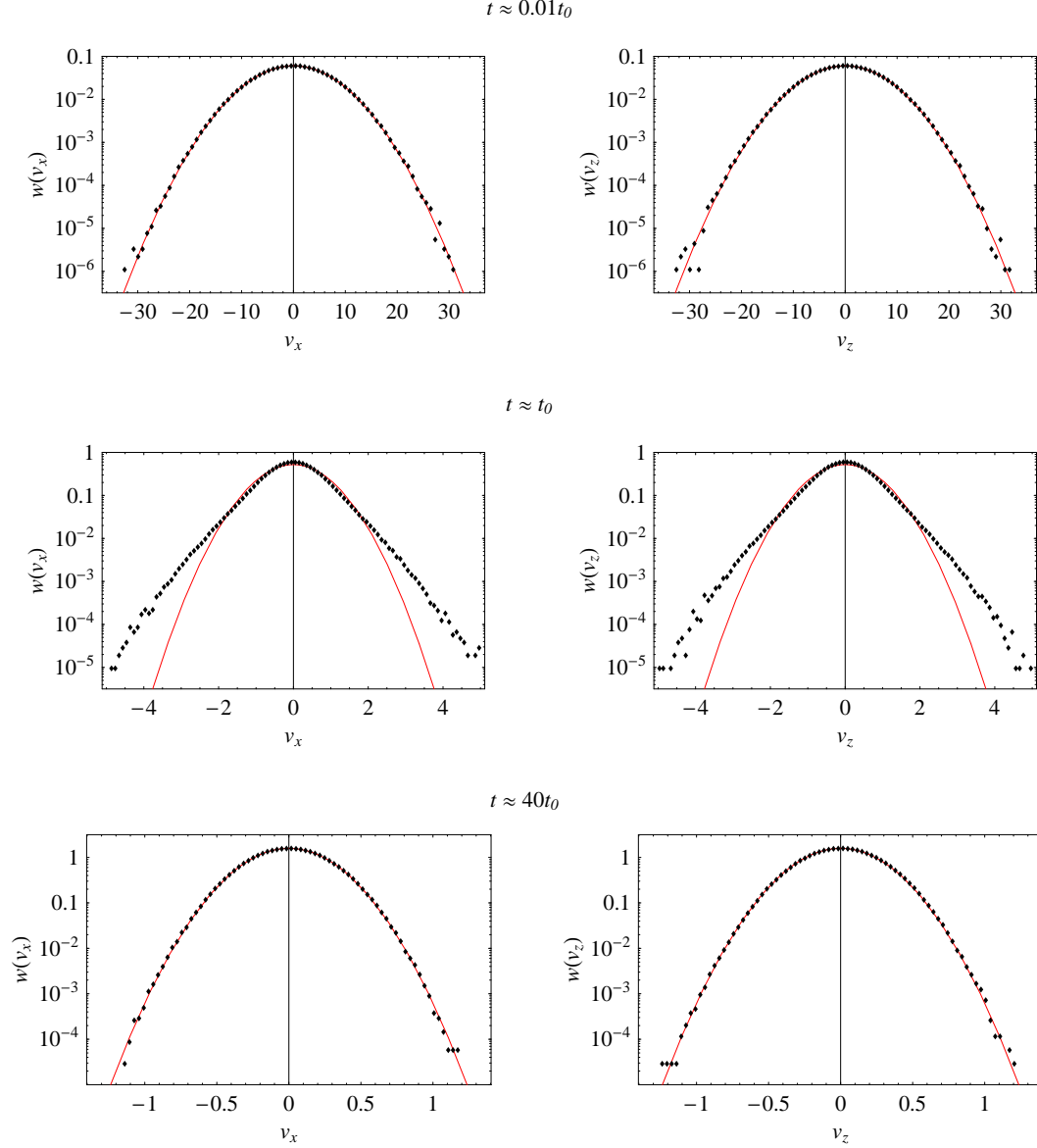


Figure 5.8: Velocity distributions $w(v_x)$ and $w(v_z)$ for times $t \approx 0.01t_0$ in the free cooling regime (top), $t \approx t_0$ in the transition state (center) and $t \approx 40t_0$ in the highly clustered state (bottom). The corresponding temperatures are $T/\Delta E = 44$ (top), 0.59 (center) and 0.063 (bottom). The data points show the distribution for a system with $N = 1048576$ particles and $\phi = 1.96\%$ and the solid lines are Maxwell distributions for the corresponding temperature at the given system time.

In order to characterize the deviations from a Gaussian profile, one typically uses the *kurtosis*, defined as the ratio of the fourth moment and the variance squared [see Balanda & MacGillivray, 1988; Chissom, 1970]:

$$\beta_2 = \frac{\langle v^4 \rangle}{\langle v^2 \rangle^2}. \quad (5.25)$$

For a Gaussian distribution, one obtains $\beta_2 = 3$; thus deviations from 3 quantify deviations from the Gaussian distribution, in particular $\beta_2 > 3$ typically corresponds to over-populated tails and peaked centers (so called *leptokurtic* distributions) and $\beta_2 < 3$ vice versa (*platykurtic* distributions). For an exponential profile $e^{-|v|/v_0}$, the kurtosis is $\beta_2 = 6$.

In our three dimensional case, every spatial dimension yields a separate value for the kurtosis. To see to what extent the anisotropy of the system plays a role, we keep the three separate values:

$$\beta_{2,\nu} = \frac{\langle v_\nu^4 \rangle}{\langle v_\nu^2 \rangle^2} \quad \text{for } \nu = x, y, z. \quad (5.26)$$

The time evolution of the kurtosis for x - and z -dimension for the system in Fig. 5.8 is shown in Fig. 5.9. One observes a sharp peak with $\beta_2 > 3$ around the time t_0 , when clustering sets in, while the rest of the evolution has the Gaussian-like value of $\beta_2 \approx 3$. The differences between x - and z -direction are marginal.

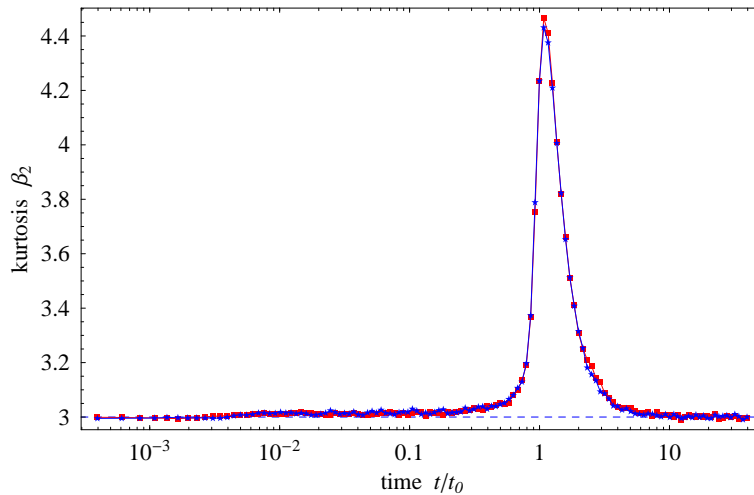


Figure 5.9: Time evolution of the kurtosis β_2 for the system of Fig. 5.8. Red boxes (■) show $\beta_{2,x}$, blue stars (★) show $\beta_{2,z}$.

The intriguing fact that the velocity distribution returns to a Maxwell distribution can be understood with the help of results of the previous section 5.3.2, where we found that bond breaking events become extremely rare in the long time limit, $P_{\text{bb}} \propto \exp(-\Delta E/T)$; consequently most collisions are elastic and the system can equilibrate.

One might consider the behavior of the particles in a cluster, which reside in “cages” determined by the gap between the hard core distance and bond breaking distance, reminiscent of glassy dynamics. Indeed, a similar model of so called “sticky hard spheres” [Baxter, 1968], where hard spheres additionally have a very short ranged attractive (but not hysteretic) square well potential, is commonly used to study glassy structures. Unlike a glass, however, it is clear that in the presented system, the rearrangements of particles do cost energy, because breakage of bonds is hysteretic; thus the long time dynamics is expected to differ from glassy systems.

Fig. 5.10 compares the time evolution of the kurtosis of different volume fractions ϕ . One can see that the value of the kurtosis around $t = t_0$ increases when going to more and more dilute systems. Since a kurtosis value of 3 is no guarantee for a Gaussian distribution, it was checked that every system indeed approaches a Maxwellian for large times and this is fulfilled very well.

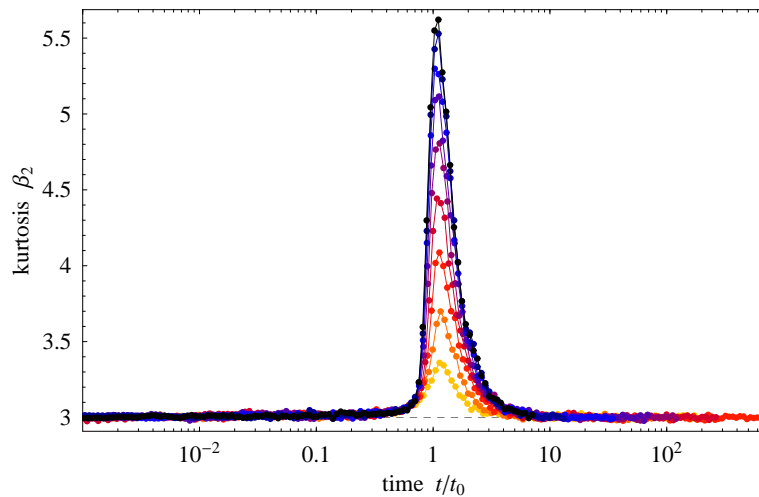


Figure 5.10: Time evolution of the kurtosis $\beta_{2,x}$ in x -direction for systems with $N = 262144$ particles and volume fractions (from bottom to top) $\phi = 15.6\%$, 7.81% , 3.90% , 1.95% , 0.98% , 0.49% , 0.24% , 0.12% , 0.061% . (same color coding as in Fig. 5.3)

5.3.4 Partitioning of the Energy into Translational, Rotational and Internal Degrees of Freedom

After the time t_0 has passed, stable clusters emerge. For the definition of a cluster, we define particles as *neighbors*, if a bridge is formed and the relative kinetic energy is not sufficient to break it. This makes sure that particles which are just “passing by”, are not considered neighbors. A *cluster* is a set of particles connected through this neighbor-relationship. We refer to the number of particles a cluster contains as the cluster mass m . One should keep in mind that clusters defined in this way are not truly stable. Particles belonging to the cluster are occasionally kicked out, if hit by a very energetic particle.

For a more detailed understanding of the system, we investigate the cooling dynamics on the cluster level, and determine how energy is partitioned among the degrees of freedom. We split the total temperature T into three constituents, the *translational* temperature defined via the center-of-mass velocities of the clusters, the *rotational* temperature defined via the angular momenta of the clusters, and the *internal* temperature describing the relative movement of the particles inside a cluster. These three temperatures are defined as follows.

Our definition of neighborhood relations gives rise to n_{cl} distinct clusters numbered by $i = 1, \dots, n_{\text{cl}}$. We denote by \mathcal{N}_i the i -th cluster with m_i particles. Its center of mass position and velocity are given by:

$$\mathbf{R}_i = \frac{1}{m_i} \sum_{\nu \in \mathcal{N}_i} \mathbf{r}_\nu \quad \text{and} \quad \mathbf{V}_i = \frac{1}{m_i} \sum_{\nu \in \mathcal{N}_i} \mathbf{v}_\nu. \quad (5.27)$$

Note that single particles with $m_i = 1$ are also considered as clusters.

The center of mass movement of each cluster has $f_{\text{trans},i} = 3$ translational degrees of freedom, so that the total number of translational degrees of freedom of these clusters is simply $3n_{\text{cl}}$. Homogeneous cluster translations are thus characterized by the *translational* temperature

$$T_{\text{trans}} := \frac{2}{3n_{\text{cl}}} \sum_{i=1}^{n_{\text{cl}}} \frac{m_i}{2} \mathbf{V}_i^2. \quad (5.28)$$

Analogously, the *rotational* temperature describes the energy in homogeneous cluster rotations. The angular momentum, \mathbf{L}_i , of cluster i is given in terms of the relative particle positions $\tilde{\mathbf{r}}_{i,\nu} = \mathbf{r}_\nu - \mathbf{R}_i$ and velocities $\tilde{\mathbf{v}}_{i,\nu} = \mathbf{v}_\nu - \mathbf{V}_i$

$$\mathbf{L}_i = \sum_{\nu \in \mathcal{N}_i} \tilde{\mathbf{r}}_{i,\nu} \times \tilde{\mathbf{v}}_{i,\nu}. \quad (5.29)$$

The rotational energy of cluster \mathcal{N}_i with $m_i > 2$ is thus given by

$$E_{\text{rot},i} = \frac{1}{2} \mathbf{L}_i \underline{\mathbf{I}}_i^{-1} \mathbf{L}_i, \quad (5.30)$$

where the moment of inertia tensor $\underline{\mathbf{I}}_i$ is defined in the usual way⁴. The case $m_i = 2$, requires special treatment, since the inertia tensor is singular. The rotational energy of a dimer can be easily calculated to $E_{\text{rot},i} = \Delta \mathbf{v}_{i\perp}^2 / 4$, where $\Delta \mathbf{v}_{i\perp}$ denotes the relative velocity perpendicular to the axis of the dimer. The *rotational* temperature is thus

$$T_{\text{rot}} := \frac{2}{\sum_{i=1}^{n_{\text{cl}}} f_{\text{rot},i}} \sum_{i=1}^{n_{\text{cl}}} E_{\text{rot},i}, \quad (5.31)$$

with $f_{\text{rot},i} = 2$ for dimers and $f_{\text{rot},i} = 3$ for larger clusters.

All the left-over kinetic energy E_{int} describes the relative movement of particles inside a cluster and contributes to the *internal* temperature. Each cluster has a

⁴ $I_{i,\mu\nu} = \sum_{j \in \mathcal{N}_i} m_j (\delta_{\mu\nu} r_j^2 - r_{j,\mu} r_{j,\nu})$.

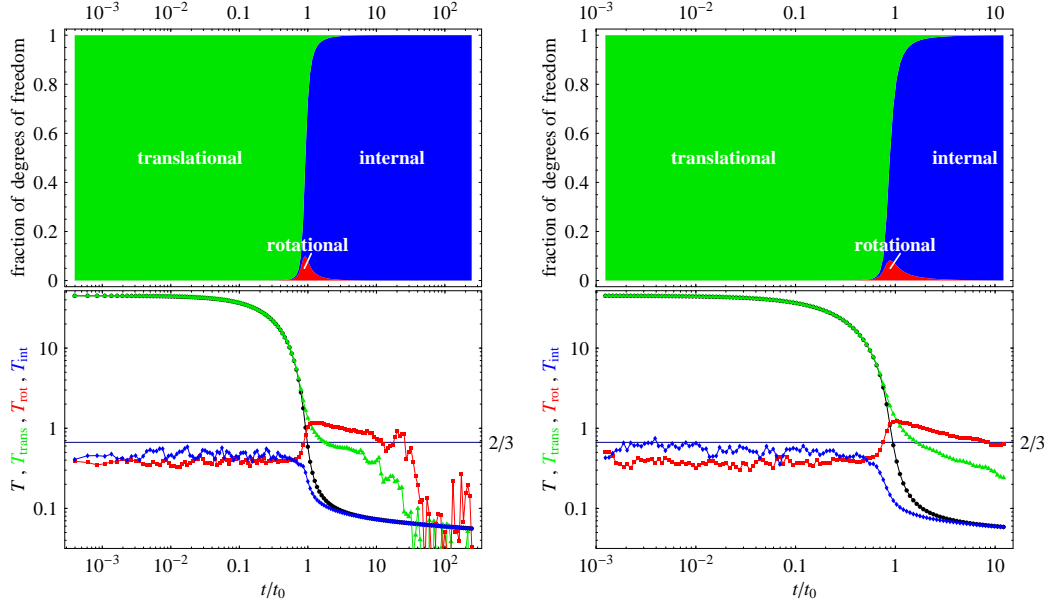


Figure 5.11: Top: Division of the total $3N$ degrees of freedom into the *translational*, *rotational* and *internal* parts, dependent on time. Bottom: Evolution of the *total* (\bullet), *translational* (\blacktriangle), *rotational* (\blacksquare), and *internal* (\blacklozenge) granular temperatures. Data for $N = 262144$ particles and volume fraction $\phi = 1.95\%$ (left) and $\phi = 0.12\%$ (right) are shown; the behavior is qualitatively the same for all investigated system sizes. The horizontal line at $2/3$ corresponds to the bond breaking energy.

total of $3m_i$ degrees of freedom, so that the remaining number for internal degrees of freedom is $f_{\text{int},i} = 3m_i - f_{\text{trans},i} - f_{\text{rot},i}$. The internal temperature T_{int} is:

$$T_{\text{int}} := \frac{2}{\sum_{i=1}^{n_{\text{cl}}} f_{\text{int},i}} \sum_{i=1}^{n_{\text{cl}}} E_{\text{int},i}. \quad (5.32)$$

Fig. 5.11 (top) shows how the total of $3N$ degrees of freedom divide up into *translational*, *rotational*, and *internal* degrees of freedom. The corresponding temperatures are shown in the lower half of the figure. As one might expect, for $t \ll t_0$ almost all degrees of freedom are *translational*, since most clusters are just single particles, and $T_{\text{trans}} \approx T$. Keeping in mind that two particles are only defined as neighbors if their relative velocity is not sufficient to break the bond, only *stable* clusters (mostly dimers) enter the *internal* and *rotational* temperatures, and therefore $T_{\text{rot}}, T_{\text{int}} < \frac{2}{3}\Delta E = \frac{2}{3}$ for $t/t_0 < 1$.⁵

In the transitional regime $t \approx t_0$, when the number of intermediate size clusters increases, the rotational degrees of freedom become important. Larger objects can have higher rotational energies without rupture⁶, therefore the growing clusters

⁵The factor $\frac{2}{3}$ is due to the relation $\frac{2}{3}\bar{E}_{\text{kin}} = T$ between temperature and energy.

⁶Roughly speaking, the maximum rotational energy $E_{\text{rot,max}}$ of a cluster with radius r and mass M is $E_{\text{rot,max}} \sim Mr^2\omega^2$, where the maximum rotational frequency ω is limited by the centrifugal

obtain energy from caught particles, and thus T_{rot} increases until reaching the value of T_{trans} . After that, the energy of the incoming lumps is not sufficient to increase T_{rot} any further.

In contrast to the homogeneous cluster *rotations*, the *internal* degrees of freedom which have higher energies than ΔE will in most cases result in a bond rupture, independent of the cluster size. Therefore, T_{int} decreases monotonically. At late times $t \gg t_0$, large clusters have formed, thus almost all degrees of freedom are *internal* and $T \approx T_{\text{int}}$.

5.4 Aggregation

When the average kinetic energy per particle is comparable to the bond breaking energy, $t \approx t_0$, the system starts to form aggregates, which seem to grow in a self-similar process. In the following we are going to analyze these aggregates and compare them to cluster-cluster aggregation models [Jullien & Botet, 1987]. As time proceeds, larger and larger clusters are formed. A spanning or percolating cluster is observed for all finite densities (provided the system size L is large enough), and ultimately all particles and clusters have merged into a single cluster.

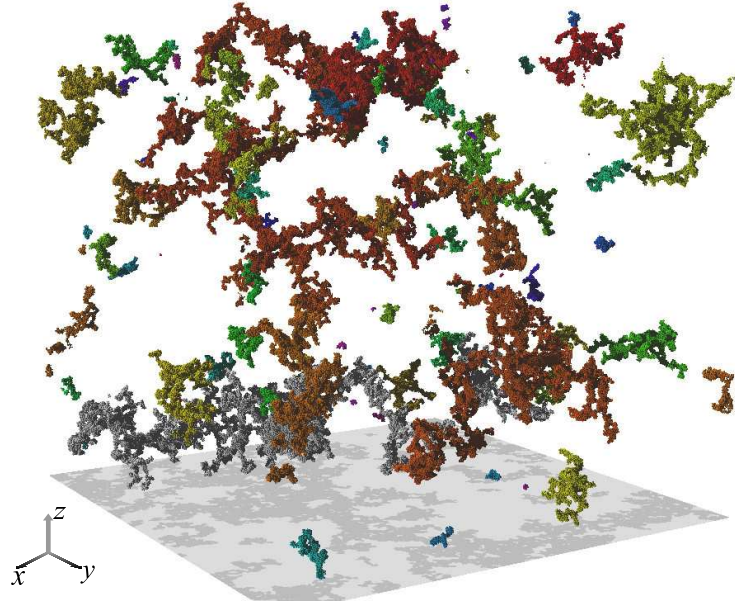


Figure 5.12: Snapshot of the system with volume fraction $\phi = 0.48\%$ and $N = 262\,144$ particles taken at time $t \approx 12t_0$; the largest cluster (gray) contains 22% of the particles. Particles of the same cluster have the same color shade.

force $F \sim M\omega^2 r$. This yields $E_{\text{rot,max}} \sim r$. In our case the bond breaking energy ΔE is related to the maximum force F on the particles by $\Delta E \sim F \cdot (d_c - d)$, with the freely movable distance of a particle ($d_c - d$).

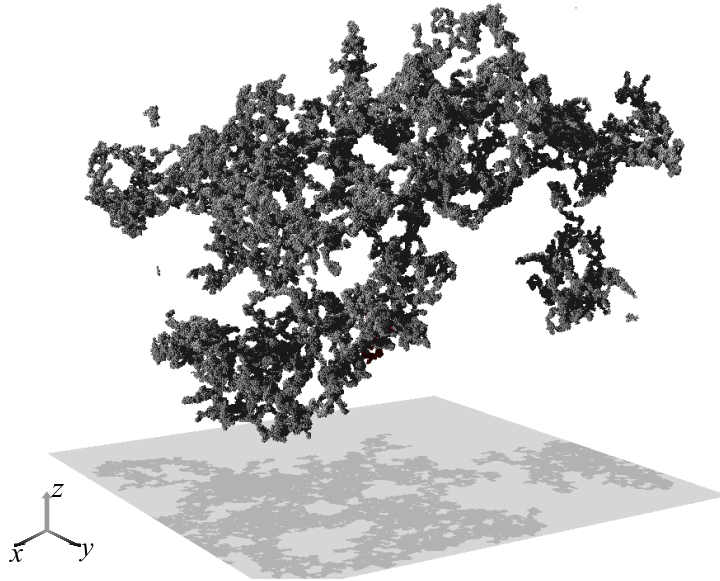


Figure 5.13: Same as Fig. 5.12 for $t \approx 52t_0$; the largest cluster contains 99% of the particles.

Figs. 5.12 and 5.13 show snapshots of a system at $t = 12t_0$ and $t = 52t_0$ with small volume fraction, $\phi = 0.48\%$. At the smaller time the system is not yet percolating, even though rather large clusters have already formed, the largest one (in gray) contains 22% of all particles. The second snapshot, taken at a much longer time, shows a spanning cluster. At such large times the average kinetic energy is much smaller than the bond breaking energy ($T \approx 0.06\Delta E$), so that bonds almost never break up. The cluster shown is already well beyond the critical time for percolation with 99% of the particles in the cluster.

Fig. 5.14 shows the evolution of the cluster mass distribution $N_m(t)$, which is the number of clusters containing m particles at time t . One can clearly see that after some time, $t \approx 2.5t_0$, which depends on volume fraction, the largest cluster emerges from the rest of the distribution. For all volume fractions a gelation transition was observed at the percolation time $t_c > t_0$. The critical behavior of the gelation transition is still controversial. Since aggregation is a *nonequilibrium* process, there is *a priori* no reason that it should be in the same universality class as the corresponding equilibrium percolation transition. Yet there is some evidence in favor of this conjecture. Gimel *et al.* [1995] observe a cross-over from self-similar growth at small times and volume fractions – called the flocculation regime – to the percolation regime around t_c . In the latter they observe critical exponents⁷ as in standard percolation theory. Kolb & Herrmann [1985] on the other hand obtain values for the fractal dimension of the percolating cluster, distinct from percolation

⁷like fractal dimension, or scaling of the mean cluster mass with the distance from the critical concentration

theory as well as from flocculation theory. Both studies refer to diffusion limited cluster-cluster aggregation.

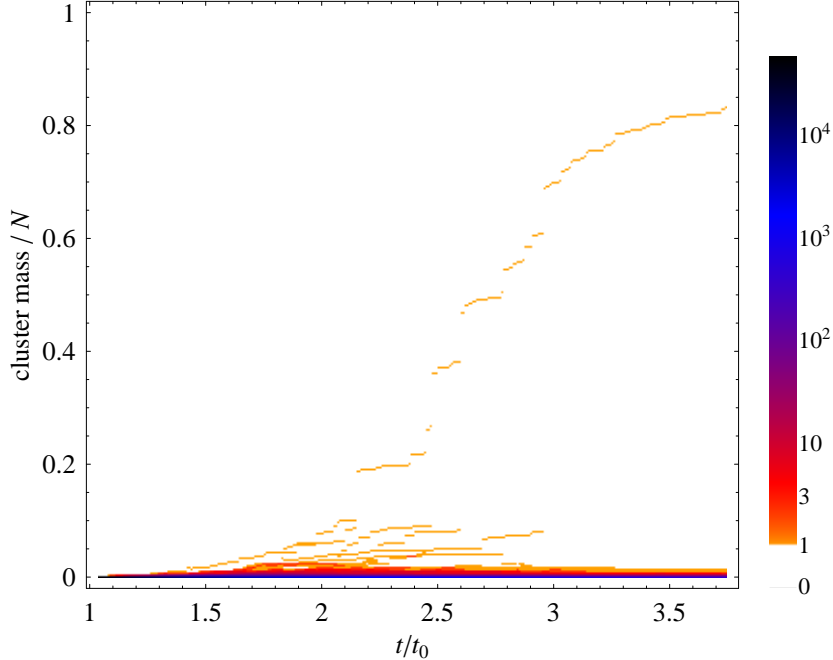


Figure 5.14: Histogram of the cluster mass distribution dependent on time, for volume fraction $\phi = 3.9\%$ and $N = 262144$. The number of clusters at the respective time and size is color coded on a logarithmic scale so that the single largest cluster is visible. At $t \approx 2.5t_0$ one can see the large cluster emerging, clearly distinguishable from the rest of the distribution.

Here we do not analyze the gelation transition in detail, however, work along these lines is in progress. Instead we investigate two regimes in detail in the following:

- a) The **self-similar growth process**, or flocculation regime, which is present for small times and volume fractions.
- b) The **properties of the final cluster** which emerges, when (almost) all particles have aggregated to form one large cluster.

5.4.1 Self-Similar Growth

5.4.1.1 Fractal Dimension of the Aggregates

A central quantity of aggregation models is the fractal dimension of the aggregates. It is usually determined from the radius of gyration as a function of cluster mass. We consider a cluster of m particles with positions $(\mathbf{r}_1, \dots, \mathbf{r}_m)$ and define its radius

of gyration by [see e.g. Stanley & Ostrowsky, 1986]

$$r_g^2(m) = \frac{1}{m} \sum_{i=1}^m (\mathbf{r}_i - \bar{\mathbf{r}})^2 \quad \text{with} \quad \bar{\mathbf{r}} = \frac{1}{m} \sum_{i=1}^m \mathbf{r}_i. \quad (5.33)$$

If the clusters are fractal we expect a scaling relation for large m of the form

$$r_g \sim m^{1/D_f}, \quad (5.34)$$

which yields the fractal dimension D_f . This method is commonly used in aggregation models, where particles move diffusively, ballistically, or are interacting and stick to the aggregate once they touch it [Jullien & Kolb, 1984; Meakin, 1991; Westbrook *et al.*, 2004; Alves & Ferreira, 2006].

Fig. 5.15 shows the radius of gyration for a system of 262144 particles at volume fraction $\phi = 1.95\%$. Several snapshots of the ensemble of growing clusters have been taken at times $t_0 < t < t_c$ with the percolation time t_c , when a spanning cluster is first observed. The data scale well according to Eq. (5.34), some scatter is observed for the largest masses, corresponding to times close to the percolation transition.

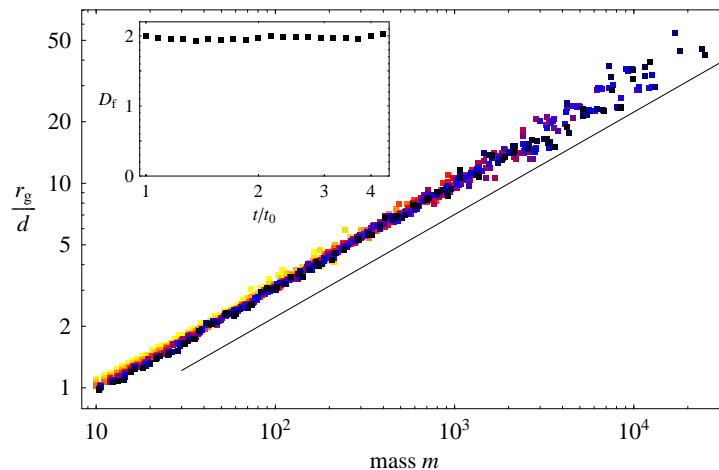


Figure 5.15: Radius of gyration as a function of cluster size for a system of 262144 particles at volume fraction $\phi = 1.95\%$; different colors/shades correspond to simulation times between t_0 (yellow) and $4t_0 < t_c$ (black)); The slope of the solid line corresponds to $D_f = 2$. Inset: fractal dimension as a function of time, extracted from the slope of the curves in the main figure.

In contrast to aggregation models, where the clusters are static and do not break up, we occasionally do observe the breaking of bonds. In addition there are internal deformations of the clusters during growth, so that the fractal dimension could depend on time. Therefore the relation between m and $r_g(m)$ is checked for many instances of time and the resulting fractal dimension is shown as a function of time in the inset of Fig. 5.15. As can be seen from the Figure, there is no systematic dependence on time, and the fractal dimension is close to $D_f = 2$.

5.4.1.2 Cluster Mass Distribution

All information about the connectivity of the clusters is contained in the cluster mass distribution $N_m(t)$, the number of clusters of size m at time t . Fig. 5.16 shows $N_m(t)$ for a system with $\phi = 1.96\%$ and $N = 1048576$. The time interval has been chosen such that $t_0 < t < 2t_0 < t_c \approx 4t_0$ (for this volume fraction). In this time interval the mean cluster mass increases roughly by a factor of 30.

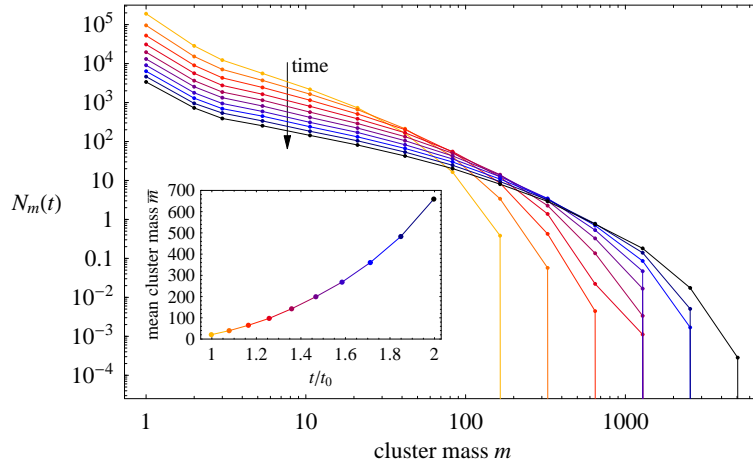


Figure 5.16: The cluster mass distribution $N_m(t)$. The different graphs represent different times, which are increasing from top to bottom (left side of the graph). The inset shows how the mean cluster mass increases during the investigated time period, where the colors correspond to the times of the main plot.

It has been suggested [e.g. Meakin, 1991] that for aggregating systems the mass distribution evolves towards a self-preserving scaling form, independent of the initial distribution:

$$N_m(t) = m^{-\theta} f(m/\bar{m}(t)), \quad (5.35)$$

where the time dependence is only contained in the mean cluster mass

$$\bar{m}(t) = \frac{\sum_{m=1}^{\infty} m^2 N_m(t)}{\sum_{m=1}^{\infty} m N_m(t)}. \quad (5.36)$$

This scaling form has been applied successfully to various aggregating systems [Vicsek & Family, 1984; Botet & Jullien, 1984; Jiang & Leyvraz, 1993, 1994; van Dongen & Ernst, 1985; Meakin, 1991; Trizac & Hansen, 1995], involving fractal as well as non-fractal objects. Mass conservation requires $\theta = 2$ [Meakin, 1991].

In Fig. 5.17 the scaling function $f(m/\bar{m}) = N_m(t)m^2$ is plotted for the same data sets as in Fig. 5.16. We expect scaling to hold only in the aggregation regime, *i.e.* for times not too close to t_c , where the system gels (see next Sec. 5.4.1.3); that is why we restrict ourselves to times $t_0 < t < 2t_0$. Also the data points for $m = 1$ have been left out, *i.e.* clusters consisting of single particles. As can be seen from

Fig. 5.17 the data scale very well. Deviations occur only for times close to the percolation transition (not shown here), where they should be expected.

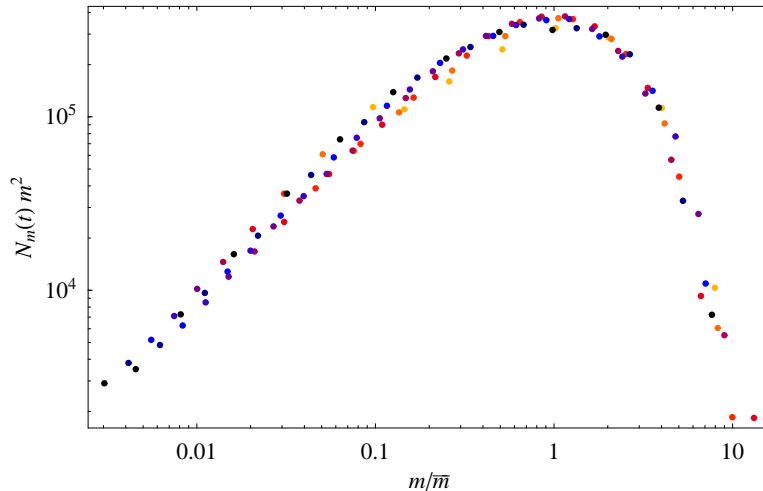


Figure 5.17: Rescaled cluster mass distribution $f(m/\bar{m}) = N_m(t) \cdot m^2$ from Eq. (5.35) versus the normalized cluster mass m/\bar{m} . The color coding as in Fig. 5.16 is used.

5.4.1.3 Number of Clusters and Mean Cluster Mass

Another characteristic of a realization of clusters is simply the total number of clusters $n_{\text{cl}}(t) = \sum_{m=1}^{\infty} N_m(t)$, which decreases as aggregation proceeds. As long as the system is in the scaling regime (*i.e.* relation (5.35) is fulfilled), the mean cluster mass, $\bar{m}(t)$ and the number of clusters are simply related: $\bar{m}(t) \sim n_{\text{cl}}^{-1}$. However, as mentioned above, the scaling relation (5.35) only holds in the aggregation regime and is expected to break down as the percolation transition is approached. At that point, \bar{m} should diverge due to the formation of a spanning cluster. On the other hand, there is still a large number of smaller clusters coexisting with the macroscopic cluster, so that n_{cl}/N remains finite at the percolation transition.

The aggregation of particles to larger objects has been investigated for various *ballistic aggregation* models [Family & Vicsek, 1985; Carnevale *et al.*, 1990; Jiang & Leyvraz, 1994; Trizac & Krapivsky, 2003], where spherical particles of mass $m = 1$ and diameter $d = d_0$ move ballistically, until two of them collide to form clusters irreversibly. In a particularly simple model, one assumes that two colliding particles form one larger spherical particle with conserved momentum and a mass m equal to the sum of the two particles masses, so that m is always equal to the number of *initial particles* contained in a given cluster. For spatial dimension D , the diameter increases like $d = m^{1/D} d_0$, assuming the particles to be compact spheres which conserve volume when merging. For this model, a mean field theory [Jiang & Leyvraz, 1994] and simple scaling arguments [Carnevale *et al.*, 1990; Trizac & Krapivsky, 2003] yield the dependence of the expected average mass \bar{m} on time like $\bar{m} \sim t^\xi$

with an exponent $\xi = 2D/(D + 2)$ (assuming $t_0 = 0$).

Since the aggregating clusters in our system are not compact, but fractal objects with fractal dimension D_f , the assumption for the diameter $d \sim m^{1/D}$ does not hold and must be changed to $d \sim m^{1/D_f}$. With this assumption, we follow the scaling arguments of [Trizac & Krapivsky \[2003\]](#), and find the scaling relation between \bar{m} and t .

We assume that the number of clusters per volume, n_{cl} , is reduced by one whenever two clusters collide:

$$\frac{dn_{\text{cl}}}{dt} \sim -f_{\text{coll}} \cdot n_{\text{cl}}. \quad (5.37)$$

The collision frequency [[Brilliantov & Pöschel, 2004](#)] is approximately given by $f_{\text{coll}} \sim d^{D-1} n_{\text{cl}} v$ with $d \propto r_g$ the linear dimension of the cluster and v its typical velocity. The average momentum should scale as $p \sim m^{1/2}$ [[Trizac & Krapivsky, 2003](#)], and therefore

$$v = p/m \sim m^{-1/2} \sim n_{\text{cl}}^{1/2}. \quad (5.38)$$

Plugging in all these scaling relations as well as $m \sim r_g^{D_f}$, one obtains:

$$\frac{dn_{\text{cl}}}{dt} \sim -n_{\text{cl}}^2 \cdot v \cdot d^{D-1} \sim -n_{\text{cl}}^{5/2-(D-1)/D_f}, \quad (5.39)$$

which is solved by

$$n_{\text{cl}} \sim (t - t^*)^{-2D_f/(3D_f-2D+2)}, \quad (5.40)$$

where the integration constant t^* is the onset of cluster growth. In our context $t^* \approx t_0$.⁸ This implies the following growth law for the mean cluster mass in the scaling regime:

$$\bar{m} \sim (t - t^*)^\xi \quad \text{with} \quad \xi = \frac{2D_f}{3D_f - 2D + 2}, \quad (5.41)$$

which generalizes the result for compact objects, $\xi = 2D/(D + 2)$ with $D = D_f$ to fractal ones with $D \neq D_f$.

Fig. 5.18 shows how the number of clusters decreases over time as larger and larger aggregates form for $t > t_0$. The inset of Fig. 5.18 investigates the scaling behavior (5.40), with the origin of the time axis shifted to the transition point t^* . One can see that the slope of $\xi = 2$, obtained from (5.41) for $D = 3$ and $D_f = 2$ is in good agreement with the simulation.

5.4.2 Properties of the Asymptotic Cluster

The largest cluster – well beyond the percolation transition for most volume fractions – will be the main focus of this section. In particular we determine its fractal dimensions and local structure. For the fractal dimension, there are various definitions, which in general may yield different results and three of which are presented. Particularly illuminative is the box counting algorithm: simple considerations make a prediction about a cross-over length scale at which the cluster structure changes from fractal to compact behavior, which is indeed observed in the simulation data.

⁸As one can see in the main plot of Fig. 5.18, the actual onset of cluster growth is not exactly at t_0 , but a little bit earlier.

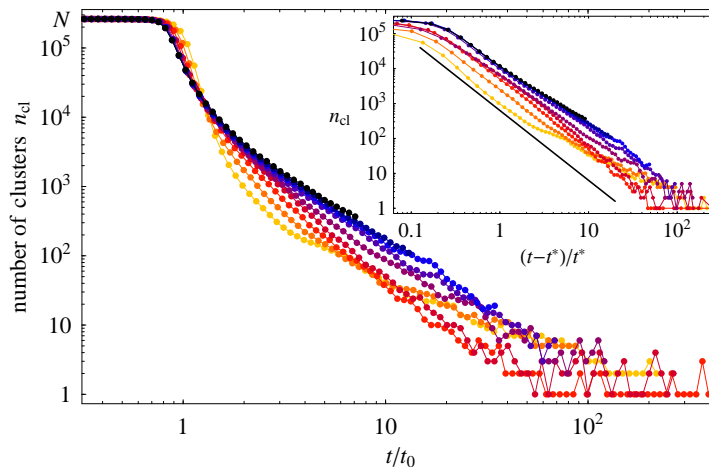


Figure 5.18: Evolution of the mean cluster mass. Labeling and parameters as in Fig. 5.3. For the inset, the origin of the time-axis has been shifted to the transition point t^* to investigate the scaling relation $n_{cl} \sim (t - t^*)^{-\xi}$. The solid line has a slope of -2 .

5.4.2.1 Fractal Dimension from Radius of Gyration

One way to determine the fractal dimension is the radius of gyration, as was done in Sec. 5.4.1.1 for aggregates. Here, however, we only have one large cluster and have to find a way to obtain the function $r_g(m)$ as a function of cluster size m . This is implemented in following way: Starting from a random particle of the cluster, we mark all particles that can be reached through i or less neighbor-to-neighbor steps (see Fig. 5.19). Thus, for every i , we get a *partial cluster* with $m(i)$ particles and radius of gyration $r_g(i)$, which yields the scaling relation $r_g \sim m^{1/D_f}$ and the fractal dimension D_f . For good statistics, this procedure is repeated 100 times (each with a different initial particle) and the obtained values of r_g are averaged. Note furthermore that the procedure takes care that no particle is marked a second time, in order to make sure that one does not go through the cluster several times because of the periodic boundary conditions.

Fig. 5.20 shows the results of this procedure for the radius of gyration r_g as a function of m for different densities. For high volume fractions we are well beyond the percolation transition and hence expect $D_f = 3$ on the *largest* length scales of the cluster. This is clearly seen in Fig. 5.20, e.g. for $\phi = 15.6\%$ and $10^3 < m < 10^5$. On smaller length scales, however, we find a fractal dimension $D_f \approx 2$. For smaller volume fractions, the cross-over to $D_f = 3$ happens at larger masses and hence the fractal behavior extends to larger scales. In the following section we will see, why there must be a cross-over from fractal to compact behavior and how the volume fraction determines the cross-over point.

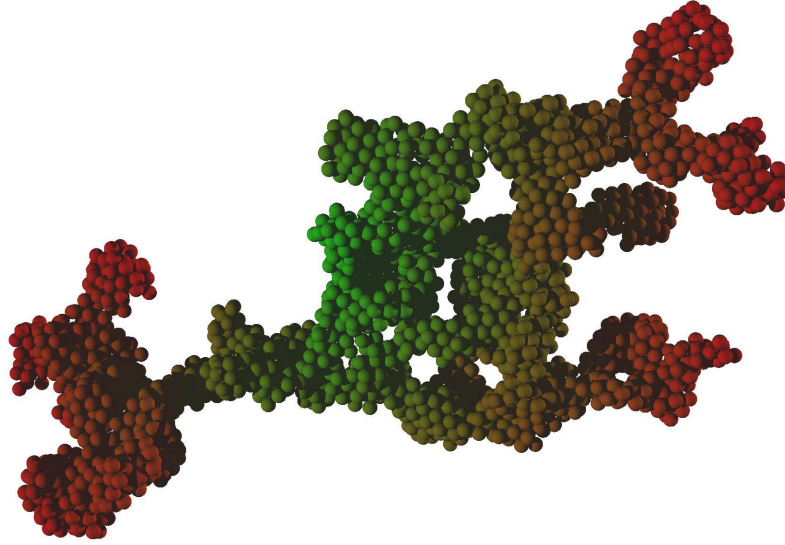


Figure 5.19: Illustration of the procedure to obtain *partial clusters*. The initial particle is marked in pure green. The number of neighbor-to-neighbor steps i necessary to reach another particle is indicated by the color shade of the particles: Larger i corresponds to a gradual shift towards red. The outermost particles (fully red) correspond to $i = 36$ steps.

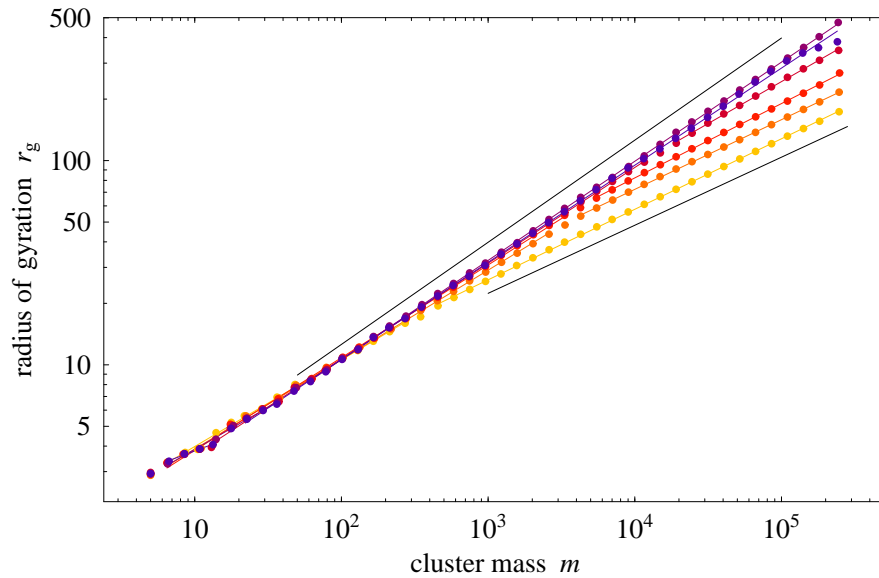


Figure 5.20: Radius of gyration dependent on the mass of the partial cluster at simulation time $t \approx 27t_0$. The particle number is fixed $N = 262144$ and the volume fractions are (from bottom to top) $\phi = 15.6\%$, 7.81% , 3.90% , 1.95% , 0.98% , 0.49% . The lines along the data points are the respective fits. The outer solid lines have slopes $1/2$ (top) and $1/3$ (bottom) corresponding to fractal dimensions of 2 and 3, respectively.

5.4.2.2 Fractal Dimension from Box Counting Algorithm

To further investigate the fractal dimension of the largest cluster at *intermediate* length scales, we use the box counting algorithm [Grassberger, 1983; Hentschel & Procaccia, 1983]. The system is divided into sub-boxes of edge length L_{box} . Then each box which contains or hits at least one particle is marked. In this way, we find the number of boxes N_{box} necessary to cover the whole cluster. This number should scale with L_{box} like

$$N_{\text{box}} \sim L_{\text{box}}^{-D_f}, \quad (5.42)$$

with the Hausdorff dimension D_f .

On length scales much smaller than the particle diameter, $L_{\text{box}} \ll d$, the system obviously behaves three-dimensionally. In this regime, the number of filled boxes N_{box} is just the volume fraction ϕ times the total number of boxes $N_{\text{box,tot}} = L^3/L_{\text{box}}^3$, therefore:

$$N_{\text{box}} = \frac{\phi L^3}{L_{\text{box}}^3}. \quad (5.43)$$

Since our system is finite and contains a system-spanning cluster, the scaling behavior on large length scales $L_{\text{box}} \approx L$ should also be three dimensional. On this length scale, almost all the boxes should be filled, so that

$$N_{\text{box}} = \frac{L^3}{L_{\text{box}}^3}. \quad (5.44)$$

In particular, the relation must include the point $(L_{\text{box}}, N_{\text{box}}) = (L, 1)$, since a box of the system size includes all particles and will certainly be marked.

Only in the regime between these two limiting cases is it possible to observe the fractal dimension with the box-counting method. A schematic plot is given on the left side of Fig. 5.21, where the number N_{box} of boxes containing particles is plotted against the edge length L_{box} of a box. Comparing (5.43) and (5.44) shows that the interesting range is proportional to $|\log \phi|$, which only depends on the volume fraction, but not on the particular choice of the system size.

For numerical reasons, it is very tedious to observe the expected slope of -3 for small L_{box} , because of the vast amount of boxes to account for. Since this regime is not relevant anyway, it has only been investigated exemplarily and is reached for $L_{\text{box}} \lesssim 0.03d$ (see right side of Fig. 5.21). For all other runs we simplify the algorithm and only use the *centers* of the particles, *i.e.* a box is only marked, if a particle center is inside. With this definition the number of boxes needed to cover the system for small box sizes $L_{\text{box}} \lesssim d$ is just the particle number N , resulting in a horizontal line on the left side of the graph (dotted line in Fig. 5.21), instead of the slope -3 .

Fig. 5.22 (top) shows the outcome of the box-counting algorithm, at a time $t \approx 27t_0$, where roughly all particles are inside the largest cluster. It yields the relation between the box size L_{box} and the number of boxes of that size, needed to

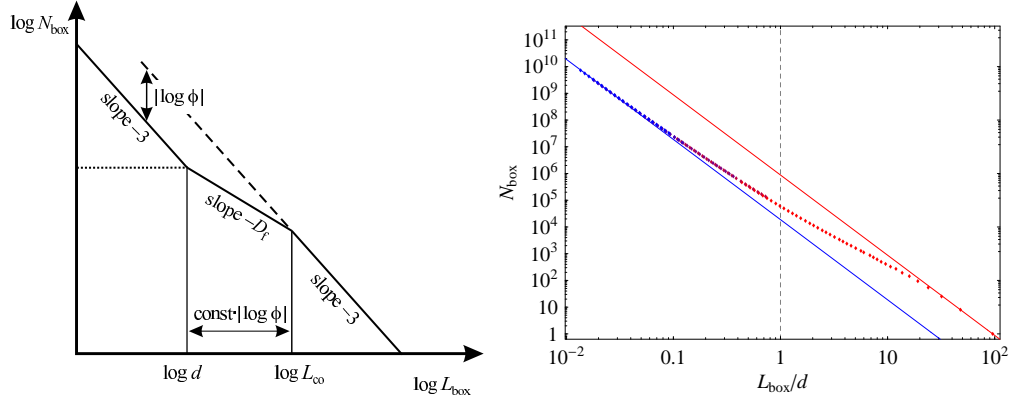


Figure 5.21: Left: Schematic double logarithmic plot of the box size L_{box} versus the number of boxes N_{box} of that size needed to cover the cluster. The negative slope is the fractal dimension. We expect three scaling regions: For small and large L_{box} , the system should behave three dimensionally, and the region in between yields the non-trivial fractal dimension. If only the particle centers are considered, the algorithm simply counts the number of particles in the cluster for $L_{\text{box}} \lesssim d$ resulting in a horizontal line (dotted line). Right: Exemplary application of the box counting algorithm, considering the extent of the spheres. Only a small system ($N = 32768$ particles and $\phi = 1.95\%$) was used and on small length scales, only fractions of the system were investigated and the resulting data was rescaled: the different colors of the data points correspond to cubic sections of the whole system with edge length $L/30$, $L/24$, $L/8$, $L/4$, and L (whole system). The solid lines are fits to the data points on the largest and smallest length scales. As proposed, their vertical distance of 1.67 roughly equals $|\log_{10} \phi| = 1.71$.

cover the cluster. The slope of that curve is the negative fractal dimension. The result for different system sizes, but with the same volume fraction are presented. As proposed, for all system sizes, there is a cross-over point L_{co} , at which the slope changes. On length scales between d and L_{co} , the fractal dimension is roughly 2 (the fits yield values between 1.92 and 2.03). Above L_{co} the fractal dimension has a trivial value of about 3 (fit values between 2.95 and 3.00), which means that on these large length scales all the boxes are filled and is therefore an indication that the cluster is system-spanning.

In the lower half of Fig. 5.22 we see the number of boxes normalized by the cluster mass. The data collapse well onto a single curve, obviously with the same slopes. Here one can see very well that for systems with the same volume fraction, the slopes as well as the cross-over point do not depend on the absolute system size.

Results of the box counting algorithm for *different densities* are presented in Fig. 5.23 (top). Only densities for which a spanning cluster has developed are included. The *slopes* of -2 and -3 in the two scaling regions are not affected by the volume fraction ϕ , but the size of the non-trivial region (with $D_f \approx 2$) is seen to increase significantly as the density decreases. Even for the lowest density, the

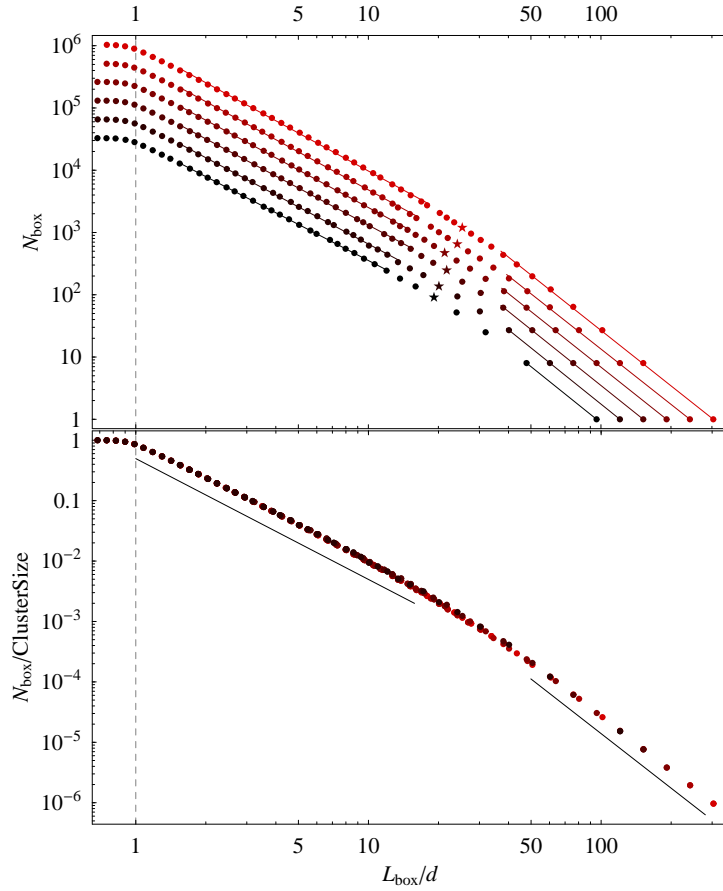


Figure 5.22: Top: N_{box} versus L_{box} at time $t \approx 27t_0$ and volume fraction $\phi = 1.95\%$ for the box counting algorithm; particle number is varied from bottom to top, according to $N = 32768, 65536, 131072, 262144, 524288, 1048576$. The straight lines are fits to the data to the left and right of the cross-over point $L_{\text{co}} \approx 25d$, which is also a fitting parameter and shown as a star (\star). Bottom: As top, but N_{box} normalized by cluster mass; the solid lines have slopes -2 (left) and -3 (right); the vertical dashed line represents the particle size.

size of the scaling region is less than 2 decades, which makes it difficult to extract precise values for the fractal dimension. For the three most dense systems, the scaling region is less than one decade. As discussed earlier, this is an intrinsic feature of the “high” density systems, which can not be resolved by taking larger systems ($N, L \rightarrow \infty$ with constant ϕ). As the bottom panel of Fig. 5.23 shows, we can collapse all data on a single curve by rescaling N_{box} with ϕ^{-2} and L_{box} with ϕ in agreement with the dependence of the cross-over length on $|\log \phi|$.

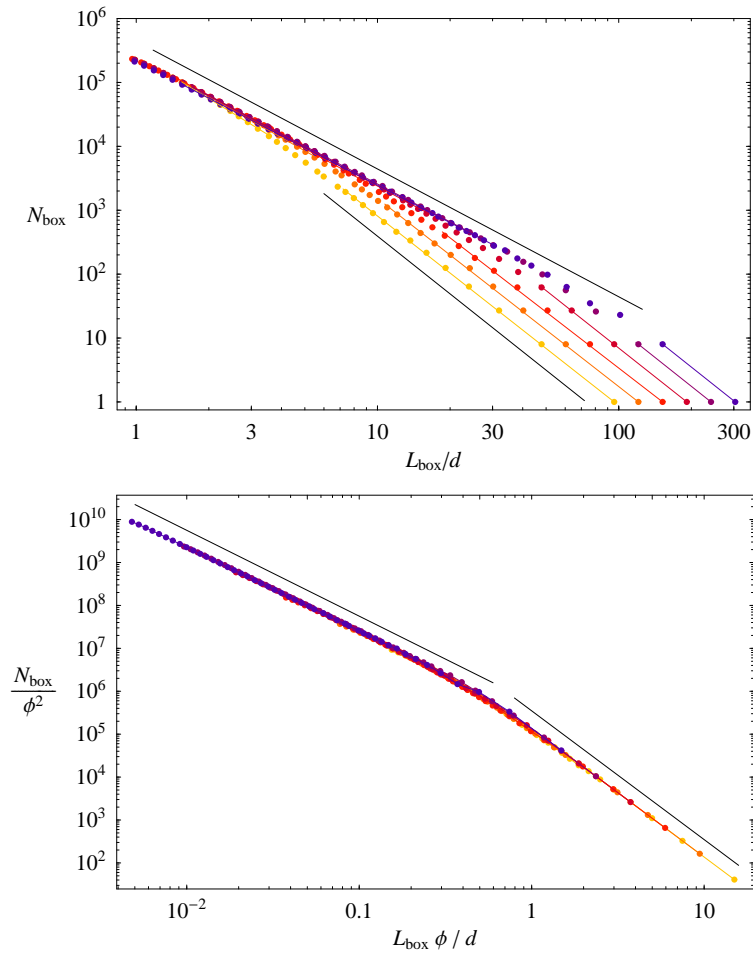


Figure 5.23: Top: Result of the box counting algorithm, as in Fig. 5.22 at $t \approx 45t_0$; the particle number is fixed $N = 262144$ and the volume fraction varies from left to right according to $\phi = 15.6\%$, 7.81% , 3.90% , 1.95% , 0.98% , 0.49% ; the lines along the data points are the respective fits; the outer lines have slope -2 (upper line) and -3 (lower line). Bottom: As top, but with rescaled axes.

5.4.2.3 Pair Correlation Function and Correlation Dimension

A fundamental quantity for granular (or atomic) systems is the **pair correlation function** $g(\mathbf{r})$ which specifies the probability that the centers of two particles are separated by a vector \mathbf{r} [Hansen & McDonald, 1986]:

$$g(\mathbf{r}) = \frac{1}{n} \left\langle \frac{1}{N} \sum_{i \neq j} \delta(\mathbf{r}_i - \mathbf{r}_j - \mathbf{r}) \right\rangle. \quad (5.45)$$

Since the interactions of the particles do not give rise to anisotropies, one typically only considers the *radial* component, i.e. the **radial distribution function (RDF)** $g(r)$; specifically, it is the average of $g(\mathbf{r})$ over all directions of \mathbf{r} but with fixed $|\mathbf{r}| = r$, and describes how the particle density varies as a function of the distance from a given particle. Due to the normalization with the particle density n , $g(r) \rightarrow 1$ for $r \gg d$, if there are no long range correlations. Furthermore, since hard spheres cannot overlap, obviously $g(r) = 0$ for $r < d$.

In between these two limits, $g(r)$ can give information about the structure of the system. If e.g. there is short range order and close-by particles arrange in small lattice structures, $g(r)$ shows peaks at distances corresponding to a few next neighbor steps of the respective lattice.

In Fig. 5.24 the radial distribution function is shown for a system with $N \approx 10^6$ and $\phi = 1.96\%$ at different times. For details on the computation of $g(r)$, see Appendix D.3. As a consistency check, we confirm that the simulation data for the (equilibrated) start configuration (Fig. 5.24(a): $t = 0$) agrees with the well known Percus-Yevick solution for hard spheres [Boon & Yip, 1991], an approximation for $g(r)$ that is presumably excellent for low volume fractions.

For $t \approx 3/f_{\text{coll}}$, Fig. 5.24(b), each particle had on average three collisions, and the temperature $T \approx 44\Delta E$ is still much higher than ΔE . However, one can already see the effect of the attractive bonds, since $g(r)$ has significantly increased beyond 1 between d and d_c . This observation has also been made for similar systems with hard core repulsion, but *energy conserving* square-well potential: For this system, Dawson *et al.* [2000] also find a $g(r)$ much larger than 1 for $d < r < d_c$, with discontinuous jumps at d and d_c .

During the cooling regime, as long as $T \gg \Delta E$, this graph remains qualitatively the same, with an increasing area under the curve between d and d_c . As the transition to the clustering regime is approached (Fig. 5.24(c), $t \approx t_0$ and $T \approx \Delta E$) and larger stable clusters emerge, peaks arise in $g(r)$ also for $r > d_c$, corresponding to second and further next neighbor distances in the clusters.

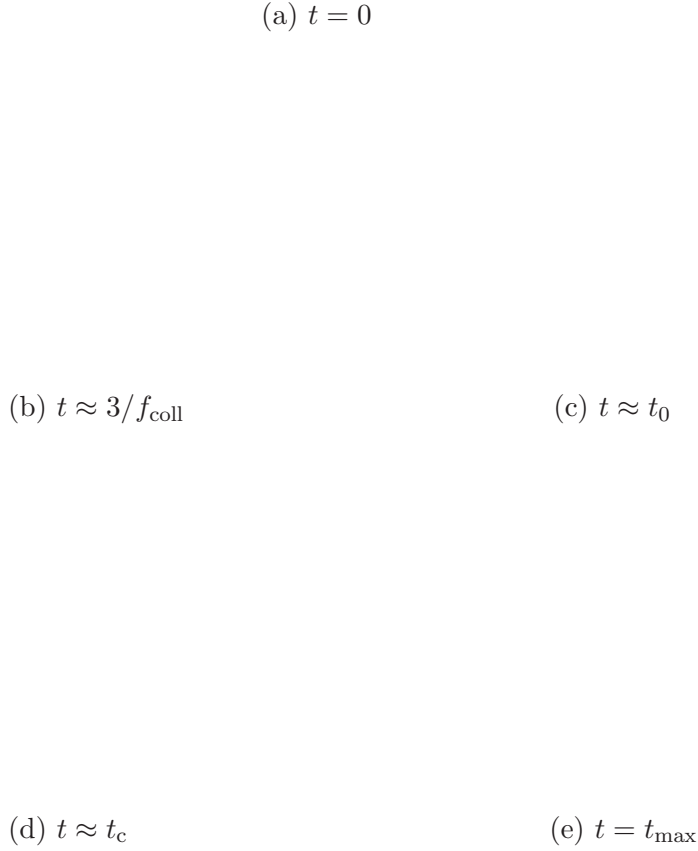


Figure 5.24: Radial distribution function $g(r)$ for a system with $N = 1048576$ particles and volume fraction $\phi = 1.96\%$. In (a), the equilibrated start configuration is shown and compared to the Percus-Yevick solution (blue) for $g(r)$. In (b), $t \approx 3/f_{\text{coll}}$ and the temperature $T/\Delta E \approx 44 \gg 1$. In (c), data is shown for $t \approx t_0$, the transition point between cooling and clustering regime, with $T/\Delta E \approx 1$. In (d), $t \approx t_c$, i.e. a percolating cluster has emerged for the first time. And in (e), the latest simulated time is shown, in which the largest cluster contains 99.6% of all particles. In (a)-(c), all particles are considered; in (d) and (e) data for *all particles* (green) and for *the percolating cluster only* (blue) are superimposed, but they are virtually identical in the shown region of small r . The vertical red lines in (b)-(e) show the hard core distance d , the bond breaking distance d_c , and the second nearest neighbor distance d_{HCP} of a hexagonal close packed structure. The dashed horizontal gray line in all plots corresponds to $g(r) = 1$.

In Fig. 5.24(d), $g(r)$ is shown at the percolation time $t_c \approx 4t_0$, at which a system-spanning cluster has developed. Here one can see very well that the correlations extend further than the next-neighbor distance. The first peak for $r > d_c$ appears at the second nearest neighbor distance of the hexagonal closed packed lattice $d_{\text{HCP}} = \frac{1}{2}\sqrt{3} \cdot \frac{d+d_c}{2}$; ⁹ the further peaks appear in accordance with other lattice distances. For the green and blue curve, respectively, *all* particles and only particles of the *percolating cluster* are considered; however, there seems to be no difference in the shown “short range” structure ($r \leq 3d$), as the two curves are almost identical.

Fig. 5.24(e) shows $g(r)$ at the latest simulated time ($t \approx 40t_0$), again considering *all* particles (green) and only the percolating cluster (blue), which now contains 99.6% of all particles. One can see that the area under the curve between d and d_c has marginally increased, but other than that there are no significant differences to (d), suggesting that the characteristics of the local structure has not significantly changed on the *small length scales* ($d_c < r \leq 3d$), even though the simulation time has increased by a factor of 10 and extensive restructuring of the cluster has taken place.

On length scales much larger than the particle diameter, the RDF can give information about the dimensionality of the respective structure. Let us, for example, consider a *planar object* embedded in a three dimensional system, and have a look at some point on this object. A volume element at distance r from this point is less and less likely to contain a part of that object, as r is increased; this is because, the *surface* of a sphere grows with its radius r like r^2 , but the intersection of that sphere with the planar object only like r ; hence $g(r)$ should fall off like $\frac{r}{r^2} = \frac{1}{r}$. In this regard, one defines the so called **correlation dimension** (which we also denote D_f) [Grassberger, 1983; Rottureau *et al.*, 2004] for fractal objects by

$$g(r) \propto r^{D_f-D} \quad \text{for } r \gg d. \quad (5.46)$$

As we already noted before, for a structure with homogeneous density (*i.e.* $D_f = D$), we get $g(r) \rightarrow 1$ in the limit of large r .

Fig. 5.25 shows the decay of $g(r)$ in the large r regime for the final cluster of the same system as above. Consistent with the fractal behavior with previous definitions of the fractal dimension (Secs. 5.4.2.1 and 5.4.2.2), there appears to be a cross-over at about $r \approx 20d$ from the fractal behavior on smaller length scales to a compact behavior on larger length scales. Due to the oscillations of $g(r)$ up to about $r = 5d$ and the large cross-over region, the regime with well-defined slope only covers half a decade, which makes the *precise* identification of D_f impossible. Nevertheless this regime appears to be consistent with a fractal dimension of $D_f = 2$ (blue line), the result of the box-counting and the radius-of-gyration method.

Beyond the cross-over point, we expect the structure to be compact, which would result in a horizontal progression of $g(r)$. However, due to the finite system size, $g(r)$ is eventually cut off, hence the cross-over to a horizontal line cannot be observed well. As we found in Sec. 5.4.2.2, the size of the fractal regime is proportional to

⁹ We assumed that the lattice constant is the mean between d and d_c .

Figure 5.25: Radial distribution function $g(r)$ for the same system as in Fig. 5.24, however, in the regime $r \gg d$. The blue line has a slope of -1 corresponding to a correlation dimension $D_f = 2$. The system size is $L = 304d$. The inset shows data of the final cluster for a system with higher density $\phi = 7.8\%$ and $N = 2\,097\,152$ particles.

$|\log \phi|$ and hence should decrease as the density ϕ is increased. Thus, the inset of Fig. 5.25 shows data for a denser system with $\phi = 7.8\%$ and $N \approx 2 \cdot 10^6$, where the cross-over to compact behavior can be observed well.

5.4.2.4 Coordination Number

Given the definition of a neighborhood relation (two particles are neighbors if they have built a bridge and their kinetic energy is not sufficient to break it), we can extract the average number of neighbors of a particle, *i.e.* the average coordination number. A good understanding of the development of the system can be obtained from the *evolution* of the coordination numbers. This is shown for the thick film model in Fig. 5.26. Here the clustering transition can be seen very well: Initially

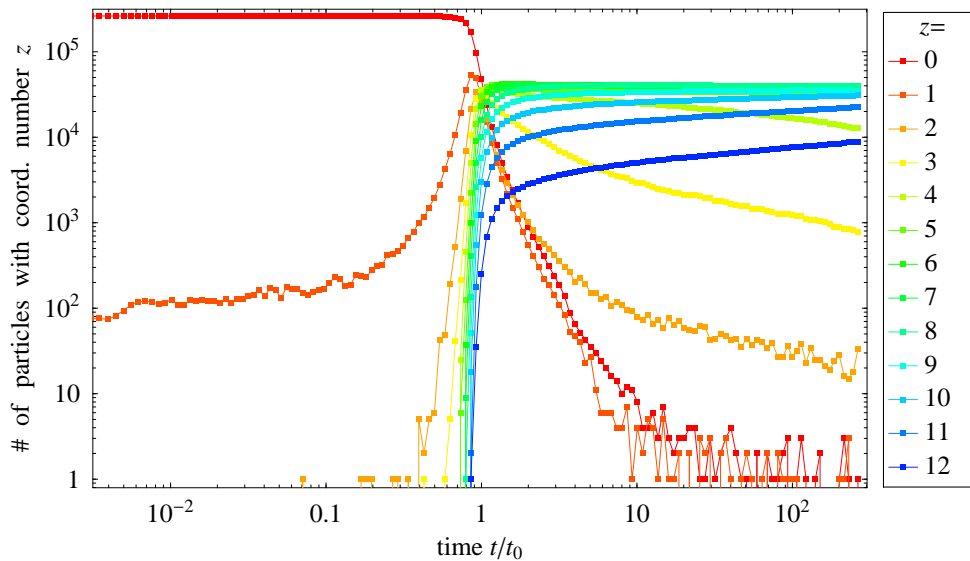


Figure 5.26: Time evolution of the coordination numbers, considering all particles. The data points show the number of particles with coordination number z , where different colors correspond to different z , as indicated by the legend on the right. The system has $N = 262144$ particles, volume fraction $\phi = 1.95\%$, and bond breaking distance $d_c = 1.07d$.

mostly single particles are found, as the dominant coordination number is $z = 0$. Only a small fraction appears as dimers ($z = 1$) and larger aggregates are negligible. At the transition $t = t_0$, larger coordination numbers $z = 2, 3, 4, \dots$ increase sharply, one after another, indicating that larger aggregates are forming. After this sharp increase at the transition point, a coordination number of about $z \approx 6$ is most prevailing. As time proceeds smaller coordination numbers slowly diminish, and larger ones slowly continue to increase.

Let us now have a closer look at the nature of the asymptotic cluster: Fig. 5.27 (top) shows histograms for the coordination number in the percolating cluster for the thick film model with two different bond-breaking distances. As one would expect, these distributions are rather broad with coordination numbers between

one and twelve¹⁰. The smaller bond-breaking distance (left) gives rise to a more asymmetric distribution with more weight for smaller coordination numbers. The bottom panel of Fig. 5.27 shows the typical histogram of coordination numbers for the thin film model. As we can see, the dependence on d_c is weaker for the thin film model and the asymmetry for small d_c is less pronounced.

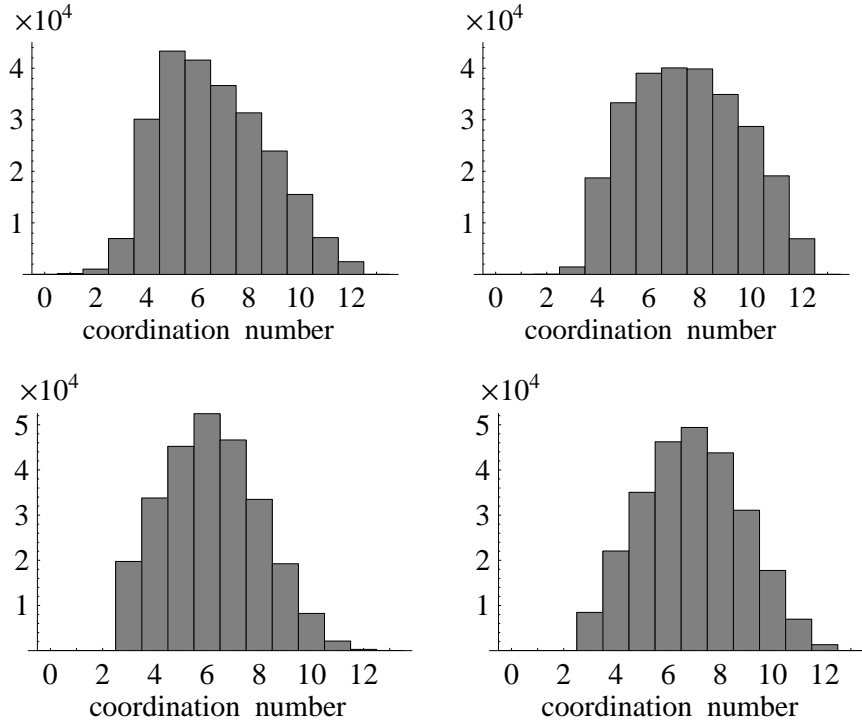


Figure 5.27: Top: Histogram of the coordination number of the asymptotic cluster for two different bond breaking distances $d_c = 1.01d$ (left) and $d_c = 1.07d$ (right); both plots show data from the thick film model with $N = 262144$ and $\phi = 1.95\%$. Bottom: As top, but for the thin film model.

More comprehensible than the full histogram is the *average* coordination number of the largest cluster, whose time evolution is shown in Fig. 5.28 for different bond-breaking distances d_c . After a strong increase at the time t_0 , the average coordination number continues to grow slowly. This slow increase is strongly suppressed in the thin film model (right) as compared to the thick film model (left). Within the thin film model the slow growth with time is further suppressed for decreasing bond-breaking distance d_c .

As one can see in Figs. 5.27, 5.28, the coordination number becomes smaller for smaller d_c . This is reasonable, because the particles can more easily collect neighbors for higher d_c . As $d_c \rightarrow d$ the average coordination number of the thin film model approaches 6, which is the isostatic value. This is demonstrated in

¹⁰Very rarely, a coordination number of $z = 13$ is observed for $d_c = 1.07d$, however not in the presented system.

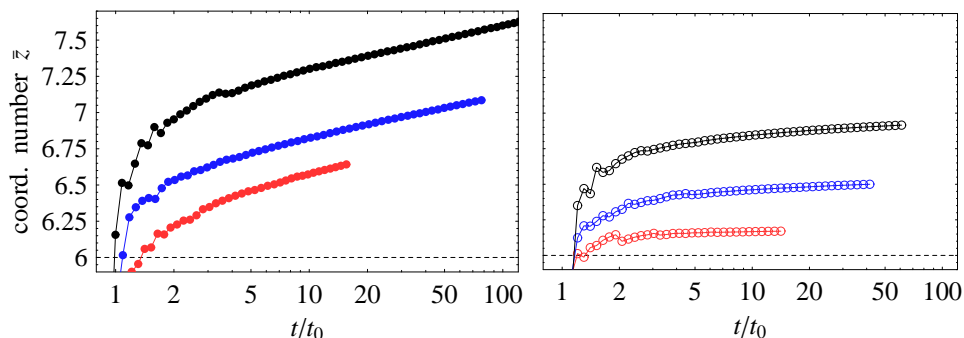


Figure 5.28: Time evolution of the average coordination number \bar{z} of particles in the largest cluster ($N = 262144$ and $\phi = 1.95\%$); the critical break-off distances are $d_c = 1.07d, 1.035d$ and $1.01d$ from top to bottom; thick film model (left) in comparison to thin film model (right).

Fig. 5.29, where we plot the asymptotic coordination number as a function of d_c . Here the asymptotic value is taken, when $T < 0.06\Delta E$ for the first time.

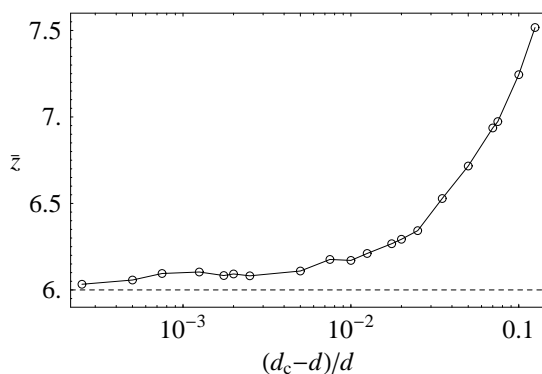


Figure 5.29: Influence of the bond-breaking distance d_c on the final value of the average coordination number \bar{z} for the thin film model. $\phi = 0.24\%$ and $N = 10648$.

Naively one might expect that the increase of the coordination number with larger d_c is caused by a compactification and therefore accompanied by an increase of the fractal dimension. However, as can be seen in Fig. 5.30, there is no significant influence of d_c on the development of the fractal dimension. Thus, we conclude that this compactification is mostly occurring on the single particle length scale and therefore increasing the average coordination number, but not influencing the structure on larger length scales.¹¹

¹¹Note that there is also a very slight increase of the fractal dimension and therefore also a very slow compactification on larger length scales. However these effects are much less pronounced than the change of the coordination number.

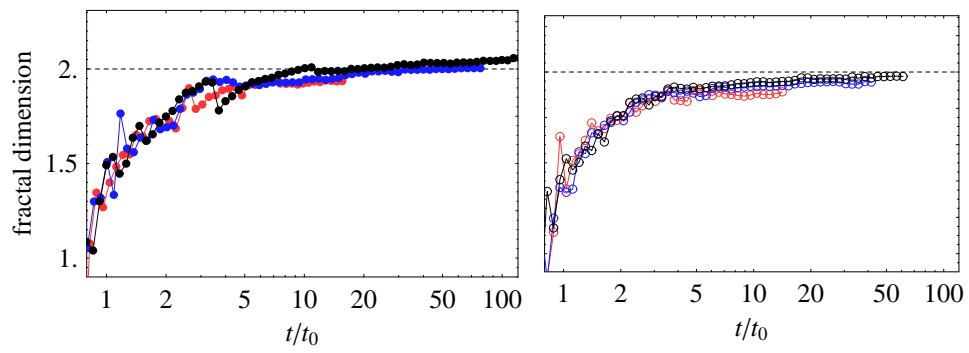


Figure 5.30: Time evolution of the fractal dimension D_f of the largest cluster. System and colors as in Fig. 5.28.

5.4.2.5 Inertia Tensor and Spatial Extension

Throughout this section 5.4.2 we repeatedly observe a fractal dimension of $D_f \approx 2$ for the asymptotic cluster; this raises the question, whether or not the cluster largely consists of *planar* topologies. To gain quantitative information about the shape of an object, it is useful to compute its **moment of inertia tensor** [see e.g. Landau & Lifshitz, 1976] and look at its eigenvalues, the **principal moments of inertia** $I_1 \leq I_2 \leq I_3$. The latter quantify the mass distribution perpendicular to the respective **principal axes** $\mathbf{e}_1, \mathbf{e}_2, \mathbf{e}_3$, as they obey the equation:

$$I_1 = \int \varrho(\mathbf{r}) (r_2^2 + r_3^2) \, \mathrm{d}\mathbf{r}, \quad (5.47)$$

and equivalent equations for I_2, I_3 . Here $\mathbf{r} = \sum_{\nu=1}^3 r_\nu \mathbf{e}_\nu$ is the distance from the center of mass in the coordinate system of the principal axes and $\varrho(\mathbf{r})$ the density at that point.

The principal moments of inertia can easily be related to a more intuitively comprehensible quantity, the spatial extensions Δr_ν along the three principal axes:

$$\Delta r_\nu^2 := \frac{\int \varrho(\mathbf{r}) r_\nu^2 \, \mathrm{d}\mathbf{r}}{\int \varrho(\mathbf{r}) \, \mathrm{d}\mathbf{r}} \quad \text{for } \nu = 1, 2, 3. \quad (5.48)$$

While the denominator is simply the mass of the object, the numerator can be expressed in terms of the principal moments; e.g. for $\nu = 1$:

$$\begin{aligned} \frac{-I_1 + I_2 + I_3}{2} &= \int \varrho(\mathbf{r}) \frac{-(r_2^2 + r_3^2) + (r_2^2 + r_3^2) + (r_1^2 + r_2^2)}{2} \, \mathrm{d}\mathbf{r} \\ &= \int \varrho(\mathbf{r}) r_1^2 \, \mathrm{d}\mathbf{r}. \end{aligned} \quad (5.49)$$

The reason for not calculating Eq. (5.48) directly, but using the detour *via* the principal moments of inertia, is that *a priori* the principal axes are not known. Note furthermore that from these spatial extensions one can trivially obtain the radius of gyration $r_g^2 = \sum_{\nu=1}^3 \Delta r_\nu^2$, as one can easily see in the definition (5.48). The choice of the principal axes is always such that $\Delta r_1 \geq \Delta r_2 \geq \Delta r_3$. For spherical symmetric objects obviously $\Delta r_1 = \Delta r_2 = \Delta r_3$; for plate-like, or so-called *oblate* objects $\Delta r_1 \approx \Delta r_2 \gg \Delta r_3$, and for rod-like, or *prolate* objects $\Delta r_1 \gg \Delta r_2 \approx \Delta r_3$.

In order to obtain information about the structure on a wide range of length scales, we analyze *partial clusters* deduced from the largest cluster, as done in Sec. 5.4.2.1: starting from a random initial particle of the cluster, we define all particles that can be reached by i or less neighbor-to-neighbor steps as partial cluster $\mathcal{N}(i)$ (see Fig. 5.19). The spatial extensions $\Delta r_{1,2,3}$ of $\mathcal{N}(i)$ are calculated for different values of i and the data are averaged over several initial particles.¹² The result is shown in Fig. 5.31. As one can see, the structures are clearly not spher-

¹²For simplicity only the particle *centers* are considered for the calculation of the inertia tensor. As we can easily convince ourselves in Eq. (5.49), due to the Huygens-Steiner theorem, accounting for the extension of the individual particles would just lead to a constant shift of $\Delta r_{1,2,3}^2 \rightarrow \Delta r_{1,2,3}^2 + \frac{1}{2} I_{\text{sphere}}$ with the moment of inertia of a single sphere $I_{\text{sphere}} = \frac{1}{10} d^2$.

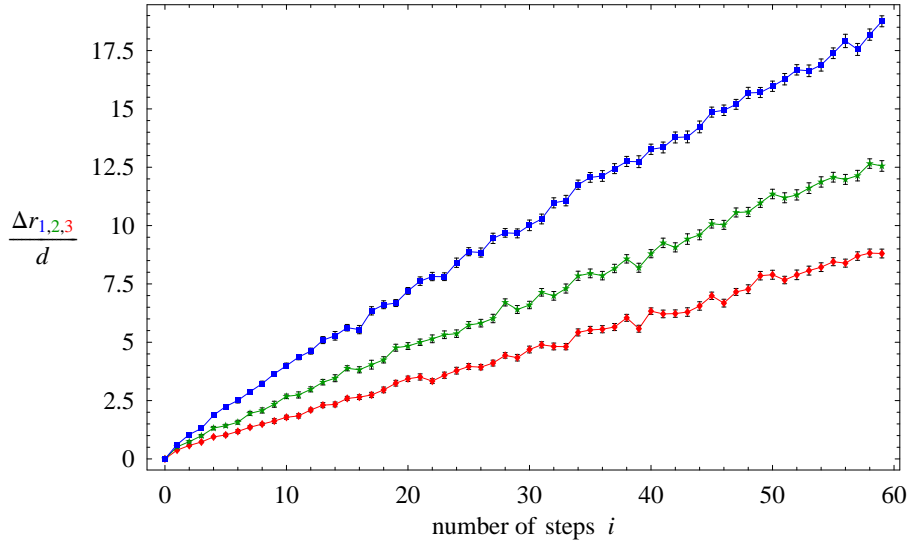


Figure 5.31: Spatial extension of partial clusters $\mathcal{N}(i)$ along the principal axes. The blue curve shows the largest component Δr_1 , the green curve the second largest Δr_2 , and the red curve the smallest Δr_3 . The inset shows data for the same procedure, for which, however, the particle positions are not obtained from the simulation, but randomly placed into the system.

ically symmetric, as the spatial extensions in the principal directions are different for all shown length scales. However, also the assumption that the cluster involves *planar* structures cannot be supported, since that would imply $\Delta r_1 \approx \Delta r_2 \gg \Delta r_3$ at least on some intermediate length scales. Indeed, we observe the most general case $\Delta r_1 \not\approx \Delta r_2 \not\approx \Delta r_3$ throughout the investigated range.

5.5 Conclusions

We have analyzed a simple model of a wet granulate allowing for large scale event driven simulations. A central feature of wet granulates is the existence of an energy scale ΔE associated with the rupture of a capillary bridge between two grains. This energy scale has important consequences not only for the phase diagram [Fingerle *et al.*, 2008] but also for the free cooling dynamics investigated in this paper. The most important feature is a well defined transition at a time t_0 , when the kinetic energy T of the particles becomes equal to ΔE .

For $t < t_0$ the particles are energetic enough to supply the bond breaking energy ΔE , so that very few collisions result in bound pairs and most particles are unbound. Cooling is very effective in this regime, but drastically different from a dry granulate. Whereas in dry granulates the dissipated energy is proportional to the energy of the colliding particles, in wet granulates the dissipated energy is ΔE , independent of the energy of the colliding particles so that $\dot{T} \propto \sqrt{T}$. Consequently Haff's law does

not hold and is replaced by $T(t) = T(0)(1 - t/t_0)^2$ for $t < t_0$. The simulations are in very good agreement with this cooling law for $t < t_0$.

For $t > t_0$, the kinetic energy of the particles is too small to provide the bond breaking energy, so that larger and larger clusters form. We call this regime the aggregation regime and analyze the properties of the aggregates. For not too long times and sufficiently small volume fractions, we observe flocculation characterized by nonoverlapping, weakly interacting clusters. The fractal dimension of the aggregates is $D_f \approx 2$. The cluster mass distribution follows a simple scaling form, $N_m(t) \propto m^{-2} f(m/\bar{m}(t))$, which has been applied successfully to different aggregation models before. The increase of the typical cluster size $\bar{m}(t)$ can be understood by a simple scaling analysis: Assuming that clusters irreversibly stick together when they hit upon each other and that their radius r grows with the number of particles m like $r^{D_f} \propto m$, yields a cluster growth $\bar{m} \propto t^{2D_f/(3D_f-2D+2)}$. This scaling relation shows good agreement with the simulation for fractal dimension $D_f = 2$.

At larger times, a spanning cluster forms, and a gelation transition is observed for all finite volume fractions. At the gelation transition a spanning cluster coexists with many small ones, whereas at very long times almost all particles are connected to one large cluster. On the largest length scales the final cluster is no longer a fractal but compact, as one would expect for a spanning cluster in the percolating phase. On smaller length scales, however, we still find fractal structures with $D_f \approx 2$. The range where a nontrivial fractal dimension can be observed increases with decreasing density as $|\log \phi|$.

Even on the longest time-scales, the temperature continues to decay. In this regime the limiting process is the breaking of a bond. The probability for this process becomes exponentially small $P_{bb} \propto \sqrt{\Delta E/T} e^{-\Delta E/T}$ as the temperature goes to zero. Hence the cooling law for high temperatures is replaced by $\dot{T} \propto e^{-\Delta E/T}$, yielding a slow (logarithmic) temperature decay, which is in very good agreement with the data.

Several extensions of the presented work might be interesting. So far we have completely neglected all inelasticities except for the bond rupture. One expects the collisions at the hard core to be dissipative as they are in dry granular media. In the simplest model these could be described by normal restitution. Furthermore, real wet grains experience frictional forces, coupling translational and rotational motion of the grains [Brilliantov *et al.*, 2007]. However, the author is not aware of any such studies for wet granulates at the present time. In the context of gelation it would be interesting to see, if and how a finite shear modulus emerges in the system, as the percolation transition is passed.

Properties of \perp - and \parallel -vectors

The properties of \perp - and \parallel -vectors discussed here apply to the calculations in chapters 2 and 3. For any given replicated vector $\hat{x} = (\mathbf{x}^{(0)}, \dots, \mathbf{x}^{(n)})^T$, define:

$$\mathbf{x}_{\parallel} := \frac{\sum_{\alpha=0}^n \mathbf{x}^{(\alpha)}}{\sqrt{n+1}} \quad (\text{A.1})$$

and

$$x_{\perp} := \begin{pmatrix} \mathbf{x}^{(0)} \\ \vdots \\ \mathbf{x}^{(n)} \end{pmatrix} - \frac{1}{n+1} \begin{pmatrix} \sum_{\alpha=0}^n \mathbf{x}^{(\alpha)} \\ \vdots \\ \sum_{\alpha=0}^n \mathbf{x}^{(\alpha)} \end{pmatrix} = \hat{x} - \frac{\mathbf{x}_{\parallel} \otimes \hat{\varepsilon}}{\sqrt{n+1}} \quad (\text{A.2})$$

with $\hat{\varepsilon} = (1, \dots, 1)^T$. Then these vectors have the following properties:

- The mean of all components of x_{\perp} is $\mathbf{x}_{\perp}^{(0)} + \dots + \mathbf{x}_{\perp}^{(n)} = \mathbf{0}$
- Therefore $(\mathbf{x}_{\parallel} \otimes \hat{\varepsilon}) \cdot x_{\perp} = 0$
- $\hat{x} \cdot \hat{y} = \mathbf{x}_{\parallel} \cdot \mathbf{y}_{\parallel} + x_{\perp} \cdot y_{\perp}$

When integrating over a function $f(\hat{x}) = f(\mathbf{x}_{\parallel}, x_{\perp})$, we might ask the the question how to change integration variables

$$\int_{V^n} f(\hat{x}) d\hat{x} \rightarrow \int_{V^n} f(\mathbf{x}_{\parallel}, x_{\perp}) d\mathbf{x}_{\parallel} dx_{\perp}, \quad (\text{A.3})$$

whereby

$$\begin{aligned} d\hat{x} &= d\mathbf{x}^{(0)} \dots d\mathbf{x}^{(n)} \quad \text{and} \\ dx_{\perp} &= d\mathbf{x}_{\perp}^{(0)} \dots d\mathbf{x}_{\perp}^{(n)} \delta(\mathbf{x}_{\perp}^{(0)} + \dots + \mathbf{x}_{\perp}^{(n)}) \\ &= d\mathbf{x}_{\perp}^{(1)} \dots d\mathbf{x}_{\perp}^{(n)}. \end{aligned} \quad (\text{A.4})$$

We already noted that the mean component of x_{\perp} must be zero, therefore it is sufficient to integrate over only n of the $n+1$ components, or use the δ -function to satisfy this constraint. For the variable transformation (A.3), we now successively

change the individual variables:

$$\begin{aligned}
& d\mathbf{x}_{\parallel} d\mathbf{x}_{\perp}^{(1)} d\mathbf{x}_{\perp}^{(2)} \cdots d\mathbf{x}_{\perp}^{(n)} \\
&= \left| \frac{\partial \mathbf{x}_{\parallel}}{\partial \mathbf{x}^{(0)}} \right|_{\mathbf{x}_{\perp}^{(1), \dots, (n)}} d\mathbf{x}^{(0)} d\mathbf{x}_{\perp}^{(1)} d\mathbf{x}_{\perp}^{(2)} \cdots d\mathbf{x}_{\perp}^{(n)} \\
&= \left| \frac{\partial \mathbf{x}_{\parallel}}{\partial \mathbf{x}^{(0)}} \right|_{\mathbf{x}_{\perp}^{(1), \dots, (n)}} \left| \frac{\partial \mathbf{x}_{\perp}^{(1)}}{\partial \mathbf{x}^{(1)}} \right|_{\mathbf{x}^{(0)}, \mathbf{x}_{\perp}^{(2), \dots, (n)}} d\mathbf{x}^{(0)} d\mathbf{x}^{(1)} d\mathbf{x}_{\perp}^{(2)} \cdots d\mathbf{x}_{\perp}^{(n)} \\
&\quad \vdots \\
&= \left| \frac{\partial \mathbf{x}_{\parallel}}{\partial \mathbf{x}^{(0)}} \right|_{\mathbf{x}_{\perp}^{(1), \dots, (n)}} \left| \frac{\partial \mathbf{x}_{\perp}^{(1)}}{\partial \mathbf{x}^{(1)}} \right|_{\mathbf{x}^{(0)}, \mathbf{x}_{\perp}^{(2), \dots, (n)}} \cdots \left| \frac{\partial \mathbf{x}_{\perp}^{(n)}}{\partial \mathbf{x}^{(n)}} \right|_{\mathbf{x}^{(0), \dots, (n-1)}} d\mathbf{x}^{(0)} d\mathbf{x}^{(1)} d\mathbf{x}^{(2)} \cdots d\mathbf{x}^{(n)}
\end{aligned} \tag{A.5}$$

In each transformation the determinant of the Jacobian matrix for (e.g. $\left| \frac{\partial \mathbf{x}_{\perp}^{(n)}}{\partial \mathbf{x}^{(n)}} \right|$) has to be evaluated keeping the variables constant, which are currently not transformed (denoted by $|\cdot|_{\mathbf{x}^{(0), \dots, (n-1)}}$). With the definition (A.1) and (A.2) one can easily confirm that

$$\left| \frac{\partial \mathbf{x}_{\parallel}}{\partial \mathbf{x}^{(0)}} \right|_{\mathbf{x}_{\perp}^{(1), \dots, (n)}} = (n+1)^{D/2}, \tag{A.6}$$

and for $1 \leq j \leq n$:

$$\left| \frac{\partial \mathbf{x}_{\perp}^{(j)}}{\partial \mathbf{x}^{(j)}} \right|_{\mathbf{x}^{(0), \dots, (j-1)}, \mathbf{x}_{\perp}^{(j+1), \dots, (n)}} = \left(\frac{j}{j+1} \right)^D. \tag{A.7}$$

Therefore we get:

$$\begin{aligned}
& \left| \frac{\partial \mathbf{x}_{\parallel}}{\partial \mathbf{x}^{(0)}} \right|_{\mathbf{x}_{\perp}^{(1), \dots, (n)}} \left| \frac{\partial \mathbf{x}_{\perp}^{(1)}}{\partial \mathbf{x}^{(1)}} \right|_{\mathbf{x}^{(0)}, \mathbf{x}_{\perp}^{(2), \dots, (n)}} \cdots \left| \frac{\partial \mathbf{x}_{\perp}^{(n)}}{\partial \mathbf{x}^{(n)}} \right|_{\mathbf{x}^{(0), \dots, (n-1)}} = \\
&= (n+1)^{D/2} \left(\frac{1}{2} \right)^D \left(\frac{2}{3} \right)^D \cdots \left(\frac{n}{n+1} \right)^D \\
&= (n+1)^{D/2} \left(\frac{1}{n+1} \right)^D = (n+1)^{-D/2}
\end{aligned} \tag{A.8}$$

Hence the transformation can be written as:

$$\int_{V^n} f(\hat{x}) d\hat{x} = (n+1)^{D/2} \int_{V^n} f(\mathbf{x}_{\parallel}, x_{\perp}) d\mathbf{x}_{\parallel} dx_{\perp}, \tag{A.9}$$

Calculations for the Randomly Cross-Linked Particle Model

Contents

| | | |
|------------|---|------------|
| B.1 | Calculation of the Replica Free Energy | 128 |
| B.1.1 | Emergence of the Replica Free Energy | 128 |
| B.1.2 | Introduction of the Replicated Density Field | 129 |
| B.2 | Hubbard-Stratonovich Transformation | 130 |
| B.2.1 | Applying the Hubbard-Stratonovich Transformation | 130 |
| B.2.2 | Meaning of the Order Parameter $\Omega(\hat{q})$ | 132 |
| B.3 | Expansion of the replica free energy to 3rd Order in Ω | 134 |
| B.4 | Fluctuations around the Saddle-Point Solution | 135 |
| B.5 | Replica Free Energy with Shear Deformations | 136 |
| B.5.1 | Preparing the Replica Free Energy for the Ansatz | 136 |
| B.5.2 | Expansion of the one-particle partition function \mathfrak{z} | 137 |
| B.5.3 | Insertion of the Order Parameter in \mathfrak{z} | 137 |
| B.5.4 | Insertion of the Order Parameter in $f_{n+1}\{\Omega\}$ | 145 |
| B.5.5 | Recomposing the Replica Free Energy | 146 |
| B.5.6 | Analysis of the quantity Ξ_{r,a^2} | 149 |
| B.6 | Auxiliary Calculations for the Results Section | 150 |
| B.6.1 | Distribution of the Number of Cross-Links | 150 |
| B.6.2 | Shear modulus | 152 |
| B.6.3 | Square of the expansion for $(1 - Q)$ | 153 |
| B.7 | The Order Parameter in the One Replica Sector | 154 |

B.1 Calculation of the Replica Free Energy

B.1.1 Emergence of the Replica Free Energy

Here we plug Eq. (2.12) into Eq. (2.11) and define $\tilde{\mu}^2 := \frac{\mu^2}{2N\phi}$ to get:

$$\begin{aligned}
[Z_C^n] &= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{(\tilde{\mu}^2)^M}{M!} \\
&\times \int \left(\prod_{j=1}^N D\hat{r}_j \right) \exp \left(-\frac{1}{2a^2} \sum_{e=1}^M (\hat{r}_{i_e} - \hat{r}_{j_e})^2 - H_{\text{ev}}^{(n+1)} \right)
\end{aligned} \tag{B.1}$$

A part of this expression can be simplified:

$$\begin{aligned}
&\sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{(\tilde{\mu}^2)^M}{M!} \exp \left(-\frac{1}{2a^2} \sum_{e=1}^M (\hat{r}_{i_e} - \hat{r}_{j_e})^2 \right) \\
&= \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2)^M}{M!} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \prod_{e=1}^M \exp \left(-\frac{1}{2a^2} (\hat{r}_{i_e} - \hat{r}_{j_e})^2 \right) \\
&= \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2)^M}{M!} \left(\sum_{i,j=1}^N \exp \left(-\frac{(\hat{r}_i - \hat{r}_j)^2}{2a^2} \right) \right)^M \\
&= \exp \left(\tilde{\mu}^2 \sum_{i,j=1}^N \exp \left(-\frac{(\hat{r}_i - \hat{r}_j)^2}{2a^2} \right) \right)
\end{aligned} \tag{B.2}$$

Hence Eq. (B.1) becomes:

$$\begin{aligned}
[Z^n] &= \frac{1}{\mathcal{Z}_1} \int \left(\prod_{j=1}^N D\hat{r}_j \right) \exp \left(\tilde{\mu}^2 \sum_{i,j=1}^N \exp \left(-\frac{(\hat{r}_i - \hat{r}_j)^2}{2a^2} \right) - H_{\text{ev}}^{(n+1)} \right) \\
&=: \frac{\mathcal{Z}_{n+1}}{\mathcal{Z}_1}
\end{aligned} \tag{B.3}$$

where

$$\mathcal{Z}_{n+1} = \int \left(\prod_{j=1}^N D\hat{r}_j \right) \exp(-N\tilde{f}_{n+1}\{\hat{r}_j\}) \tag{B.4a}$$

$$\text{and } \tilde{f}_{n+1}\{\hat{r}_j\} = -\frac{\tilde{\mu}^2}{N} \sum_{i,j=1}^N \exp \left(-\frac{(\hat{r}_i - \hat{r}_j)^2}{2a^2} \right) + \frac{H_{\text{ev}}^{(n+1)}}{N} \tag{B.4b}$$

B.1.2 Introduction of the Replicated Density Field

In terms of the replicated density field $O(\hat{x}) = \frac{1}{N} \sum_{j=1}^N \delta(\hat{x} - \hat{r}_j)$, Eq. (2.16), we can rewrite the cross-linking term of Eq. (2.14c)/(B.4b):

$$\begin{aligned}
& \sum_{i,j=1}^N \exp\left(-\frac{(\hat{r}_i - \hat{r}_j)^2}{2a^2}\right) \\
&= \sum_{i,j=1}^N \int d\hat{x} d\hat{y} \delta(\hat{x} - \hat{r}_i) \delta(\hat{y} - \hat{r}_j) \exp\left(-\frac{(\hat{x} - \hat{y})^2}{2a^2}\right) \\
&= N^2 \int d\hat{x} d\hat{y} O(\hat{x}) O(\hat{y}) \exp\left(-\frac{(\hat{x} - \hat{y})^2}{2a^2}\right)
\end{aligned} \tag{B.5}$$

The Fourier transform of the Gaussian is

$$\int d\hat{x} \exp(-i\hat{q}\hat{x}) \exp\left(-\frac{\hat{x}^2}{2a^2}\right) = (2\pi a^2)^{D(n+1)/2} \exp\left(-\frac{\hat{q}^2 a^2}{2}\right), \tag{B.6}$$

and thus we can use Parseval's identity to transform the whole term (B.5) into Fourier space:

$$\begin{aligned}
& N^2 \int d\hat{x} d\hat{y} O(\hat{x}) O(\hat{y}) \exp\left(-\frac{(\hat{x} - \hat{y})^2}{2a^2}\right) \\
&= N^2 \frac{(2\pi a^2)^{D(n+1)/2}}{V^{n+1}} \sum_{\hat{q}} O(\hat{q}) O(-\hat{q}) \exp\left(-\frac{\hat{q}^2 a^2}{2}\right) \\
&= N^2 \phi^{n+1} \sum_{\hat{q}} |O(\hat{q})|^2 \Delta(\hat{q}),
\end{aligned} \tag{B.7}$$

with

$$\Delta(\hat{q}) := \exp\left(-\frac{\hat{q}^2 a^2}{2}\right) \tag{B.8}$$

Analogously we can rewrite the excluded volume Hamiltonian in terms of the replicated density field:

$$\begin{aligned}
H_{\text{ev}}^{(n+1)} &= \frac{\lambda}{2} \sum_{\alpha=0}^n \sum_{i,j=1}^N U(\mathbf{r}_i^{(\alpha)} - \mathbf{r}_j^{(\alpha)}) \\
&= \frac{\lambda}{2} \sum_{\alpha=0}^n \sum_{i,j=1}^N \int d\mathbf{x} d\mathbf{y} \delta(\mathbf{x} - \mathbf{r}_i^{(\alpha)}) \delta(\mathbf{y} - \mathbf{r}_j^{(\alpha)}) U(\mathbf{x} - \mathbf{y}) \\
&= \frac{\lambda N^2}{2} \sum_{\alpha=0}^n \int d\mathbf{x} d\mathbf{y} O^{(\alpha)}(\mathbf{x}) O^{(\alpha)}(\mathbf{y}) U(\mathbf{x} - \mathbf{y}) \\
&= \frac{\lambda N^2}{2V} \sum_{\alpha=0}^n \sum_{\mathbf{q}} |O^{(\alpha)}(\mathbf{q})|^2 U(\mathbf{q})
\end{aligned} \tag{B.9}$$

Now, we can plug Eqs. (B.7) and (B.9) back into Eq. (2.14c) and get:

$$\tilde{f}_{n+1}\{\hat{r}_j\} = -\phi^n \frac{\mu^2}{2} \sum_{\hat{q}} |\text{O}(\hat{q})|^2 \Delta(\hat{q}) + \frac{\lambda n_0}{2} \sum_{\alpha=0}^n \sum_{\mathbf{q}} |\text{O}^{(\alpha)}(\mathbf{q})|^2 U(\mathbf{q}), \quad (\text{B.10})$$

with the mean particle density $n_0 = N/V$.

The sums in Eq. (B.10) include $\hat{q} = \hat{0}$ and $\mathbf{q} = \mathbf{0}$. Those terms are mere constants, since $\text{O}(\hat{0}) = \text{O}^{(\alpha)}(\mathbf{0}) = 1$. To make that clearer in the main calculation, it is useful to exclude those terms from the sums and rewrite Eq. (B.10) as

$$\begin{aligned} \tilde{f}_{n+1}\{\hat{r}_j\} = & \overbrace{-\phi^n \frac{\mu^2}{2} + (n+1) \frac{\lambda n_0}{2}}{=:f_0} \\ & - \phi^n \frac{\mu^2}{2} \sum_{\hat{q} \neq \hat{0}} |\text{O}(\hat{q})|^2 \Delta(\hat{q}) + \frac{n_0 \lambda}{2} \sum_{\alpha=0}^n \sum_{\mathbf{q} \neq \mathbf{0}} |\text{O}^{(\alpha)}(\mathbf{q})|^2 U(\mathbf{q}) \end{aligned} \quad (\text{B.11})$$

B.2 Hubbard-Stratonovich Transformation

B.2.1 Applying the Hubbard-Stratonovich Transformation

The Hubbard-Stratonovich transformation is used to **linearize a quadratic term in an exponential**, whereby one has to put up with the integration over a new variable. In its basic form, the Hubbard-Stratonovich transformation is nothing more than a complex Gaussian integral:

$$\exp(c|w|^2) = \frac{c}{\pi} \int_{\mathbb{C}} dz \exp(-c|z|^2 + 2c \text{Re}(zw^*)), \quad (\text{B.12a})$$

$$\exp(-c|w|^2) = \frac{c}{\pi} \int_{\mathbb{C}} dz \exp(-c|z|^2 + 2ic \text{Re}(zw^*)). \quad (\text{B.12b})$$

Here, $c > 0$ and real, and w^* is the complex conjugate of w . Furthermore $dz = d(\text{Re } z) d(\text{Im } z)$, *i.e.* one has to integrate twice.

Now we want to apply this relation to Eq. (2.24a), with $w = \text{O}(\hat{q})$ successively for all \hat{q} . Looking at the definition of the density $\text{O}(\hat{q})$, Eq. (2.17a), we see that it fulfills the relation $\text{O}(-\hat{q}) = \text{O}^*(\hat{q})$, and hence $\text{O}(\hat{q})$ and $\text{O}(-\hat{q})$ are not independent variables. In particular $|\text{O}(\hat{q})|^2 = |\text{O}(-\hat{q})|^2$. That means, we automatically transform the $|\text{O}(\hat{q})|^2$ -term along with the corresponding $|\text{O}(-\hat{q})|^2$ -term.

To account for that particularity, it is useful to restrict the sums of Eq. (2.24a) to a half-space with $\hat{q} \cdot \hat{e} > 0$ with $\hat{e} = (\mathbf{e}, \dots, \mathbf{e})$ for arbitrary $\mathbf{e} \neq \mathbf{0}$. Thereby we combine the terms for \hat{q} and $-\hat{q}$ in the sum and furthermore use $\Delta(\hat{q}) + \Delta(-\hat{q}) = 2\Delta(\hat{q})$ and $\tilde{\lambda}(\hat{q}) + \tilde{\lambda}(-\hat{q}) = 2\tilde{\lambda}(\hat{q})$ (this is true, because all interactions are symmetric). Thus

we can write the replica partition function (2.14c), with the free energy (2.24a), as:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} &= \int \left(\prod_{j=1}^N D\hat{r}_j \right) \exp \left(N\phi^n \mu^2 \sum_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} |\text{O}(\hat{q})|^2 \Delta(\hat{q}) - N \sum_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} |\text{O}(\hat{q})|^2 \tilde{\lambda}(\hat{q}) \right). \end{aligned} \quad (\text{B.13})$$

Now we apply the Hubbard-Stratonovich transformation (B.12) with $w = \text{O}(\hat{q})$. For any $\hat{q} \in \text{HRS}$, $\hat{q} \cdot \hat{e} > 0$ we use (B.12a) and set $c = N\phi^n \mu^2 \Delta(\hat{q})$:

$$\begin{aligned} &\exp \left(N\phi^n \mu^2 |\text{O}(\hat{q})|^2 \Delta(\hat{q}) \right) \\ &= \int \frac{N\phi^n \mu^2 \Delta(\hat{q}) d\Omega(\hat{q})}{\pi} \exp \left(-N\phi^n \mu^2 \Delta(\hat{q}) |\Omega(\hat{q})|^2 + 2N\phi^n \mu^2 \Delta(\hat{q}) \text{Re}(\Omega(\hat{q})\text{O}(-\hat{q})) \right) \end{aligned} \quad (\text{B.14a})$$

and for $\hat{q} \in \text{1RS}$, $\hat{q} \cdot \hat{e} > 0$, we use (B.12b) with $c = N\tilde{\lambda}(\hat{q})$:

$$\begin{aligned} &\exp \left(-N |\text{O}(\hat{q})|^2 \tilde{\lambda}(\hat{q}) \right) \\ &= \int \frac{N\tilde{\lambda}(\hat{q}) d\Omega(\hat{q})}{\pi} \exp \left(-N |\Omega(\hat{q})|^2 \tilde{\lambda}(\hat{q}) + 2iN\tilde{\lambda}(\hat{q}) \text{Re}(\Omega(\hat{q})\text{O}(-\hat{q})) \right), \end{aligned} \quad (\text{B.14b})$$

With those two relations, Eq. (B.13) becomes:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} &= \int \mathcal{D}\Omega \exp \left(-N\phi^n \mu^2 \sum_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} |\Omega(\hat{q})|^2 \Delta(\hat{q}) - N \sum_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} |\Omega(\hat{q})|^2 \tilde{\lambda}(\hat{q}) \right) \\ &\quad \times \int \left(\prod_{j=1}^N D\hat{r}_j \right) \exp \left(2N\phi^n \mu^2 \sum_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} \Delta(\hat{q}) \text{Re}(\Omega(\hat{q})\text{O}(-\hat{q})) \right. \\ &\quad \left. - 2iN \sum_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} \tilde{\lambda}(\hat{q}) \text{Re}(\Omega(\hat{q})\text{O}(-\hat{q})) \right), \end{aligned} \quad (\text{B.15})$$

where we introduced the measure

$$\mathcal{D}\Omega = \prod_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} \frac{N\tilde{\lambda}(\hat{q}) d\Omega(\hat{q})}{\pi} \cdot \prod_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} \frac{N\phi^n \mu^2 \Delta(\hat{q}) d\Omega(\hat{q})}{\pi}. \quad (\text{B.16})$$

Now we abolish the constraint $\hat{q} \cdot \hat{e} > 0$ in the sums. To this end, we define the field $\Omega(\hat{q})$ also for $\hat{q} \cdot \hat{e} < 0$ by

$$\Omega(-\hat{q}) = \Omega^*(\hat{q}). \quad (\text{B.17})$$

We also plug in the definition of $O(\hat{q})$, Eq. (2.17a), and get:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} &= \int \mathcal{D}\Omega \exp\left(-N\frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} |\Omega(\hat{q})|^2 \Delta(\hat{q}) - \frac{N}{2} \sum_{\hat{q} \in \text{IRS}} |\Omega(\hat{q})|^2 \tilde{\lambda}(\hat{q})\right) \\ &\times \int \left(\prod_{j=1}^N D\hat{r}_j\right) \exp\left(\phi^n \mu^2 \sum_{j=1}^N \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}_j} + i \sum_{j=1}^N \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}_j}\right) \end{aligned} \quad (\text{B.18})$$

At this point, we can see that the Hubbard-Stratonovich transformation pays off: Now, the density $\sum_{j=1}^N e^{-i\hat{q}\hat{r}_j}$ appears linearly in the exponential and we can rewrite the difficult integral $\int (\prod_{j=1}^N D\hat{r}_j)$ in the last line of Eq. (B.18) in the following way:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} &= \int \mathcal{D}\Omega \exp\left(-N\frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) |\Omega(\hat{q})|^2 - \frac{N}{2} \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) |\Omega(\hat{q})|^2\right) \\ &\times \underbrace{\left(\int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}} + i \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}}\right)\right)}_{=: \mathfrak{z}}^N. \end{aligned} \quad (\text{B.19})$$

And this form can be written in a very descriptive way as:

$$\mathcal{Z}_{n+1} = \exp(-Nf_0) \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\}) \quad (\text{B.20a})$$

with the *replica free energy*

$$f_{n+1}\{\Omega\} = \frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) |\Omega(\hat{q})|^2 + \frac{1}{2} \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) |\Omega(\hat{q})|^2 - \frac{1}{\mathcal{N}} \ln \mathfrak{z}^{\mathcal{N}}, \quad (\text{B.20b})$$

and an *effective one-particle partition function*

$$\mathfrak{z} = \int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}} + i \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}}\right). \quad (\text{B.20c})$$

B.2.2 Meaning of the Order Parameter $\Omega(\hat{q})$

In the basic form of the Hubbard-Stratonovich transformation with plus sign, Eq. (B.12a), one can calculate how the two variables w and z are connected. To see

this, we have a look at the expectation value:

$$\begin{aligned}
\langle |w - z|^2 \rangle_+ &:= \frac{\int dz |w - z|^2 \exp(-c|z|^2 + 2c \operatorname{Re}(zw^*))}{\int dz \exp(-c|z|^2 + 2c \operatorname{Re}(zw^*))} \\
&= \frac{\int dz |w - z|^2 \exp(-c|z|^2 + 2c \operatorname{Re}(zw^*))}{\frac{\pi}{c} \exp(c|w|^2)} \\
&= \frac{c}{\pi} \int dz |w - z|^2 \exp(-c|w - z|^2)
\end{aligned} \tag{B.21}$$

where we used $|w - z|^2 = |z|^2 + |w|^2 - 2 \operatorname{Re}(zw^*)$. Then:

$$\begin{aligned}
\langle |w - z|^2 \rangle_+ &= \frac{c}{\pi} \left(-\frac{\partial}{\partial c} \right) \int dz \exp(-c|w - z|^2) = -\frac{c}{\pi} \frac{\partial}{\partial c} \frac{\pi}{c} \\
&= \frac{1}{c}
\end{aligned} \tag{B.22}$$

Using this relation for the transformation (B.14a), we get with $c \propto N$:

$$\langle |O(\hat{q}) - \Omega(\hat{q})|^2 \rangle_{\Omega(\hat{q})} \propto \frac{1}{N} \quad \text{for all } \hat{q} \in \text{HRS}, \tag{B.23}$$

for an arbitrary particle configuration $\{\hat{r}_j\}$. The angled brackets $\langle \cdot \rangle_{\Omega(\hat{q})}$ mean averaging over all $\Omega(\hat{q})$ with the statistical weight as in Eq. (B.14a). If we additionally average Eq. (B.23) over all particle configurations $\{\hat{r}_j\}$, in analogy to the step from Eq. (B.14) to Eq. (B.15), we immediately get:

$$\boxed{\langle |O(\hat{q}) - \Omega(\hat{q})|^2 \rangle_{f_{n+1}} \propto \frac{1}{N} \quad \text{for all } \hat{q} \in \text{HRS},} \tag{B.24}$$

where $\langle \cdot \rangle_{f_{n+1}}$ is now averaging with respect to the replica free energy $f_{n+1}\{\Omega\}$ (before reducing the integral $\int d\hat{r}_1 \cdots d\hat{r}_N$ to an integral over one particle). This result is very convenient: It means that the difference between $O(\hat{q})$ and $\Omega(\hat{q})$ vanishes in the thermodynamic limit and hence, both can be studied equivalently.

For $\hat{q} \in \text{1RS}$ we used the Hubbard-Stratonovich transformation with minus sign, Eq. (B.12b). For that case one can show an analogous relation:

$$\langle |iw - z|^2 \rangle_- := \frac{\int dz |iw - z|^2 \exp(-c|z|^2 + 2ic \operatorname{Re}(zw^*))}{\int dz \exp(-c|z|^2 + 2ic \operatorname{Re}(zw^*))} = \frac{1}{c}. \tag{B.25}$$

We apply this relation to the transformation (B.14b) and get:

$$\boxed{\langle |iO(\hat{q}) - \Omega(\hat{q})|^2 \rangle_{f_{n+1}} \propto \frac{1}{N} \quad \text{for all } \hat{q} \in \text{1RS},} \tag{B.26}$$

which means that in the 1RS, $\Omega(\hat{q})$ and $iO(\hat{q})$ are the same in the thermodynamic limit.

B.3 Expansion of the replica free energy to 3rd Order in Ω

Here, we want to expand the replica free energy, Eq. (2.28), to the lowest necessary order, which is $\mathcal{O}(\Omega^3)$. As one can easily see, the expansion of \mathfrak{z} , Eq. (2.28b), yields

$$\begin{aligned}
\mathfrak{z} &= \int D\hat{r} \left(1 + \phi^n \mu^2 \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}) e^{-i\hat{q}\hat{r}} \right. \\
&\quad + \frac{1}{2!} \phi^{2n} \mu^4 \sum_{\hat{q}_1, \hat{q}_2 \in \text{HRS}} \Delta(\hat{q}_1) \Delta(\hat{q}_2) \Omega(\hat{q}_1) \Omega(\hat{q}_2) e^{-i(\hat{q}_1 + \hat{q}_2)\hat{r}} \\
&\quad \left. + \frac{1}{3!} \phi^{3n} \mu^6 \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Delta(\hat{q}_1) \Delta(\hat{q}_2) \Delta(\hat{q}_3) \Omega(\hat{q}_1) \Omega(\hat{q}_2) \Omega(\hat{q}_3) e^{-i(\hat{q}_1 + \hat{q}_2 + \hat{q}_3)\hat{r}} \right) \\
&= 1 + \frac{\phi^{2n} \mu^4}{2!} \sum_{\hat{q}_1, \hat{q}_2 \in \text{HRS}} \Delta(\hat{q}_1) \Delta(\hat{q}_2) \Omega(\hat{q}_1) \Omega(\hat{q}_2) \delta_{\hat{q}_1 + \hat{q}_2, \hat{0}} \\
&\quad + \frac{\phi^{3n} \mu^6}{3!} \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Delta(\hat{q}_1) \Delta(\hat{q}_2) \Delta(\hat{q}_3) \Omega(\hat{q}_1) \Omega(\hat{q}_2) \Omega(\hat{q}_3) \delta_{\hat{q}_1 + \hat{q}_2 + \hat{q}_3, \hat{0}} + \mathcal{O}(\Omega^4), \tag{B.27}
\end{aligned}$$

and with the expansion $\Delta(\hat{q}) = \exp(-a^2 \hat{q}^2 / 2) = 1 - a^2 \hat{q}^2 / 2 + \mathcal{O}(\hat{q}^4)$, we get:

$$\begin{aligned}
\ln \mathfrak{z} &= \frac{\phi^{2n} \mu^4}{2} \sum_{\hat{q} \in \text{HRS}} \left(1 - \frac{2a^2 \hat{q}^2}{2} \right) \Omega(\hat{q}) \Omega(-\hat{q}) \\
&\quad + \frac{\phi^{3n} \mu^6}{6} \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Omega(\hat{q}_1) \Omega(\hat{q}_2) \Omega(\hat{q}_3) \delta_{\hat{q}_1 + \hat{q}_2 + \hat{q}_3, \hat{0}} + \mathcal{O}(\Omega^4, \Omega^3 \hat{q}^2, \Omega^2 \hat{q}^4). \tag{B.28}
\end{aligned}$$

With Eq. (2.28) the complete replica free energy becomes:

$$\begin{aligned}
f_{n+1}\{\Omega\} &= \frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} \left(1 - \frac{a^2 \hat{q}^2}{2} \right) |\Omega(\hat{q})|^2 - \frac{\phi^{2n} \mu^4}{2} \sum_{\hat{q} \in \text{HRS}} (1 - a^2 \hat{q}^2) \Omega(\hat{q}) \Omega(-\hat{q}) \\
&\quad - \frac{\phi^{3n} \mu^6}{6} \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Omega(\hat{q}_1) \Omega(\hat{q}_2) \Omega(\hat{q}_3) \delta_{\hat{q}_1 + \hat{q}_2 + \hat{q}_3, \hat{0}} + \mathcal{O}(\Omega^4, \Omega^3 \hat{q}^2, \Omega^2 \hat{q}^4) \tag{B.29}
\end{aligned}$$

and in the limit $n \rightarrow 0$:

$$\begin{aligned}
f_{n+1}\{\Omega\} &= \frac{1}{2} \sum_{\hat{q} \in \text{HRS}} \left(\mu^2 - \mu^4 + \left(\mu^4 - \frac{\mu^2}{2} \right) a^2 \hat{q}^2 \right) |\Omega(\hat{q})|^2 \\
&\quad - \mu^6 \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Omega(\hat{q}_1) \Omega(\hat{q}_2) \Omega(\hat{q}_3) \delta_{\hat{q}_1 + \hat{q}_2 + \hat{q}_3, \hat{0}} + \mathcal{O}(\Omega^4, \Omega^3 \hat{q}^2, \Omega^2 \hat{q}^4), \tag{B.30}
\end{aligned}$$

or

$$\begin{aligned}
 f_{n+1}\{\Omega\} &= \sum_{\hat{q} \in \text{HRS}} \left(-A\varepsilon + \frac{B}{2} \hat{q}^2 \right) |\Omega(\hat{q})|^2 \\
 &\quad - C \sum_{\hat{q}_1, \hat{q}_2, \hat{q}_3 \in \text{HRS}} \Omega(\hat{q}_1) \Omega(\hat{q}_2) \Omega(\hat{q}_3) \delta_{\hat{q}_1 + \hat{q}_2 + \hat{q}_3, \hat{0}} + \mathcal{O}(\Omega^4, \Omega^3 \hat{q}^2, \Omega^2 \hat{q}^4)
 \end{aligned} \tag{B.31}$$

with

$$\varepsilon = \mu^2 - 1, \quad A = \frac{\mu^2}{2}, \quad B = \left(\mu^4 - \frac{\mu^2}{2} \right) a^2, \quad C = \frac{\mu^6}{6}. \tag{B.32}$$

A discussion of the *universal* free energy, Eq. (B.31), is found in Sec. 2.4.5.

B.4 Fluctuations around the Saddle-Point Solution

In Sec. 2.6.2, we derived an Ansatz for the order parameter, Eqs. (2.45):

$$\bar{\Omega}_u(\hat{q}) = (1 - Q) \delta_{\hat{q}, \hat{0}} + Q \omega(\hat{q}) \int D\mathbf{x}_{\parallel} \exp(i\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + iq_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) \tag{B.33}$$

Let us have a look at the integral over \mathbf{x}_{\parallel} of the gel part (marked in blue). Without deformations, $u_{\perp}(\mathbf{x}_{\parallel}) \equiv \hat{0}$, it can be performed and yields a δ -function. For this case, the Ansatz reproduces the saddle point solution:

$$\bar{\Omega}_{u \equiv 0}(\hat{q}) = (1 - Q) \delta_{\hat{q}, \hat{0}} + Q \omega(\hat{q}) \delta_{\mathbf{q}_{\parallel}, \mathbf{0}} = \bar{\Omega}_{\text{sp}}(\hat{q}) \tag{B.34}$$

where $\bar{\Omega}_{\text{sp}}(\hat{q})$ is the Fourier representation of Eq. (2.33).

In the limit of small deformations u_{\perp} and long wavelengths, i.e. small \hat{q} , we can expand the exponential to the lowest order in $q_{\perp} u_{\perp}$:

$$\begin{aligned}
 &\int D\mathbf{x}_{\parallel} \exp(i\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + iq_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) \\
 &= \int D\mathbf{x}_{\parallel} \exp(i\mathbf{q}_{\parallel} \mathbf{x}_{\parallel}) (1 + iq_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) \\
 &= \delta_{\mathbf{q}_{\parallel}, \mathbf{0}} + iq_{\perp} u_{\perp}(\mathbf{q}_{\parallel}),
 \end{aligned} \tag{B.35}$$

with the Fourier transform $u_{\perp}(\mathbf{q}_{\parallel})$ of $u_{\perp}(\mathbf{x}_{\parallel})$. That means, we can write the Ansatz (B.33) as:

$$\bar{\Omega}_u(\hat{q}) = \bar{\Omega}_{\text{sp}}(\hat{q}) + \delta\Omega_u(\hat{q}) \tag{B.36a}$$

$$\text{with } \delta\Omega_u(\hat{q}) = \mathcal{O}(q_{\perp} u_{\perp}), \tag{B.36b}$$

as one can see by comparing Eqs. (B.34) and (B.33), with (B.35) inserted.

B.5 Replica Free Energy with Shear Deformations

In this section we will insert the Ansatz (2.45)

$$\bar{\Omega}_u(\hat{q}) = (1 - Q)\delta_{\hat{q},\hat{0}} + Q\omega(\hat{q}) \int D\mathbf{x}_{\parallel} \exp(\mathbf{i}q_{\parallel}\mathbf{x}_{\parallel} + \mathbf{i}q_{\perp}u_{\perp}(\mathbf{x}_{\parallel})) \quad (\text{B.37a})$$

$$\text{with } \omega(\hat{q}) = \int_0^{\infty} d\xi^2 \mathcal{P}(\xi^2) \exp\left(-\frac{\hat{q}^2\xi^2}{2}\right) \quad (\text{B.37b})$$

into the general expression (2.28) for the replica free energy. We will only consider small, long-wavelength deformations and keep u_{\perp} only to the lowest order, $(\partial u_{\perp}/\partial \mathbf{x}_{\parallel})^2$.

B.5.1 Preparing the Replica Free Energy for the Ansatz

The derived Ansatz for the order parameter (B.37) suggests to define a slightly altered version of the order parameter

$$\Omega'(\hat{q}) := \Omega(\hat{q}) - (1 - Q)\delta_{\hat{q},\hat{0}} \quad (\text{B.38})$$

so that the Ansatz for this version of the order parameter,

$$\boxed{\bar{\Omega}'_u(\hat{q}) = Q\omega(\hat{q}) \int D\mathbf{x}_{\parallel} \exp(\mathbf{i}q_{\parallel}\mathbf{x}_{\parallel} + \mathbf{i}q_{\perp}u_{\perp}(\mathbf{x}_{\parallel}))} \quad (\text{B.39})$$

does not involve a δ -function and hence becomes easier to handle when summing over \hat{q} .

When inserting the Ansatz $\bar{\Omega}_u$ into the general replica free energy $f_{n+1}\{\Omega\}$, Eq. (2.28), we note that $f_{n+1}\{\Omega\}$ only involves sums over the higher replica sector (which do not include $\hat{q} = \hat{0}$), so we can just swap $\bar{\Omega}_u(\hat{q})$ with $\bar{\Omega}'_u(\hat{q})$:

$$f_{n+1}\{\bar{\Omega}_u\} = \frac{\phi^n \mu^2}{2} \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) |\bar{\Omega}'_u(\hat{q})|^2 - \ln \mathfrak{z} \quad (\text{B.40a})$$

$$\text{with } \mathfrak{z} = \int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-\mathbf{i}\hat{q}\hat{r}}\right) \quad (\text{B.40b})$$

To simplify further calculations, we include the 1RS in the sums. We do not need correction terms, since volume conserving deformations keep a constant density and yield $\bar{\Omega}'_u(\hat{q}) = 0$ for $\hat{q} \in \text{1RS}$, as we show in Appendix B.7. Furthermore, we also include $\hat{q} = \hat{0}$ in the sums and subtract the generated terms (involving $\bar{\Omega}'_u(\hat{0}) = Q$) again. With that we can omit the unpleasant constraint $\hat{q} \in \text{HRS}$ and sum over

all \hat{q} :

$$f_{n+1}\{\bar{\Omega}_u\} = -\frac{\phi^n \mu^2}{2} Q^2 + \frac{\phi^n \mu^2}{2} \sum_{\hat{q}} \Delta(\hat{q}) |\bar{\Omega}'_u(\hat{q})|^2 - \ln \mathfrak{z}, \quad (\text{B.41a})$$

$$\mathfrak{z} = \exp(-\phi^n \mu^2 Q) \int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}}\right) \quad (\text{B.41b})$$

B.5.2 Expansion of the one-particle partition function \mathfrak{z}

Now we expand the exponential in the integral of the one-particle partition function \mathfrak{z} :

$$\begin{aligned} \frac{\mathfrak{z}}{\exp(-\phi^n \mu^2 Q)} &= \int D\hat{r} \exp\left(\phi^n \mu^2 \sum_{\hat{q}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}}\right) \\ &= \int D\hat{r} \sum_{r=0}^{\infty} \frac{1}{r!} \left\{ \phi^n \mu^2 \sum_{\hat{q}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}} \right\}^r \\ &= \sum_{r=0}^{\infty} \frac{(\phi^n \mu^2)^r}{r!} \int D\hat{r} \left\{ \sum_{\hat{q}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}} \right\}^r \\ &= \sum_{r=0}^{\infty} \frac{(\phi^n \mu^2)^r}{r!} A_r \end{aligned} \quad (\text{B.42})$$

with

$$A_r := \int_{V^{n+1}} D\hat{r} \left\{ \sum_{\hat{q}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}} \right\}^r. \quad (\text{B.43})$$

which we now try to calculate.

B.5.3 Insertion of the Order Parameter in \mathfrak{z}

The term A_0 is very easy:

$$A_0 = \int_{V^{n+1}} D\hat{r} \left\{ \dots \right\}^0 = 1 \quad (\text{B.44})$$

For A_1 we note that $\int D\hat{r} e^{-i\hat{q}\hat{r}} = \delta_{\hat{q},\hat{0}}$ and thus:

$$A_1 = \int D\hat{r} \sum_{\hat{q}} \Delta(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}} = \Delta(\hat{0}) \bar{\Omega}'_u(\hat{0}) = Q \quad (\text{B.45})$$

Now, let us consider the ‘‘difficult’’ case $r \geq 2$ and plug in the Ansatz (B.39):

$$\begin{aligned} A_r &= \frac{Q^r}{V^{r(n+1)Dr/2}} \int D\hat{r} \left\{ \sum_{\hat{q}} \int d\mathbf{x}_{\parallel} \int_0^{\infty} d\xi^2 \mathcal{P}(\xi^2) \right. \\ &\quad \left. \exp\left(-\frac{a^2 \hat{q}^2}{2}\right) \exp\left(-\frac{\xi^2 \hat{q}^2}{2}\right) \exp\left(i(\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + q_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) - i\hat{q}\hat{r}\right) \right\}^r \end{aligned} \quad (\text{B.46})$$

Let us have a closer look at the sum $\sum_{\hat{q}}$ in the first line: It runs over elements $\hat{q} = \frac{2\pi}{L} \cdot \hat{n}$ with $\hat{n} \in \mathbb{Z}^{D(n+1)}$ and $L = V^{1/D}$ being the linear dimension of the system. In the thermodynamic limit $L \rightarrow \infty$ the discrete values of this sum move closer and closer and the sum approaches a Riemann integral

$$\sum_{\hat{q}}(\dots) \longrightarrow \left(\frac{L}{2\pi}\right)^{D(n+1)} \int d\hat{q}(\dots) \quad (\text{B.47})$$

To simplify the notation we also define a $D(n+1)$ -dimensional hatted function $\hat{v}(\mathbf{x}_{\parallel})$ such that

$$\mathbf{v}_{\parallel}(\mathbf{x}_{\parallel}) = \mathbf{x}_{\parallel} \quad \text{and} \quad v_{\perp}(\mathbf{x}_{\parallel}) = u_{\perp}(\mathbf{x}_{\parallel}) \quad (\text{B.48})$$

so that

$$\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + q_{\perp} u_{\perp}(\mathbf{x}_{\parallel}) = \hat{q} \hat{v}(\mathbf{x}_{\parallel}), \quad (\text{B.49})$$

and with that we get:

$$\begin{aligned} A_r &= \frac{Q^r}{V^{r(n+1)Dr/2}} \frac{V^{r(n+1)}}{(2\pi)^{Dr(n+1)}} \int D\hat{r} \\ &\quad \left\{ \int d\hat{q} \int d\mathbf{x}_{\parallel} \int_0^{\infty} d\xi^2 \mathcal{P}(\xi^2) \exp\left(-\frac{(a^2 + \xi^2)\hat{q}^2}{2}\right) \exp\left(i\hat{q}\hat{v}(\mathbf{x}_{\parallel}) - i\hat{q}\hat{r}\right) \right\}^r \end{aligned} \quad (\text{B.50})$$

The integral over \hat{q} is a simple Gaussian integral, which we perform:

$$\begin{aligned} A_r &/ \left(\frac{Q^r V^{rn}}{(2\pi)^{Dr(n+1)} (n+1)^{Dr/2}} \right) = \\ &= \int D\hat{r} \left\{ \int d\mathbf{x}_{\parallel} \int_0^{\infty} d\xi^2 \mathcal{P}(\xi^2) \left(\frac{2\pi}{a^2 + \xi^2} \right)^{D(n+1)/2} \exp\left(-\frac{(\hat{v}(\mathbf{x}_{\parallel}) - \hat{r})^2}{2(a^2 + \xi^2)}\right) \right\}^r \end{aligned} \quad (\text{B.51})$$

Now we expand the r -th power:

$$\begin{aligned} A_r &/ \left(\frac{Q^r V^{rn}}{(2\pi)^{Dr(n+1)} (n+1)^{Dr/2}} \right) = \\ &= \int_0^{\infty} d\xi_1^2 \mathcal{P}(\xi_1^2) \cdots d\xi_r^2 \mathcal{P}(\xi_r^2) \int_V D\hat{r} \int d\mathbf{x}_{\parallel 1} \cdots d\mathbf{x}_{\parallel r} \\ &\quad \prod_{\gamma=1}^r \left\{ \left(\frac{2\pi}{a^2 + \xi_{\gamma}^2} \right)^{D(n+1)/2} \exp\left(-\frac{(\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{r})^2}{2(a^2 + \xi_{\gamma}^2)}\right) \right\} \end{aligned} \quad (\text{B.52})$$

To simplify the notation we furthermore define

$$\int_{\xi_1^2, \dots, \xi_r^2} := \int_0^{\infty} d\xi_1^2 \mathcal{P}(\xi_1^2) \cdots d\xi_r^2 \mathcal{P}(\xi_r^2) \quad (\text{B.53})$$

$$C_{\gamma} := \frac{1}{a^2 + \xi_{\gamma}^2} \quad \text{and} \quad C := \sum_{\gamma=1}^r C_{\gamma} \quad (\text{B.54})$$

With that, Eq. (B.52) becomes:

$$A_r / \left(\frac{Q^r V^{rn}}{(2\pi)^{Dr(n+1)} (n+1)^{Dr/2}} \right) = \int_{\xi_1^2, \dots, \xi_r^2} \left(\prod_{\gamma=1}^r (2\pi C_\gamma)^{D(n+1)/2} \right) \times \int d\mathbf{x}_{\parallel 1} \cdots d\mathbf{x}_{\parallel r} \int D\hat{r} \exp \left(- \sum_{\gamma=1}^r \frac{C_\gamma}{2} (\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{r})^2 \right) \quad (\text{B.55})$$

Now we try to perform the integral over \hat{r} and remember that $D\hat{r} = d\hat{r}/V^{n+1}$:

$$\begin{aligned} & \int D\hat{r} \exp \left(- \sum_{\gamma=1}^r \frac{C_\gamma}{2} (\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{r})^2 \right) \\ &= \int D\hat{r} \exp \left(- \sum_{\gamma=1}^r \frac{C_\gamma}{2} (\hat{v}^2(\mathbf{x}_{\parallel \gamma}) - 2\hat{r}\hat{v}(\mathbf{x}_{\parallel \gamma}) + \hat{r}^2) \right) \\ &= \exp \left(- \sum_{\gamma=1}^r \frac{C_\gamma}{2} \hat{v}^2(\mathbf{x}_{\parallel \gamma}) \right) \int D\hat{r} \exp \left(- \frac{C}{2} \hat{r}^2 + \hat{r} \sum_{\gamma=1}^r C_\gamma \hat{v}(\mathbf{x}_{\parallel \gamma}) \right) \\ &= \exp \left(- \sum_{\gamma=1}^r \frac{C_\gamma}{2} \hat{v}^2(\mathbf{x}_{\parallel \gamma}) \right) \frac{1}{V^{n+1}} \left(\frac{2\pi}{C} \right)^{D(n+1)/2} \exp \left(\frac{1}{2C} \left(\sum_{\gamma=1}^r C_\gamma \hat{v}(\mathbf{x}_{\parallel \gamma}) \right)^2 \right) \\ &= \frac{1}{V^{n+1}} \left(\frac{2\pi}{C} \right)^{D(n+1)/2} \exp \left(- \frac{1}{2C} \left\{ C \sum_{\gamma=1}^r C_\gamma \hat{v}^2(\mathbf{x}_{\parallel \gamma}) - \left(\sum_{\gamma=1}^r C_\gamma \hat{v}(\mathbf{x}_{\parallel \gamma}) \right)^2 \right\} \right) \end{aligned} \quad (\text{B.56})$$

Have a look at the term in the curly brackets { }:

$$\begin{aligned} & C \sum_{\gamma=1}^r C_\gamma \hat{v}^2(\mathbf{x}_{\parallel \gamma}) - \left(\sum_{\gamma=1}^r C_\gamma \hat{v}(\mathbf{x}_{\parallel \gamma}) \right)^2 \\ &= \sum_{\gamma, \delta=1}^r C_\gamma C_\delta \hat{v}^2(\mathbf{x}_{\parallel \gamma}) - \sum_{\gamma, \delta=1}^r C_\gamma C_\delta \hat{v}(\mathbf{x}_{\parallel \gamma}) \hat{v}(\mathbf{x}_{\parallel \delta}) \\ &= \sum_{\gamma, \delta=1}^r C_\gamma C_\delta \hat{v}(\mathbf{x}_{\parallel \gamma}) \left(\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{v}(\mathbf{x}_{\parallel \delta}) \right) \\ &= \sum_{\gamma < \delta} C_\gamma C_\delta \left(\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{v}(\mathbf{x}_{\parallel \delta}) \right) \left(\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{v}(\mathbf{x}_{\parallel \delta}) \right) \\ &= \sum_{\gamma < \delta} C_\gamma C_\delta \left(\hat{v}(\mathbf{x}_{\parallel \gamma}) - \hat{v}(\mathbf{x}_{\parallel \delta}) \right)^2 \\ &= \sum_{\gamma < \delta} C_\gamma C_\delta \left((u_\perp(\mathbf{x}_{\parallel \gamma}) - u_\perp(\mathbf{x}_{\parallel \delta}))^2 + (\mathbf{x}_{\parallel \gamma} - \mathbf{x}_{\parallel \delta})^2 \right) \end{aligned} \quad (\text{B.57})$$

Hence Eq. (B.56) becomes:

$$\int D\hat{r} \exp\left(-\sum_{\gamma=1}^r \frac{C_\gamma}{2} (\hat{v}(\mathbf{x}_{\parallel\gamma}) + \hat{r})^2\right) = \frac{1}{V^{n+1}} \left(\frac{2\pi}{C}\right)^{D(n+1)/2} \\ \times \exp\left(-\frac{1}{2C} \left\{ \sum_{\gamma<\delta} C_\gamma C_\delta \left((u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}))^2 + (\mathbf{x}_{\parallel\gamma} - \mathbf{x}_{\parallel\delta})^2 \right) \right\}\right) \quad (\text{B.58})$$

and we can plug that in Eq. (B.55) to get:

$$A_r / \left(\frac{Q^r V^{rn}}{(2\pi)^{Dr(n+1)} (n+1)^{Dr/2} V^{n+1}} \right) \\ = \int_{\xi_1^2, \dots, \xi_r^2} \left(\prod_{\gamma=1}^r (2\pi C_\gamma)^{D(n+1)/2} \right) \left(\frac{2\pi}{C} \right)^{D(n+1)/2} \\ \times \int d\mathbf{x}_{\parallel 1} \cdots d\mathbf{x}_{\parallel r} \exp\left(-\frac{1}{2C} \sum_{\gamma<\delta} C_\gamma C_\delta (\mathbf{x}_{\parallel\gamma} - \mathbf{x}_{\parallel\delta})^2\right) \\ \times \exp\left(-\frac{1}{2C} \sum_{\gamma<\delta} C_\gamma C_\delta (u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}))^2\right) \quad (\text{B.59})$$

In the last line one can see, as one would expect, that strong deformations, meaning a high $(u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}))^2$, will have a high energy cost; unless $\mathbf{x}_{\parallel\gamma}$ and $\mathbf{x}_{\parallel\delta}$ are far away from each other, then the contribution will be suppressed by the exponential in the second last line and the magnitude of $(u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}))^2$ does not matter.

As already mentioned, we are only interested in the lowest order of the deformation, $\mathcal{O}(\partial^2 u_\perp^2)$, and in this spirit we make an expansion in u_\perp :

$$u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}) \\ = u_\perp(\mathbf{x}_{\parallel\gamma}) - \left(u_\perp(\mathbf{x}_{\parallel\gamma}) + \sum_{\tau=1}^D \frac{\partial u_\perp(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\tau)}} (x_{\parallel\delta}^{(\tau)} - x_{\parallel\gamma}^{(\tau)}) \right) + \mathcal{O}(\partial^2 u_\perp) \\ = \sum_{\tau=1}^D \frac{\partial u_\perp(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\tau)}} (x_{\parallel\gamma}^{(\tau)} - x_{\parallel\delta}^{(\tau)}) + \mathcal{O}(\partial^2 u_\perp) \quad (\text{B.60})$$

where $x_{\parallel\gamma}^{(\tau)}$ is the τ -th spatial component of $\mathbf{x}_{\parallel\gamma}$. We furthermore define $u_\perp^{(\alpha,\nu)}(\mathbf{x}_{\parallel\gamma})$ as the ν -th spatial component in replica α of $u_\perp(\mathbf{x}_{\parallel\gamma})$ and obtain:

$$(u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}))^2 \\ = \sum_{\alpha,\nu} \sum_{\tau,\sigma} \frac{\partial u_\perp^{(\alpha,\nu)}(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\tau)}} \frac{\partial u_\perp^{(\alpha,\nu)}(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\sigma)}} (x_{\parallel\gamma}^{(\tau)} - x_{\parallel\delta}^{(\tau)}) (x_{\parallel\gamma}^{(\sigma)} - x_{\parallel\delta}^{(\sigma)}) + \mathcal{O}(\partial^4 u_\perp^2) \quad (\text{B.61})$$

and therefore:

$$\begin{aligned}
& \exp\left(-\frac{1}{2C} \sum_{\gamma < \delta} C_\gamma C_\delta (u_\perp(\mathbf{x}_{\parallel\gamma}) - u_\perp(\mathbf{x}_{\parallel\delta}))^2\right) \\
&= 1 - \frac{1}{2C} \sum_{\gamma < \delta} C_\gamma C_\delta \sum_{\alpha, \nu} \sum_{\tau, \sigma} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\tau)}} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\sigma)}} (x_{\parallel\gamma}^{(\tau)} - x_{\parallel\delta}^{(\tau)})(x_{\parallel\gamma}^{(\sigma)} - x_{\parallel\delta}^{(\sigma)}) \\
&\quad + \mathcal{O}(\partial^4 u_\perp^2). \tag{B.62}
\end{aligned}$$

With the expanded u_\perp and omitting the $\mathcal{O}(\partial^4 u_\perp^2)$, Eq. (B.59) becomes:

$$\begin{aligned}
& A_r \left/ \left(\frac{Q^r V^{(r-1)n-1}}{(2\pi)^{Dr(n+1)} (n+1)^{Dr/2}} (2\pi)^{Dr(n+1)/2 + D(n+1)/2} \right) \right. \\
&= \int_{\xi_1^2, \dots, \xi_r^2} \left(\prod_{\gamma=1}^r C_\gamma \right)^{D(n+1)/2} \frac{1}{C^{D(n+1)/2}} \\
&\quad \times \int d\mathbf{x}_{\parallel 1} \cdots d\mathbf{x}_{\parallel r} \exp\left(-\frac{1}{2C} \sum_{\gamma < \delta} C_\gamma C_\delta (\mathbf{x}_{\parallel\gamma} - \mathbf{x}_{\parallel\delta})^2\right) \\
&\quad \times \left(1 - \frac{1}{2C} \sum_{\gamma < \delta} C_\gamma C_\delta \sum_{\alpha, \nu} \sum_{\tau, \sigma} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\tau)}} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel\gamma})}{\partial x_{\parallel\gamma}^{(\sigma)}} (x_{\parallel\gamma}^{(\tau)} - x_{\parallel\delta}^{(\tau)})(x_{\parallel\gamma}^{(\sigma)} - x_{\parallel\delta}^{(\sigma)}) \right) \tag{B.63}
\end{aligned}$$

When we look at the last two lines, we realize that the expression is symmetric with respect to interchanging any two γ and δ . Therefore, after performing the integral $\int d\mathbf{x}_{\parallel 1} \cdots d\mathbf{x}_{\parallel r}$, each summand of the sum $\sum_{\gamma < \delta}$ in the last line will have the same result. Since there are $r(r-1)/2$ ways to choose $1 \leq \gamma < \delta \leq r$, we can replace the sum by $r(r-1)/2$ and take $\gamma = r, \delta = 1$:

$$\begin{aligned}
& A_r \left/ \left(\frac{Q^r V^{(r-1)n-1}}{(2\pi)^{D(r-1)(n+1)/2} (n+1)^{Dr/2}} \right) \right. = \int_{\xi_1^2, \dots, \xi_r^2} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{D(n+1)/2} \\
&\quad \times \int d\mathbf{x}_{\parallel 1} \cdots d\mathbf{x}_{\parallel r} \exp\left(-\frac{1}{2C} \sum_{\gamma < \delta} C_\gamma C_\delta (\mathbf{x}_{\parallel\gamma} - \mathbf{x}_{\parallel\delta})^2\right) \\
&\quad \times \left(1 - \frac{r(r-1)}{2} \frac{C_r C_1}{2C} \sum_{\alpha, \nu} \sum_{\tau, \sigma} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel r})}{\partial x_{\parallel r}^{(\tau)}} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel r})}{\partial x_{\parallel r}^{(\sigma)}} (x_{\parallel r}^{(\tau)} - x_{\parallel 1}^{(\tau)})(x_{\parallel r}^{(\sigma)} - x_{\parallel 1}^{(\sigma)}) \right) \tag{B.64}
\end{aligned}$$

It is clear that we will not be able to perform the integral over $\mathbf{x}_{\parallel r}$, because we do not know the actual form of $u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel r})$ at this point. However, for $\mathbf{x}_{\parallel 1}, \dots, \mathbf{x}_{\parallel (r-1)}$ we can try to perform the integral. Let us first make the substitution $\mathbf{y}_{\parallel\gamma} = \mathbf{x}_{\parallel\gamma} - \mathbf{x}_{\parallel r}$ for $\gamma < r$ and make the simplifying definition

$$U_{\tau\sigma}(\mathbf{x}_{\parallel r}) := \frac{r(r-1)}{2} \frac{C_r C_1}{2C} \sum_{\alpha, \nu} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel r})}{\partial x_{\parallel r}^{(\tau)}} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_{\parallel r})}{\partial x_{\parallel r}^{(\sigma)}}. \tag{B.65}$$

Hence we get:

$$\begin{aligned} A_r & \Big/ \left(\frac{Q^r V^{(r-1)n-1}}{(2\pi)^{D(r-1)(n+1)/2} (n+1)^{Dr/2}} \right) \\ & = \int_{\xi_1^2, \dots, \xi_r^2} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{D(n+1)/2} \int d\mathbf{x}_{\parallel r} B_r(\mathbf{x}_{\parallel r}) \end{aligned} \quad (\text{B.66a})$$

with

$$\begin{aligned} B_r(\mathbf{x}_{\parallel r}) & = \int d\mathbf{y}_{\parallel 1} \cdots d\mathbf{y}_{\parallel (r-1)} \exp \left(-\frac{1}{2C} \left(\sum_{\gamma < \delta < r} C_\gamma C_\delta (\mathbf{y}_{\parallel \gamma} - \mathbf{y}_{\parallel \delta})^2 + \sum_{\gamma < r} C_\gamma C_r \mathbf{y}_{\parallel \gamma}^2 \right) \right) \\ & \quad \times \left(1 - \sum_{\tau, \sigma} U_{\tau\sigma}(\mathbf{x}_{\parallel r}) y_{\parallel 1}^{(\tau)} y_{\parallel 1}^{(\sigma)} \right). \end{aligned} \quad (\text{B.66b})$$

which we try to calculate now. Let us have a look at the sum $\sum_{\tau, \sigma}$ in the last line. For $\tau \neq \sigma$, integration over $\mathbf{y}_{\parallel 1}$ yields zero due to point symmetry of the integrand.

To perform the integral over $\mathbf{y}_{\parallel 1}, \dots, \mathbf{y}_{\parallel (r-1)}$, we can write the sums in the exponential such that it takes the form of a well known Gaussian integral:

$$\begin{aligned} T & = \sum_{\gamma < \delta < r} C_\gamma C_\delta (\mathbf{y}_{\parallel \gamma} - \mathbf{y}_{\parallel \delta})^2 + \sum_{\gamma < r} C_\gamma C_r \mathbf{y}_{\parallel \gamma}^2 \\ & = \sum_{\gamma < \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma}^2 - 2 \sum_{\gamma < \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma} \mathbf{y}_{\parallel \delta} + \sum_{\gamma < \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \delta}^2 + \sum_{\gamma < r} C_\gamma C_r \mathbf{y}_{\parallel \gamma}^2 \end{aligned} \quad (\text{B.67})$$

The first and third term can be combined to one sum over $\gamma \neq \delta$. The second sum can also be changed to a sum over $\gamma \neq \delta$, eliminating the factor of 2:

$$T = \sum_{\substack{\gamma \neq \delta \\ \gamma, \delta < r}} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma}^2 - \sum_{\substack{\gamma \neq \delta \\ \gamma, \delta < r}} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma} \mathbf{y}_{\parallel \delta} + C_r \sum_{\gamma < r} C_\gamma \mathbf{y}_{\parallel \gamma}^2 \quad (\text{B.68})$$

We can also include the term $\gamma = \delta$ in the first and second sum. Since they make the same contribution in both sums, with opposite sign, they cancel out exactly.

$$\begin{aligned} T & = \sum_{\gamma, \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma}^2 - \sum_{\gamma, \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma} \mathbf{y}_{\parallel \delta} + C_r \sum_{\gamma < r} C_\gamma \mathbf{y}_{\parallel \gamma}^2 \\ & = (C - C_r) \sum_{\gamma < r} C_\gamma \mathbf{y}_{\parallel \gamma}^2 - \sum_{\gamma, \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma} \mathbf{y}_{\parallel \delta} + C_r \sum_{\gamma < r} C_\gamma \mathbf{y}_{\parallel \gamma}^2 \\ & = C \sum_{\gamma < r} C_\gamma \mathbf{y}_{\parallel \gamma}^2 - \sum_{\gamma, \delta < r} C_\gamma C_\delta \mathbf{y}_{\parallel \gamma} \mathbf{y}_{\parallel \delta} \end{aligned} \quad (\text{B.69})$$

Hence we can write Eq. (B.66b) as:

$$B_r(\mathbf{x}_{\parallel r}) = \int d\mathbf{y}_{\parallel 1} \cdots \mathbf{y}_{\parallel(r-1)} \exp\left(-\sum_{\gamma, \delta < r} M_{\gamma\delta} \mathbf{y}_{\parallel\gamma} \mathbf{y}_{\parallel\delta}\right) \left(1 - \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) (y_{\parallel 1}^{(\tau)})^2\right) \quad (\text{B.70a})$$

$$\text{with a Matrix} \quad M_{\gamma\delta} = \frac{\delta_{\gamma\delta} C C_{\gamma} - C_{\gamma} C_{\delta}}{2C} = \delta_{\gamma\delta} \frac{C_{\gamma}}{2} - \frac{C_{\gamma} C_{\delta}}{2C} \quad (\text{B.70b})$$

To perform the integral, we introduce a D -dimensional vector $\mathbf{b}_{\parallel} = (b_{\parallel}^{(1)}, \dots, b_{\parallel}^{(D)})^T$ coupling to $\mathbf{y}_{\parallel 1}$ and rewrite that expression as:

$$B_r(\mathbf{x}_{\parallel r}) = \lim_{\mathbf{b}_{\parallel} \rightarrow \mathbf{0}} \left(1 - \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) \left(\frac{\partial}{\partial b_{\parallel}^{(\tau)}}\right)^2\right) \int d\mathbf{y}_{\parallel 1} \cdots \mathbf{y}_{\parallel(r-1)} \exp\left(-\sum_{\gamma, \delta < r} M_{\gamma\delta} \mathbf{y}_{\parallel\gamma} \mathbf{y}_{\parallel\delta} + \mathbf{b}_{\parallel} \mathbf{y}_{\parallel 1}\right) \quad (\text{B.71})$$

Now, the Gaussian in the second line can easily be calculated:

$$\begin{aligned} & \int d\mathbf{y}_{\parallel 1} \cdots d\mathbf{y}_{\parallel(r-1)} \exp\left(-\sum_{\gamma, \delta < r} M_{\gamma\delta} \mathbf{y}_{\parallel\gamma} \mathbf{y}_{\parallel\delta} + \mathbf{b}_{\parallel} \mathbf{y}_{\parallel 1}\right) \\ &= \frac{\pi^{D(r-1)/2}}{(\det M)^{D/2}} \exp\left(\frac{1}{4} \mathbf{b}_{\parallel}^2 M_{11}^{-1}\right), \end{aligned} \quad (\text{B.72})$$

with the determinant $\det M$ and inverse M^{-1} of the matrix M (and M_{11}^{-1} being the entry at position (1, 1) of M^{-1}). With that, Eq. (B.71) becomes:

$$\begin{aligned} B_r(\mathbf{x}_{\parallel r}) &= \frac{\pi^{D(r-1)/2}}{(\det M)^{D/2}} \lim_{\mathbf{b}_{\parallel} \rightarrow \mathbf{0}} \left(1 - \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) \left(\frac{\partial}{\partial b_{\parallel}^{(\tau)}}\right)^2\right) \exp\left(\frac{1}{4} \mathbf{b}_{\parallel}^2 M_{11}^{-1}\right) \\ &= \frac{\pi^{D(r-1)/2}}{(\det M)^{D/2}} \lim_{\mathbf{b}_{\parallel} \rightarrow \mathbf{0}} \left(1 - \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) \frac{\partial}{\partial b_{\parallel}^{(\tau)}} \frac{2}{4} b_{\parallel}^{(\tau)} M_{11}^{-1} \exp\left(\frac{1}{4} \mathbf{b}_{\parallel}^2 M_{11}^{-1}\right)\right) \\ &= \frac{\pi^{D(r-1)/2}}{(\det M)^{D/2}} \left(1 - \frac{M_{11}^{-1}}{2} \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r})\right) \end{aligned} \quad (\text{B.73})$$

To calculate the determinant and inverse of M , we take advantage of its form $M = A + \mathbf{xy}^T$ (i.e. $M_{\gamma\delta} = A_{\gamma\delta} + x_{\gamma} y_{\delta}$) with a “simple” matrix A , and use the relations [Brookes, 2009]:

$$\det(A + \mathbf{xy}^T) = \det(A)(1 + \mathbf{y}^T A^{-1} \mathbf{x}), \quad (\text{B.74a})$$

$$(A + \mathbf{xy}^T)^{-1} = A^{-1} - \frac{A^{-1} \mathbf{xy}^T A^{-1}}{1 + \mathbf{y}^T A^{-1} \mathbf{x}}. \quad (\text{B.74b})$$

In our case, we choose:

$$A_{\gamma\delta} = \delta_{\gamma\delta} \frac{C_\gamma}{2}, \quad (\text{B.75a})$$

$$x_\gamma = -\frac{C_\gamma}{2C} \quad \text{and} \quad y_\delta = C_\delta, \quad (\text{B.75b})$$

with $1 \leq \gamma, \delta \leq r-1$. Applying relation (B.74a), we easily find:

$$\begin{aligned} \det M &= \frac{C_1 \cdots C_{r-1}}{2^{r-1}} \left(1 + \sum_{\gamma=1}^{r-1} \frac{-C_\gamma}{2C} \frac{2}{C_\gamma} C_\gamma \right) \\ &= \frac{C_1 \cdots C_{r-1}}{2^{r-1}} \left(1 - \frac{1}{C} (C - C_r) \right) = \frac{1}{2^{r-1} C} \prod_{\gamma=1}^r C_\gamma \end{aligned} \quad (\text{B.76a})$$

And by applying (B.74b) one gets (after a similar calculation, which is left to the eager reader(s)):

$$M_{\gamma\delta}^{-1} = \frac{2\delta_{\gamma\delta}}{C_\gamma} - \frac{2}{C_r} \quad \text{and hence} \quad M_{11}^{-1} = \frac{2(C_1 + C_r)}{C_1 C_r} \quad (\text{B.76b})$$

Now we can plug these two results (B.76) into Eq. (B.73) and get the result for $B_r(\mathbf{x}_{\parallel r})$:

$$\begin{aligned} B_r(\mathbf{x}_{\parallel r}) &= \frac{\pi^{D(r-1)/2}}{(\det M)^{D/2}} \left(1 - \frac{M_{11}^{-1}}{2} \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) \right) \\ &= \pi^{D(r-1)/2} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{2^{r-1} C} \right)^{-D/2} \left(1 - \frac{C_1 + C_r}{C_1 C_r} \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) \right) \end{aligned} \quad (\text{B.77})$$

To get A_r , we can now plug this result for $B_r(\mathbf{x}_{\parallel r})$ back into Eq. (B.66a):

$$\begin{aligned} A_r &/ \left(\frac{Q^r V^{(r-1)n-1}}{(2\pi)^{D(r-1)(n+1)/2} (n+1)^{Dr/2}} (2\pi)^{D(r-1)/2} \right) \\ &= \int_{\xi_1^2, \dots, \xi_r^2} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{D(n+1)/2 - D/2} \int d\mathbf{x}_{\parallel r} \left(1 - \frac{C_1 + C_r}{C_1 C_r} \sum_{\tau=1}^D U_{\tau\tau}(\mathbf{x}_{\parallel r}) \right) \end{aligned} \quad (\text{B.78})$$

To simplify the notation, we can change the integration variable $\mathbf{x}_{\parallel r}$ to just \mathbf{x}_{\parallel} and insert the definition of $U_{\tau\tau}(\mathbf{x}_{\parallel})$, see Eq. (B.65):

$$\begin{aligned} A_r &/ \left(\frac{Q^r V^{(r-1)n-1}}{(2\pi)^{D(r-1)n/2} (n+1)^{Dr/2}} \right) = \int_{\xi_1^2, \dots, \xi_r^2} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{Dn/2} \\ &\times \int d\mathbf{x}_{\parallel} \left(1 - \frac{C_1 + C_r}{C_1 C_r} \cdot r(r-1) \frac{C_r C_1}{4C} \sum_{\tau} \sum_{\alpha, \nu} \frac{\partial u_{\perp}^{(\alpha, \nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \frac{\partial u_{\perp}^{(\alpha, \nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right) \end{aligned} \quad (\text{B.79})$$

After canceling $C_1 C_r$, we realize that we can arbitrarily swap C_1 and C_r in the last line with any other C_γ , because the rest of the expression is symmetric with respect to C_γ . So we can even change $C_1 \rightarrow (C_1 + \dots + C_r)/r = C/r$ and also $C_r \rightarrow C/r$. Doing that, we get:

$$A_r \left/ \left(\frac{Q^r V^{(r-1)n-1}}{(2\pi)^{D(r-1)n/2} (n+1)^{Dr/2}} \right) \right. = \int_{\xi_1^2, \dots, \xi_r^2} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{Dn/2} \\ \times \int d\mathbf{x}_\parallel \left(1 - \frac{2C}{r} \cdot r(r-1) \frac{1}{4C} \sum_\tau \sum_{\alpha, \nu} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right). \quad (\text{B.80})$$

Finally we plug in definitions (B.53) and (B.54) and end up with the result for A_r (but keep in mind that it is exact only to lowest order in the displacement field u_\perp and its derivative):

$$A_r = Q^r \frac{1}{(n+1)^{D(r-1)/2}} \left(\frac{V}{(2\pi)^{D/2}} \right)^{(r-1)n} \\ \times \int_0^\infty d\xi_1^2 \mathcal{P}(\xi_1^2) \dots d\xi_r^2 \mathcal{P}(\xi_r^2) \left(\frac{\prod_{\gamma=1}^r \frac{1}{a^2 + \xi_\gamma^2}}{\sum_{\gamma=1}^r \frac{1}{a^2 + \xi_\gamma^2}} \right)^{Dn/2} \\ \times \left(1 - \frac{r-1}{2} \int D\mathbf{x}_\parallel \sum_\alpha \sum_{\tau, \nu} \left(\frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right)^2 \right) \quad (\text{B.81})$$

$$\text{with} \quad D\mathbf{x}_\parallel = \frac{d\mathbf{x}_\parallel}{V(n+1)^{D/2}}. \quad (\text{B.82})$$

B.5.4 Insertion of the Order Parameter in $f_{n+1}\{\Omega\}$

In the previous section we calculated the one-particle partition function \mathfrak{z} , using Ansatz (B.39) for the order parameter. To get the full replica free energy, we still have to insert the Ansatz into the first part of Eq. (B.41a):

$$f_{n+1}\{\bar{\Omega}'_u\} = -\frac{\phi^n \mu^2 Q^2}{2} + \frac{\phi^n \mu^2}{2} \underbrace{\sum_{\hat{q} \in \text{HRS} \setminus \{\hat{0}\}} \Delta(\hat{q}) |\bar{\Omega}'_u(\hat{q})|^2}_{=: S_2} - \ln \mathfrak{z}. \quad (\text{B.83})$$

For the cross-link term $\Delta(\hat{q})$ we note that

$$\Delta(\hat{q}) = \exp\left(-\frac{\hat{q}^2 a^2}{2}\right) = \exp\left(-\frac{\hat{q}^2 a^2/2 + (-\hat{q})^2 a^2/2}{2}\right) = \Delta_{a^2/2}(\hat{q}) \Delta_{a^2/2}(-\hat{q}). \quad (\text{B.84})$$

where the symbol $\Delta_{a^2/2}(\hat{q})$ is the same as the original $\Delta(\hat{q})$, only with cross-link strength $a^2 \rightarrow a^2/2$. Thus we can rewrite the sum

$$\begin{aligned}
S_2 &= \sum_{\hat{q} \in \text{HRSV}\{\hat{0}\}} \Delta(\hat{q}) |\bar{\Omega}'_u(\hat{q})|^2 = \sum_{\hat{q} \in \text{HRSV}\{\hat{0}\}} \Delta_{a^2/2}(\hat{q}) \Delta_{a^2/2}(-\hat{q}) \bar{\Omega}'_u(\hat{q}) \bar{\Omega}'_u(-\hat{q}) \\
&= \sum_{\substack{\hat{q}_1, \hat{q}_2 \in \\ \text{HRSV}\{\hat{0}\}}} \delta_{\hat{q}_1 + \hat{q}_2, \hat{0}} \Delta_{a^2/2}(\hat{q}_1) \Delta_{a^2/2}(\hat{q}_2) \bar{\Omega}'_u(\hat{q}_1) \bar{\Omega}'_u(\hat{q}_2) \\
&= \int D\hat{r} \sum_{\substack{\hat{q}_1, \hat{q}_2 \in \\ \text{HRSV}\{\hat{0}\}}} e^{-i(\hat{q}_1 + \hat{q}_2)\hat{r}} \Delta_{a^2/2}(\hat{q}_1) \Delta_{a^2/2}(\hat{q}_2) \bar{\Omega}'_u(\hat{q}_1) \bar{\Omega}'_u(\hat{q}_2) \\
&= \int D\hat{r} \left\{ \sum_{\hat{q} \in \text{HRSV}\{\hat{0}\}} \Delta_{a^2/2}(\hat{q}) \bar{\Omega}'_u(\hat{q}) e^{-i\hat{q}\hat{r}} \right\}^2. \tag{B.85}
\end{aligned}$$

Comparing that expression to Eq. (B.43), we note that S_2 has the same form as A_2 with $a^2 \rightarrow a^2/2$. Therefore, using the result for A_2 , Eq. (B.81), we can immediately write down the solution:

$$\boxed{
\begin{aligned}
S_2 &= \frac{Q^2}{(n+1)^{D/2}} \left(\frac{V}{(2\pi)^{D/2}} \right)^n \int_0^\infty d\xi_1^2 \mathcal{P}(\xi_1^2) d\xi_2^2 \mathcal{P}(\xi_2^2) \left(\frac{\frac{1}{a^2/2 + \xi_1^2} \cdot \frac{1}{a^2/2 + \xi_2^2}}{\frac{1}{a^2/2 + \xi_1^2} + \frac{1}{a^2/2 + \xi_2^2}} \right)^{Dn/2} \\
&\quad \times \left(1 - \frac{1}{2} \int D\mathbf{x}_\parallel \sum_\alpha \sum_{\tau, \nu} \left(\frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right)^2 \right).
\end{aligned}
} \tag{B.86}$$

B.5.5 Recomposing the Replica Free Energy

We now recompute the expanded replica free energy $f_{n+1}\{\bar{\Omega}_u\}$ from the different components we calculated throughout the section B.5: In terms of S_2 and A_r , it is given by Eqs. (B.83) and (B.42):

$$f_{n+1}\{\bar{\Omega}_u\} = -\frac{\phi^n \mu^2 Q^2}{2} + \frac{\phi^n \mu^2}{2} S_2 - \ln \mathfrak{z}. \tag{B.87a}$$

$$\mathfrak{z} = \exp(-\phi^n \mu^2 Q) \sum_{r=0}^{\infty} \frac{(\phi^n \mu^2)^r}{r!} A_r \tag{B.87b}$$

Since we want to take the limit $n \rightarrow 0$, as seen in Eq. (2.10), we only keep the various parts up to linear order in n and omit higher orders. Therefore it is necessary to keep in mind that the sum $\sum_{\alpha=0}^n (\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel) / \partial x_\parallel^{(\tau)})^2$ is already of the order $\mathcal{O}(n)$, see Eq. (2.51). We furthermore need:

$$(n+1)^{-D(r-1)/2} = 1 - n \frac{D(r-1)}{2} + \mathcal{O}(n^2) \tag{B.88}$$

$$\mathcal{A}^n = 1 + n \ln \mathcal{A} + \mathcal{O}(n^2) \tag{B.89}$$

With that, we get for $r \geq 2$:

$$\begin{aligned}
\frac{A_r}{Q^r} &= \left\{ 1 - n \frac{D(r-1)}{2} \right\} \\
&\times \int_{\xi_1^2, \dots, \xi_r^2} \left\{ 1 + n \ln \left(\left(\frac{V}{(2\pi)^{D/2}} \right)^{r-1} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{D/2} \right) \right\} \\
&\times \left\{ 1 - \frac{r-1}{2} \int \mathrm{D}\mathbf{x}_\parallel \sum_{\alpha=0}^n \sum_{\tau, \nu} \left(\frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right)^2 \right\} + \mathcal{O}(n^2) \\
&= 1 - n \frac{D(r-1)}{2} + n \underbrace{\int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\left(\frac{V}{(2\pi)^{D/2}} \right)^{r-1} \left(\frac{\prod_{\gamma=1}^r C_\gamma}{C} \right)^{D/2} \right)}_{=: \Xi_{r, a^2}} \\
&\quad - \frac{r-1}{2} \int \mathrm{D}\mathbf{x}_\parallel \sum_{\alpha=0}^n \sum_{\tau, \nu} \left(\frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right)^2 + \mathcal{O}(n^2) \tag{B.90}
\end{aligned}$$

With that expression (for $r \geq 2$) and $A_0 = 1, A_1 = Q$, we can calculate \mathfrak{z} :

$$\begin{aligned}
\frac{\mathfrak{z}}{\exp(-\phi^n \mu^2 Q)} &= \\
&= 1 + \phi^n \mu^2 Q + \sum_{r=2}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} \frac{A_r}{Q^r} \\
&= \sum_{r=0}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} + n \cdot \sum_{r=2}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} \left(-\frac{D(r-1)}{2} + \Xi_{r, a^2} \right) \\
&\quad - \sum_{r=2}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} \cdot \frac{r-1}{2} \int \mathrm{D}\mathbf{x}_\parallel \sum_{\alpha=0}^n \sum_{\tau, \nu} \left(\frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right)^2 + \mathcal{O}(n^2). \tag{B.91}
\end{aligned}$$

To perform the sum over r , we need the relations:

$$\sum_{r=0}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} = \exp(\phi^n \mu^2 Q) \tag{B.92}$$

$$\begin{aligned}
\sum_{r=2}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} (r-1) &= 1 + \sum_{r=0}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} (r-1) \\
&\stackrel{\text{Eq. (D.2)}}{=} 1 + (\phi^n \mu^2 Q - 1) \exp(\phi^n \mu^2 Q) \tag{B.93}
\end{aligned}$$

and get:

$$\begin{aligned}
\mathfrak{z} &= 1 + n \left(-\frac{D}{2} \left(\exp(-\phi^n \mu^2 Q) + \phi^n \mu^2 Q - 1 \right) + \exp(-\phi^n \mu^2 Q) \sum_{r=2}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} \Xi_{r, a^2} \right) \\
&\quad - \frac{\exp(-\phi^n \mu^2 Q) + \phi^n \mu^2 Q - 1}{2} \int \mathrm{D}\mathbf{x}_\parallel \sum_{\alpha=0}^n \sum_{\tau, \nu} \left(\frac{\partial u_\perp^{(\alpha, \nu)}(\mathbf{x}_\parallel)}{\partial x_\parallel^{(\tau)}} \right)^2 + \mathcal{O}(n^2), \tag{B.94}
\end{aligned}$$

whereby the crossed-out terms are omitted, since they make contributions of at least quadratic order in n . In terms of expansion in n , \mathfrak{z} has the simple form $\mathfrak{z} = 1 + n \cdot \text{const.} + \mathcal{O}(n^2)$. Hence the logarithm can easily be taken and is $\ln \mathfrak{z} = n \cdot \text{const.} + \mathcal{O}(n^2) = \mathfrak{z} - 1 + \mathcal{O}(n^2)$

$$\begin{aligned} \ln \mathfrak{z} = n & \left(-\frac{D}{2} \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 \right) + \exp(-\mu^2 Q) \sum_{r=2}^{\infty} \frac{(\mu^2 Q)^r}{r!} \Xi_{r,a^2} \right) \\ & - \frac{\exp(-\mu^2 Q) + \mu^2 Q - 1}{2} \int D\mathbf{x}_{\parallel} \sum_{\alpha=0}^n \sum_{\tau,\nu} \left(\frac{\partial u_{\perp}^{(\alpha,\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 + \mathcal{O}(n^2), \end{aligned} \quad (\text{B.95})$$

Analogously, we perform the expansion of S_2 up to linear order in n . Remembering that S_2 equals A_2 with $a^2 \rightarrow a^2/2$, we can use Eq. (B.90) to write down the result immediately:

$$\begin{aligned} \frac{S_2}{Q^2} = 1 + n & \left(-\frac{D}{2} + \Xi_{2,a^2/2} \right) \\ & - \frac{1}{2} \int D\mathbf{x}_{\parallel} \sum_{\alpha=0}^n \sum_{\tau,\nu} \left(\frac{\partial u_{\perp}^{(\alpha,\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 + \mathcal{O}(n^2) \end{aligned} \quad (\text{B.96})$$

Now, we are able to write down the result for the replica free energy, according to Eq. (B.87a):

$$\begin{aligned} f_{n+1}\{\bar{\Omega}_u\} = & -\frac{\phi^n \mu^2 Q^2}{2} + \frac{\phi^n \mu^2 Q^2}{2} \\ & + n \frac{\phi^n \mu^2 Q^2}{2} \left(-\frac{D}{2} + \Xi_{2,a^2/2} \right) - \frac{\phi^n \mu^2 Q^2}{2} \cdot \frac{1}{2} \int D\mathbf{x}_{\parallel} \sum_{\alpha=0}^n \sum_{\tau,\nu} \left(\frac{\partial u_{\perp}^{(\alpha,\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 \\ & - n \left(-\frac{D}{2} \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 \right) + \exp(-\mu^2 Q) \sum_{r=2}^{\infty} \frac{(\mu^2 Q)^r}{r!} \Xi_{r,a^2} \right) \\ & + \frac{\exp(-\mu^2 Q) + \mu^2 Q - 1}{2} \int D\mathbf{x}_{\parallel} \sum_{\alpha=0}^n \sum_{\tau,\nu} \left(\frac{\partial u_{\perp}^{(\alpha,\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 + \mathcal{O}(n^2) \end{aligned} \quad (\text{B.97})$$

The fact the the two constant terms in the first line cancel out is no coincidence: As already seen in Eq. (2.28), $\lim_{n \rightarrow 0} f_{n+1}\{\Omega\}$ must be zero, because the sum over the higher replica sector does not contain any elements, since the constraint that at least two components are non-zero cannot be fulfilled, if there is only one component. Thus, there cannot be a constant term and the first terms must be at least linear in n .

The above result can be written in a very concise way as:

$$\boxed{f_{n+1}\{\bar{\Omega}_u\} = f_{n+1}^{\text{sp}} + \frac{1}{N} \cdot \frac{G}{2} \int d\mathbf{x}_{\parallel} \sum_{\alpha=0}^n \sum_{\tau,\nu} \left(\frac{\partial u_{\perp}^{(\alpha,\nu)}(\mathbf{x}_{\parallel})}{\partial x_{\parallel}^{(\tau)}} \right)^2 + \mathcal{O}(n^2),} \quad (\text{B.98a})$$

whereby the first term, f_{n+1}^{sp} , is the saddle-point value of $f_{n+1}\{\Omega\}$:

$$f_{n+1}^{\text{sp}} = n \left\{ \frac{D}{2} \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) + \frac{\mu^2 Q^2}{2} \Xi_{2,a^2/2} - \exp(-\mu^2 Q) \sum_{r=2}^{\infty} \frac{(\mu^2 Q)^r}{r!} \Xi_{r,a^2} \right\}, \quad (\text{B.98b})$$

and the second term describes the cost of fluctuations (*i.e.* shear deformations) around the saddle point, with the shear modulus

$$G = \left(\exp(-\mu^2 Q) + \mu^2 Q - 1 - \frac{\mu^2 Q^2}{2} \right) n_0. \quad (\text{B.98c})$$

B.5.6 Analysis of the quantity Ξ_{r,a^2}

In this section we have to look at the term Ξ_{r,a^2} that was defined in Eq. (B.90) and still appears in the saddle point value of the replica free energy Eq. (B.98b).

$$\begin{aligned} \Xi_{r,a^2} &:= \int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\left(\frac{V}{(2\pi)^{D/2}} \right)^{r-1} \left(\frac{\prod_{\gamma=1}^r (a^2 + \xi_\gamma^2)^{-1}}{\sum_{\gamma=1}^r (a^2 + \xi_\gamma^2)^{-1}} \right)^{D/2} \right) \\ &= \int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\left(\frac{N}{(2\pi)^{D/2}} \right)^{r-1} \left(\frac{\prod_{\gamma=1}^r \frac{n_0^{2/D}}{a^2 + \xi_\gamma^2}}{\sum_{\gamma=1}^r \frac{n_0^{2/D}}{a^2 + \xi_\gamma^2}} \right)^{D/2} \right) \\ &= (r-1) \ln N - \frac{D(r-1)}{2} \ln(2\pi) \\ &\quad + \frac{D}{2} \sum_{\gamma=1}^r \int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\frac{n_0^{2/D}}{a^2 + \xi_\gamma^2} \right) - \frac{D}{2} \int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\sum_{\gamma=1}^r \frac{n_0^{2/D}}{a^2 + \xi_\gamma^2} \right) \\ &= (r-1) \ln N - \frac{D(r-1)}{2} \ln(2\pi) \\ &\quad + \frac{Dr}{2} \int_{\xi^2} \ln \left(\frac{n_0^{2/D}}{a^2 + \xi^2} \right) - \frac{D}{2} \int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\sum_{\gamma=1}^r \frac{n_0^{2/D}}{a^2 + \xi_\gamma^2} \right) \end{aligned} \quad (\text{B.99})$$

Having the thermodynamic limit in mind, we assume that all internal length scales are small compared to the system size:

$$V^{1/D} \gg \xi, a \iff N \gg n_0 \xi^D, n_0 a^D \quad (\text{B.100})$$

In this limit, only the first term of Ξ_{r,a^2} in Eq. (B.99) remains:

$$\Xi_{r,a^2} = (r-1) \ln N + \mathcal{O}(N^0). \quad (\text{B.101})$$

B.6 Auxiliary Calculations for the Results Section

B.6.1 Distribution of the Number of Cross-Links

Here we are interested in the distribution of the number M of cross-links in the system, and in particular its mean. It is possible to calculate expectation values of the form $[M(M-1)\cdots(M-J+1)] =: \mathcal{M}_J$ using the Deam-Edwards distribution Eq. (2.8) for the disorder average $[\cdot]$. In doing so, we recall the simplifying definition $\tilde{\mu}^2 = \mu^2 / (2n_0 a^D (2\pi)^{D/2})$:

$$\begin{aligned} \mathcal{M}_J &= [M(M-1)\cdots(M-J+1)] = & (B.102) \\ &= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{(\tilde{\mu}^2)^M}{M!} Z(\{i_e, j_e\}_{e=1}^M) \cdot M(M-1)\cdots(M-J+1) \end{aligned}$$

The summands for $M = 0, 1, \dots, J-1$ obviously yield zero and can be excluded from the first sum. For $M \geq J$, we can write:

$$M(M-1)\cdots(M-J+1) = \frac{M!}{(M-J)!}, \quad (B.103)$$

and therefore:

$$\mathcal{M}_J = \frac{1}{\mathcal{Z}_1} \sum_{M=J}^{\infty} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \frac{(\tilde{\mu}^2)^M}{M!} \frac{M!}{(M-J)!} Z(\{i_e, j_e\}_{e=1}^M) \quad (B.104)$$

Now we start the sum at $M = 0$ again, whereby all occurrences of M shift to $M+J$:

$$\mathcal{M}_J = \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_{M+J}, \\ j_1, \dots, j_{M+J}=1}}^N \frac{(\tilde{\mu}^2)^{M+J}}{(M+J)!} Z(\{i_e, j_e\}_{e=1}^{M+J}) \quad (B.105)$$

Now we write out the partition function explicitly:

$$\mathcal{M}_J = \frac{1}{\mathcal{Z}_1} \int \text{D}\Gamma e^{-H_{\text{ev}}(\{\mathbf{r}_j\})} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_{M+J}, \\ j_1, \dots, j_{M+J}=1}}^N \frac{(\tilde{\mu}^2)^{M+J}}{M!} \exp\left(-\frac{1}{2a^2} \sum_{e=1}^{M+J} (\mathbf{r}_{i_e} - \mathbf{r}_{j_e})^2\right) \quad (B.106)$$

where $\int D\Gamma$ denotes integration over all particle positions $\int \prod_{j=1}^N D\mathbf{r}_j$. This form can be written in a much more convenient way:

$$\begin{aligned}
\mathcal{M}_J &= \frac{1}{\mathcal{Z}_1} \int D\Gamma e^{-H_{\text{ev}}(\{\mathbf{r}_j\})} \sum_{M=0}^{\infty} \frac{1}{M!} \left(\tilde{\mu}^2 \sum_{i,j=1}^N \exp\left(-\frac{1}{2a^2}(\mathbf{r}_i - \mathbf{r}_j)^2\right) \right)^{M+J} \\
&= \frac{1}{\mathcal{Z}_1} \int D\Gamma e^{-H_{\text{ev}}(\{\mathbf{r}_j\})} \left(\tilde{\mu}^2 \sum_{i,j=1}^N \exp\left(-\frac{(\mathbf{r}_i - \mathbf{r}_j)^2}{2a^2}\right) \right)^J \\
&\quad \times \exp\left(\tilde{\mu}^2 \sum_{i,j=1}^N \exp\left(-\frac{(\mathbf{r}_i - \mathbf{r}_j)^2}{2a^2}\right) \right) \\
&= \frac{(\tilde{\mu}^2)^J}{\mathcal{Z}_1} \left(\frac{\partial}{\partial \tilde{\mu}^2} \right)^J \int D\Gamma e^{-H_{\text{ev}}(\{\mathbf{r}_j\})} \exp\left(\tilde{\mu}^2 \sum_{i,j=1}^N \exp\left(-\frac{(\mathbf{r}_i - \mathbf{r}_j)^2}{2a^2}\right) \right) \quad (\text{B.107})
\end{aligned}$$

The integral in the last line is exactly $\lim_{n \rightarrow 0} \mathcal{Z}_{n+1} = \mathcal{Z}_1$, as we can confirm with Eq. (2.14):

$$\mathcal{M}_J = \frac{(\tilde{\mu}^2)^J}{\mathcal{Z}_1} \left(\frac{\partial}{\partial \tilde{\mu}^2} \right)^J \mathcal{Z}_1, \quad (\text{B.108})$$

and since $\tilde{\mu}^2$ is proportional to μ^2 :

$$\mathcal{M}_J = \frac{(\mu^2)^J}{\mathcal{Z}_1} \left(\frac{\partial}{\partial \mu^2} \right)^J \mathcal{Z}_1 \quad (\text{B.109})$$

Calculating \mathcal{Z}_1 is very easy: We know that $\lim_{n \rightarrow 0} f_{n+1}\{\Omega\} = 0$, because it involves only sums over the higher replica sector (see Eq. (2.28)), confirmed by the result (2.53) in Sec. 2.7. Therefore we just need to look at Eq. (2.27a), in the limit $n \rightarrow 0$:

$$\mathcal{Z}_1 = \exp(-Nf_0) \times 1 \quad (\text{B.110})$$

with f_0 given in Eq. (2.19):

$$\mathcal{Z}_1 = \exp\left(-N \left(-\frac{\mu^2}{2} + \frac{\lambda n_0}{2}\right)\right) = \exp\left(\frac{N\mu^2}{2} - \frac{N\lambda n_0}{2}\right) \quad (\text{B.111})$$

It is noteworthy that the limit of very large λ seems to be an issue here. However, since \mathcal{Z}_1 is both in the numerator and the denominator in Eq. (B.109), the dependence on λ will cancel out and the limit $\lambda \rightarrow \infty$ is possible. Now we can take the derivative:

$$\left(\frac{\partial}{\partial \mu^2} \right)^J \mathcal{Z}_1 = \left(\frac{N}{2} \right)^J \mathcal{Z}_1, \quad (\text{B.112})$$

and with Eq. (B.109) we get:

$$\mathcal{M}_J = [M(M-1)\cdots(M-J+1)] = \left(\frac{\mu^2 N}{2} \right)^J \quad (\text{B.113})$$

This result finds further use in Sec. 2.9.1.

B.6.2 Shear modulus

In Sec. 2.9.3 we wrote the shear modulus as:

$$\frac{G}{n_0 k_B T} = \frac{\mu^2}{2} - 1 + (1 - Q) - \frac{\mu^2}{2}(1 - Q)^2. \quad (\text{B.114})$$

We want to find a power series for that expression, for large cross-link densities μ^2 . Therefore we need the expansion of $(1 - Q)$ and $(1 - Q)^2$. An expression for $(1 - Q)$ is given in Eq. (2.73):

$$(1 - Q) = e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k \quad (\text{B.115})$$

For $(1 - Q)^2$, we use Eq. (B.128) derived in Appendix B.6.3 for the square of the sum:

$$\begin{aligned} (1 - Q)^2 &= e^{-2\mu^2} \left(\sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k \right)^2 \\ &= e^{-2\mu^2} \sum_{k=0}^{\infty} \frac{2(k+2)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k \end{aligned} \quad (\text{B.116})$$

With that we can calculate

$$\begin{aligned} (1 - Q) - \frac{\mu^2}{2}(1 - Q)^2 &= \\ &= e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k - \frac{\mu^2 e^{-2\mu^2}}{2} \sum_{k=0}^{\infty} \frac{2(k+2)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k \\ &= e^{-\mu^2} \left(\sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k - \sum_{k=0}^{\infty} \frac{(k+2)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^{k+1} \right) \\ &= e^{-\mu^2} \left(\sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} (\mu^2 e^{-\mu^2})^k - \sum_{k=1}^{\infty} \frac{k(k+1)^{k-2}}{k!} (\mu^2 e^{-\mu^2})^k \right) \\ &= e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-1} - k(k+1)^{k-2}}{k!} (\mu^2 e^{-\mu^2})^k \\ &= e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-2} (k+1 - k)}{k!} (\mu^2 e^{-\mu^2})^k \end{aligned} \quad (\text{B.117})$$

And hence:

$$(1 - Q) - \frac{\mu^2}{2}(1 - Q)^2 = e^{-\mu^2} \sum_{k=0}^{\infty} \frac{(k+1)^{k-2}}{k!} (\mu^2 e^{-\mu^2})^k \quad (\text{B.118})$$

In Sec. 2.9.3, we plug this expression into Eq. (2.77) to obtain the expansion of the shear modulus G for $\mu^2 \gg 1$, far from the sol-gel transition.

B.6.3 Square of the expansion for $(1 - Q)$

The goal is to calculate the square of a series of the form:

$$\tilde{W}(x) := \sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} x^k. \quad (\text{B.119})$$

This function $\tilde{W}(x)$ is related to the Lambert W function by $\tilde{W}(x) = -W(-x)/x$. To find the square of $\tilde{W}(x)$, we use the relation

$$\left(\sum_{k=0}^{\infty} a_k x^k \right)^2 = \sum_{k=0}^{\infty} \sum_{k'=0}^{\infty} a_k x^k a_{k'} x^{k'} = \sum_{k=0}^{\infty} b_k x^k \quad (\text{B.120a})$$

$$\text{with } b_k = \sum_{l=0}^k a_l a_{k-l}. \quad (\text{B.120b})$$

In our case:

$$a_k = \frac{(k+1)^{k-1}}{k!} \quad (\text{B.121})$$

and therefore

$$\begin{aligned} b_k &= \sum_{l=0}^k \frac{(l+1)^{l-1}}{l!} \frac{(k-l+1)^{k-l-1}}{(k-l)!} \\ &= \frac{1}{k!} \sum_{l=0}^k \binom{k}{l} (l+1)^{l-1} (k-l+1)^{k-l-1} \end{aligned} \quad (\text{B.122})$$

The calculation of b_k is a little bit tedious, but not difficult. Let us first consider the following relation [Knuth, 1973, p. 56]:

$$(x+y)^r = x \sum_{l=0}^r \binom{r}{l} (x-lz)^{l-1} (y+lz)^{r-l} \quad \text{for } r \geq 0, x \neq 0 \quad (\text{B.123})$$

We choose $z = -1$, $x = 1$, $y = k+1$ and $r = k, k-1$ to get:

$$(k+2)^k = \sum_{l=0}^k \binom{k}{l} (1+l)^{l-1} (k-l+1)^{k-l} \quad (\text{B.124})$$

$$(k+2)^{k-1} = \sum_{l=0}^{k-1} \binom{k-1}{l} (1+l)^{l-1} (k-l+1)^{k-l-1}. \quad (\text{B.125})$$

With the help of the two above equations, we can calculate:

$$\begin{aligned} &2(k+2)^{k-1} \\ &= (k+2)^{k-1} (k+2-k) \\ &= (k+2)^k - k(k+2)^{k-1} \end{aligned}$$

$$\begin{aligned}
&= \sum_{l=0}^k \left(\frac{k!}{l!(k-l)!} (1+l)^{l-1} (k-l+1)^{k-l} \right) \\
&\quad - k \sum_{l=0}^{k-1} \left(\frac{(k-1)!}{l!(k-l-1)!} (1+l)^{l-1} (k-l+1)^{k-l-1} \right) \\
&= \sum_{l=0}^{k-1} \left(\frac{k!}{l!(k-l)!} (1+l)^{l-1} (k-l+1)^{k-l} - \frac{k!}{l!(k-l-1)!} (1+l)^{l-1} (k-l+1)^{k-l-1} \right) \\
&\quad + (1+k)^{k-1} \\
&= \sum_{l=0}^{k-1} \left(\left\{ \frac{k!}{l!(k-l)!} (1+l)^{l-1} (k-l+1)^{k-l-1} \right\} \left\{ k-l+1 - (k-l) \right\} \right) + (1+k)^{k-1} \\
&= \sum_{l=0}^{k-1} \left(\frac{k!}{l!(k-l)!} (1+l)^{l-1} (k-l+1)^{k-l-1} \right) + (1+k)^{k-1} \\
&= \sum_{l=0}^k \binom{k}{l} (1+l)^{l-1} (k-l+1)^{k-l-1} \tag{B.126}
\end{aligned}$$

Comparing that relation with b_k in Eq. (B.122), we can write down

$$b_k = \frac{2(k+2)^{k-1}}{k!}. \tag{B.127}$$

Putting b_k into Eq. (B.120b), we get the desired result:

$$\tilde{W}^2(x) = \left(\sum_{k=0}^{\infty} \frac{(k+1)^{k-1}}{k!} x^k \right)^2 = \sum_{k=0}^{\infty} \frac{2(k+2)^{k-1}}{k!} x^k \tag{B.128}$$

B.7 The Order Parameter in the One Replica Sector

In Sec. 2.6.2 we derived an Ansatz for the order parameter $\Omega(\hat{q})$ that allows shear fluctuations around the saddle point, see Eq. (2.45):

$$\bar{\Omega}_u(\hat{q}) = (1-Q)\delta_{\hat{q},\hat{0}} + Q\omega(\hat{q}) \int D\mathbf{x}_{\parallel} \exp(i\mathbf{q}_{\parallel}\mathbf{x}_{\parallel} + iq_{\perp}u_{\perp}(\mathbf{x}_{\parallel})) \tag{B.129a}$$

$$\text{with } \omega(\hat{q}) = \int_0^{\infty} d\xi^2 \mathcal{P}(\xi^2) \exp\left(-\frac{\hat{q}^2 \xi^2}{2}\right). \tag{B.129b}$$

For $\hat{q} \in 1\text{RS}$, this order parameter describes simple density fluctuations in the respective replica. Hence, if $u_{\perp}(\mathbf{x}_{\parallel})$ describes (volume conserving) shear deformations, one would expect $\Omega(\hat{q})$ to vanish. Here, we show that this is indeed the case. For $\hat{q} = (\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}, \mathbf{0}, \dots, \mathbf{0})$, we obtain:

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$$\mathbf{q}_{\parallel} = \frac{1}{\sqrt{n+1}} \mathbf{q} \quad (\text{B.130a})$$

$$q_{\perp} = \hat{q} - \frac{1}{n+1} \begin{pmatrix} \mathbf{q} \\ \vdots \\ \mathbf{q} \end{pmatrix} \quad (\text{B.130b})$$

Hence, we can write Eq. (B.129b) in terms of \mathbf{q} :

$$\begin{aligned} & \int d\mathbf{x}_{\parallel} \exp(\mathbf{i}\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + \mathbf{i}q_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) \\ &= \int d\mathbf{x}_{\parallel} \exp\left(\frac{\mathbf{i}\mathbf{q}\mathbf{x}_{\parallel}}{\sqrt{n+1}} + \mathbf{i}\mathbf{q}\mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel}) - \frac{\mathbf{i}\mathbf{q}}{n+1} \sum_{\beta=0}^n \mathbf{u}_{\perp}^{(\beta)}(\mathbf{x}_{\parallel})\right) \end{aligned} \quad (\text{B.131})$$

The last term vanishes, because by definition, the sum over the components of a \perp -vector is zero, $\sum_{\beta=0}^n \mathbf{u}_{\perp}^{(\beta)}(\mathbf{x}_{\parallel}) = \mathbf{0}$. Now we can introduce the new integration variable

$$\mathbf{x}'_{\parallel} := \mathbf{x}_{\parallel} + \sqrt{n+1} \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel}) \quad (\text{B.132a})$$

$$\det\left(\frac{\partial \mathbf{x}'_{\parallel}}{\partial \mathbf{x}_{\parallel}}\right) = \det\left(\mathbb{1} + \sqrt{n+1} \frac{\partial \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel})}{\partial \mathbf{x}_{\parallel}}\right), \quad (\text{B.132b})$$

with which Eq. (B.131) becomes:

$$\begin{aligned} & \int d\mathbf{x}_{\parallel} \exp(\mathbf{i}\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + \mathbf{i}q_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) \\ &= \int d\mathbf{x}'_{\parallel} \exp\left(\frac{\mathbf{i}\mathbf{q}\mathbf{x}'_{\parallel}}{\sqrt{n+1}}\right) \left| \det\left(\mathbb{1} + \sqrt{n+1} \frac{\partial \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel})}{\partial \mathbf{x}_{\parallel}}\right) \right| \end{aligned} \quad (\text{B.133})$$

The prefactor $\sqrt{n+1}$ in the determinant is just coming from the difference between the usual vector \mathbf{x} and the parallel-vector $\mathbf{x}_{\parallel} = \sqrt{n+1} \mathbf{x}$. We only allow incompressible deformations (see Sec. 2.6.2), for which

$$\det\left(\mathbb{1} + \sqrt{n+1} \frac{\partial \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel})}{\partial \mathbf{x}_{\parallel}}\right) = \det\left(\mathbb{1} + \frac{\partial \mathbf{u}_{\perp}^{(\alpha)}(\mathbf{x}_{\parallel})}{\partial \mathbf{x}}\right) \equiv 1 \quad (\text{B.134})$$

and hence:

$$\begin{aligned} & \int d\mathbf{x}_{\parallel} \exp(\mathbf{i}\mathbf{q}_{\parallel} \mathbf{x}_{\parallel} + \mathbf{i}q_{\perp} u_{\perp}(\mathbf{x}_{\parallel})) \\ &= \int d\mathbf{x}'_{\parallel} \exp(\mathbf{i}\mathbf{q}_{\parallel} \mathbf{x}'_{\parallel}) = V_{\parallel} \delta_{\mathbf{q}_{\parallel}, \mathbf{0}} \end{aligned} \quad (\text{B.135})$$

Therefore, for incompressible deformations and \hat{q} -vectors in the IRS (which are non-zero), the order parameter becomes:

$$\bar{\Omega}_u(\hat{q}) = (1 - Q) \delta_{\hat{q}, \hat{0}} + Q \delta_{\mathbf{q}_{\parallel}, \mathbf{0}} = 0, \quad (\text{B.136})$$

as we already anticipated.

Calculations for the Directed Polymer Model

Contents

| | | |
|------------|---|------------|
| C.1 | Calculations for a Single Chain | 157 |
| C.1.1 | The Uncrosslinked Chain | 157 |
| C.1.2 | The Polymer Clamped in Space | 159 |
| C.2 | Examples of Single Chain Interactions | 160 |
| C.3 | Calculations for the Disorder-Averaged Free Energy | 162 |
| C.3.1 | Introduction of Replicas | 162 |
| C.3.2 | Introduction of the Replicated Density Field | 163 |
| C.3.3 | The Hubbard-Stratonovich Transformation | 164 |
| C.4 | The Saddle-Point Equation with Ansatz | 167 |
| C.4.1 | Expansion in Q to Infinite Order | 167 |
| C.4.2 | Expansion to Second Order in Q | 169 |
| C.5 | Obtaining the Equation for the Localization Length | 173 |
| C.5.1 | Normalization of Length Scales | 173 |
| C.5.2 | Laplace-Transformation of the Saddle Point Equation | 175 |
| C.6 | Calculation of a Correlator | 177 |
| C.7 | The Average Cross-Link Density | 180 |

C.1 Calculations for a Single Chain

C.1.1 The Uncrosslinked Chain

In Sec. 3.2.1 we defined the radius of gyration of an uncrosslinked polymer chain as

$$\begin{aligned}
 R_g^2 &= \left\langle \frac{1}{2L^2} \int dz_1 dz_2 (\mathbf{r}(z_1) - \mathbf{r}(z_2))^2 \right\rangle_{H_{\text{align},1}} \\
 &= \frac{1}{2L^2} \int \mathcal{D}\mathbf{r}(z) \exp\left(-\frac{\sigma}{2} \int dz \dot{\mathbf{r}}^2(z)\right) \int dz_1 dz_2 (\mathbf{r}(z_1) - \mathbf{r}(z_2))^2 \quad (\text{C.1})
 \end{aligned}$$

We discretize the functional integral into Z_L points, such that the function $\mathbf{r}(z)$ is represented by Z_L discrete vectors $\mathbf{r}_1, \dots, \mathbf{r}_{Z_L}$. With the normalization (3.4) of the functional integral we get:

$$R_g^2 = \frac{\frac{1}{2Z_L^2} \int d\mathbf{r}_1 \cdots d\mathbf{r}_{Z_L} \exp\left(-\frac{\sigma \cdot \Delta z}{2} \sum_{Z=1}^{Z_L-1} \frac{(\mathbf{r}_{Z+1} - \mathbf{r}_Z)^2}{\Delta z^2}\right) \sum_{Z_1, Z_2=1}^{Z_L} (\mathbf{r}_{Z_1} - \mathbf{r}_{Z_2})^2}{\int d\mathbf{r}_1 \cdots d\mathbf{r}_{Z_L} \exp\left(-\frac{\sigma \cdot \Delta z}{2} \sum_{Z=1}^{Z_L-1} \frac{(\mathbf{r}_{Z+1} - \mathbf{r}_Z)^2}{\Delta z^2}\right)} \quad (\text{C.2})$$

Now, instead of integrating over all *positions* of the chain segments \mathbf{r}_Z , we can integrate over the position of the bottom end of the chain \mathbf{r}_1 and over the position *differences* $\mathbf{t}_Z := \mathbf{r}_Z - \mathbf{r}_{Z-1}$ for $Z = 2, \dots, Z_L$:

$$\begin{aligned} R_g^2 &= \frac{\frac{1}{Z_L^2} \int d\mathbf{r}_1 d\mathbf{t}_2 \cdots d\mathbf{t}_{Z_L} \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=1}^{Z_L-1} \mathbf{t}_{Z+1}^2\right) \sum_{Z_1 < Z_2} \left(\sum_{Z=Z_1+1}^{Z_2} \mathbf{t}_Z\right)^2}{\int d\mathbf{r}_1 d\mathbf{t}_2 \cdots d\mathbf{t}_{Z_L} \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=1}^{Z_L-1} \mathbf{t}_{Z+1}^2\right)} \\ &= \frac{1}{Z_L^2} \sum_{Z_1 < Z_2} \frac{\int d\mathbf{t}_{Z_1+1} \cdots d\mathbf{t}_{Z_2} \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=Z_1+1}^{Z_2} \mathbf{t}_Z^2\right) \left(\sum_{Z=Z_1+1}^{Z_2} \mathbf{t}_Z^2 + \sum_{\substack{Z, Z' \neq Z_1+1 \\ Z \neq Z'}} \mathbf{t}_Z \mathbf{t}_{Z'}\right)}{\int d\mathbf{t}_{Z_1+1} \cdots d\mathbf{t}_{Z_2} \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=Z_1+1}^{Z_2} \mathbf{t}_Z^2\right)} \\ &= \frac{1}{Z_L^2} \sum_{Z_1 < Z_2} \frac{\left(\frac{2\pi \cdot \Delta z}{\sigma}\right)^{Z_2 - Z_1} \frac{\Delta z \cdot D(Z_2 - Z_1)}{2\sigma}}{\left(\frac{2\pi \cdot \Delta z}{\sigma}\right)^{Z_2 - Z_1}} \\ &= \frac{D\Delta z}{2Z_L^2\sigma} \sum_{Z_1 < Z_2} (Z_2 - Z_1) \\ &= \frac{D\Delta z}{2Z_L^2\sigma} \left(\frac{1}{6}(Z_L^2 - 1)Z_L\right) \stackrel{\text{large } Z_L}{=} \frac{D\Delta z}{2Z_L^2\sigma} \left(\frac{Z_L^3}{6}\right) = \frac{D}{2} \frac{Z_L \cdot \Delta z}{6\sigma}, \end{aligned} \quad (\text{C.3})$$

and since $\Delta z = L/Z_L$ we get:

$$\boxed{R_g^2 = \frac{D}{6} \frac{L}{2\sigma}} \quad (\text{C.4})$$

C.1.2 The Polymer Clamped in Space

Let us have a closer look at the partition function. As in the previous section 3.2.1, we discretize the functional integral into integrals over the chain segments:

$$\begin{aligned}
Z &= \int \mathcal{D}\mathbf{r}(z) \exp\left(-\frac{\sigma}{2} \int dz \dot{\mathbf{r}}^2(z) - \frac{1}{2a^2} \mathbf{r}^2(z_1) + \mathbf{b}\mathbf{r}(z_0)\right) \\
&= \frac{\int d\mathbf{r}_1 \cdots d\mathbf{r}_{Z_L} \exp\left(-\frac{\sigma \cdot \Delta z}{2} \sum_{Z=1}^{Z_L-1} \frac{(\mathbf{r}_{Z+1} - \mathbf{r}_Z)^2}{\Delta z^2} - \frac{\mathbf{r}_{Z_1}^2}{2a^2} + \mathbf{b}\mathbf{r}_{Z_0}\right)}{\int d\mathbf{r}_1 \cdots d\mathbf{r}_{Z_L} \exp\left(-\frac{\sigma \cdot \Delta z}{2} \sum_{Z=1}^{Z_L-1} \frac{(\mathbf{r}_{Z+1} - \mathbf{r}_Z)^2}{\Delta z^2}\right)} \quad (\text{C.5})
\end{aligned}$$

The denominator is easy to compute:

$$\int d\mathbf{r}_1 d\mathbf{t}_2 \cdots d\mathbf{t}_{Z_L} \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=1}^{Z_L-1} \mathbf{t}_{Z+1}^2\right) = A \left(\frac{2\pi \Delta z}{\sigma}\right)^{D(Z_L-1)/2}. \quad (\text{C.6})$$

For the numerator we keep the integration variable \mathbf{r}_{Z_1} and define $\mathbf{t}_Z = \mathbf{r}_{Z+1} - \mathbf{r}_Z$ for $1 \leq Z \leq Z_L - 1$. This change of variables yields the following substitution scheme for \mathbf{r}_Z :

$$\begin{aligned}
&\vdots \\
\mathbf{r}_{Z_1-2} &= \mathbf{r}_{Z_1} - \mathbf{t}_{Z_1-1} - \mathbf{t}_{Z_1-2} \\
\mathbf{r}_{Z_1-1} &= \mathbf{r}_{Z_1} - \mathbf{t}_{Z_1-1} \\
\mathbf{r}_{Z_1} &= \mathbf{r}_{Z_1} \\
\mathbf{r}_{Z_1+1} &= \mathbf{r}_{Z_1} + \mathbf{t}_{Z_1} \\
\mathbf{r}_{Z_1+2} &= \mathbf{r}_{Z_1} + \mathbf{t}_{Z_1} + \mathbf{t}_{Z_1+1} \\
&\vdots
\end{aligned}$$

Now we apply this scheme to the numerator of Eq. (C.5) and first consider the case $Z_0 > Z_1$:

$$\begin{aligned}
Z &= \frac{1}{A} \left(\frac{2\pi \Delta z}{\sigma}\right)^{-D(Z_L-1)/2} \int d\mathbf{t}_1 \cdots d\mathbf{t}_{Z_1-1} d\mathbf{r}_{Z_1} d\mathbf{t}_{Z_1} \cdots d\mathbf{t}_{Z_L-1} \\
&\quad \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=1}^{Z_L-1} \mathbf{t}_Z^2 - \frac{\mathbf{r}_{Z_1}^2}{2a^2} + \mathbf{b}(\mathbf{r}_{Z_1} + \mathbf{t}_{Z_1} + \dots + \mathbf{t}_{Z_0-1})\right) \\
&= \frac{1}{A} \left(\frac{2\pi \Delta z}{\sigma}\right)^{-D(Z_L-1)/2} \\
&\quad \times \left(\frac{2\pi \Delta z}{\sigma}\right)^{D(Z_L-1)/2} (2\pi a^2)^{D/2} \exp\left(\frac{\Delta z}{2\sigma} \mathbf{b}^2 \cdot (Z_0 - Z_1) + \frac{\mathbf{b}^2 a^2}{2}\right) \\
&= \frac{(2\pi a^2)^{D/2}}{A} \exp\left(\frac{\mathbf{b}^2}{2\sigma} \Delta z (Z_0 - Z_1) + \frac{\mathbf{b}^2 a^2}{2}\right) \quad (\text{C.7})
\end{aligned}$$

Analogously we get for $Z_0 < Z_1$:

$$Z = \frac{(2\pi a^2)^{D/2}}{A} \exp\left(\frac{\mathbf{b}^2}{2\sigma} \Delta z (Z_1 - Z_0) + \frac{\mathbf{b}^2 a^2}{2}\right), \quad (\text{C.8})$$

and thus we get in general:

$$Z = \frac{(2\pi a^2)^{D/2}}{A} \exp\left(\frac{\mathbf{b}^2}{2} \left(\frac{|z_1 - z_0|}{\sigma} + a^2\right)\right), \quad (\text{C.9})$$

where we re-substituted $z_i = Z_i \Delta z$ for $i = 1, 2$. For the logarithm of Z we get:

$$\ln Z = \frac{\mathbf{b}^2}{2} \left(\frac{|z_1 - z_0|}{\sigma} + a^2\right) + \ln\left(\frac{(2\pi a^2)^{D/2}}{A}\right). \quad (\text{C.10})$$

C.2 Examples of Single Chain Interactions

As mentioned in Sec. 3.3.1 on page 39, it is possible to replace the alignment interaction H_{align} by a more general interaction which also can be written as a sum over individual chains:

$$H_{\Sigma} = \sum_{j=1}^N H_{\Sigma,1}\{\mathbf{r}_j\} = \sum_{j=1}^N (H_{\text{align},1}\{\mathbf{r}_j\} + H_{X,1}\{\mathbf{r}_j\} + \dots). \quad (\text{C.11})$$

The only requirement is that $H_{\Sigma,1}\{\mathbf{r}_j\}$ only depends on the conformation $\mathbf{r}_j(z)$ of chain j . With the total Hamiltonian $H_{\mathcal{C}} = H_{\text{Xlink}} + H_{\text{ev}} + H_{\Sigma}$, the effective one-particle partition function \mathfrak{z} , Eq. (3.32c), is simply replaced by:

$$\mathfrak{z} = \int \mathcal{D}\hat{r}(z) \exp\left(\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} + i \int_0^L dz \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} - H_{\Sigma,1}^{(n+1)}\{\hat{r}(z)\}\right), \quad (\text{C.12})$$

with $H_{\Sigma,1}^{(n+1)}\{\hat{r}(z)\} = \sum_{\alpha=0}^n H_{\Sigma,1}\{\mathbf{r}_j^{(\alpha)}\}$. The other parts of Eq. (3.32) remain as they are. Beyond the alignment interaction, these interactions $\sum_j H_{X,1}\{\mathbf{r}_j\}$ may include:

- A tilt field $\mathbf{h}(z)$, creating a force on the chains:

$$H_{\text{tilt}} = \sum_{j=1}^N \int_0^L dz \mathbf{h}(z) \cdot \dot{\mathbf{r}}_j(z) \quad (\text{C.13})$$

For this Hamiltonian, a constant $\mathbf{h}(z) = \mathbf{h}$ corresponds to a linear shear force, since $\mathbf{h}(z)$ couples to the tilt $\dot{\mathbf{r}}_j(z)$ of the chains.

- A pressure field $\Pi(\mathbf{q}, z)$ that couples to the Fourier density:

$$H_{\Pi} = \sum_{\mathbf{q}} \int_0^L dz \Pi(\mathbf{q}, z) \sum_{j=1}^N e^{-i\mathbf{q}\mathbf{r}_j(z)}, \quad (\text{C.14})$$

$$\text{such that} \quad \frac{\delta H_{\Pi}}{\delta \Pi(\mathbf{q}, z)} = \sum_{j=1}^N e^{-i\mathbf{q}\mathbf{r}_j(z)} \quad (\text{C.15a})$$

$$\text{and} \quad \frac{\delta^2 H_{\Pi}}{\delta \Pi(\mathbf{q}, z_1) \delta \Pi(-\mathbf{q}, z_2)} = \sum_{i,j=1}^N e^{-i\mathbf{q}(\mathbf{r}_i(z_1) - \mathbf{r}_j(z_2))}. \quad (\text{C.15b})$$

With this field, we would be able to calculate density-density correlations.

- A “clamping interaction”, that clamps the ends of the chains, $\mathbf{r}_j(0)$ and $\mathbf{r}_j(L)$, to certain points at the bottom and top (the boundaries in z -direction) of the system:

$$H_{\text{clamp}} = \frac{1}{2b_0^2} \sum_{j=1}^N (\mathbf{r}_j(0) - \mathbf{r}_{0,j})^2 + \frac{1}{2b_L^2} \sum_{j=1}^N (\mathbf{r}_j(L) - \mathbf{r}_{L,j})^2 \quad (\text{C.16})$$

Here b_0 and b_L are the clamping strengths at the bottom ($z = 0$) and the top ($z = L$) of the system. The clamping points $\mathbf{r}_{0,j}$ and $\mathbf{r}_{L,j}$ could be included in the disorder, $\mathcal{C} = (\{(i_e, j_e, z_e)\}_{e=1}^M, \{(\mathbf{r}_{0,j}, \mathbf{r}_{L,j})\}_{j=1}^N)$, such that their distribution is chosen by the Deam-Edwards distribution. For polymer brushes, which are typically mounted with one end onto a plate, this additional interaction with only one clamping term (*i.e.* b_0 finite and $b_L \rightarrow \infty$) would be an appropriate choice.

- A bending stiffness which penalizes bending, *i.e.* high values of the second derivative of the chains:

$$H_{\text{bend}} = \frac{\kappa}{2} \sum_{j=1}^N \int_0^L dz \ddot{\mathbf{r}}_j^2(z) \quad (\text{C.17})$$

It is notable that, only this straightening interaction can produce a non-zero persistence length. Without this interaction, the slopes of a given chain at two different points z_1 and z_2 are fully uncorrelated, no matter how close those points are.

C.3 Calculations for the Disorder-Averaged Free Energy

C.3.1 Introduction of Replicas

In Eq. 3.24, we calculated the important quantity:

$$[Z_C^n] = \frac{1}{Z_1} \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2/L)^M}{M!} \sum_{\substack{i_1, \dots, i_M, \\ j_1, \dots, j_M=1}}^N \int_0^L dz_1 \cdots dz_M \\ \int \mathcal{D}\hat{r}_1(z) \cdots \mathcal{D}\hat{r}_N(z) e^{-H_{\text{ev}}^{(n+1)}\{\hat{r}_j\} - H_{\text{align}}^{(n+1)}\{\hat{r}_j\}} \prod_{e=1}^M \exp\left(-\frac{(\hat{r}_{i_e}(z_e) - \hat{r}_{j_e}(z_e))^2}{2a^2}\right).$$

Here, we perform the sum over M . For simplification, we define:

$$\mathcal{D}\Gamma := \mathcal{D}\hat{r}_1(z) \cdots \mathcal{D}\hat{r}_N(z) \quad (\text{C.18})$$

and write the product over e as a power:

$$[Z_C^n] = \frac{1}{Z_1} \sum_{M=0}^{\infty} \frac{(\tilde{\mu}^2/L)^M}{M!} \int \mathcal{D}\Gamma e^{-H_{\text{ev}}^{(n+1)}\{\hat{r}_j\} - H_{\text{align}}^{(n+1)}\{\hat{r}_j\}} \\ \left(\sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{(\hat{r}_i(z) - \hat{r}_j(z))^2}{2a^2}\right) \right)^M \\ = \frac{1}{Z_1} \int \mathcal{D}\Gamma e^{-H_{\text{ev}}^{(n+1)}\{\hat{r}_j\} - H_{\text{align}}^{(n+1)}\{\hat{r}_j\}} \\ \exp\left(\frac{\tilde{\mu}^2}{L} \sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{(\hat{r}_i(z) - \hat{r}_j(z))^2}{2a^2}\right)\right) \\ =: \frac{Z_{n+1}}{Z_1}. \quad (\text{C.19})$$

Here, the **replica partition function** Z_{n+1} has been defined:

$$Z_{n+1} = \int \mathcal{D}\Gamma \exp(-N\tilde{f}_{n+1}\{\hat{r}_j\}) \quad \text{with} \quad (\text{C.20a})$$

with the **replica free energy** $\tilde{f}_{n+1}\{\hat{r}_j\}$:

$$\tilde{f}_{n+1}\{\hat{r}_j\} = \frac{H_{\text{ev}}^{(n+1)}\{\hat{r}_j\}}{N} + \frac{H_{\text{align}}^{(n+1)}\{\hat{r}_j\}}{N} - \frac{\tilde{\mu}^2}{LN} \sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{(\hat{r}_i(z) - \hat{r}_j(z))^2}{2a^2}\right) \quad (\text{C.20b})$$

C.3.2 Introduction of the Replicated Density Field

In Sec. 3.4.2, we introduced the replicated density field, Eqs. (3.29). Here we rewrite the replica free energy (3.27c),

$$\begin{aligned} \tilde{f}_{n+1}\{\hat{r}_j\} = & -\frac{\tilde{\mu}^2}{LN} \sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{(\hat{r}_i(z) - \hat{r}_j(z))^2}{2a^2}\right) \\ & + \frac{\lambda}{2N} \sum_{\alpha=0}^n \sum_{i,j=1}^N \int_0^L dz U(\mathbf{r}_i^{(\alpha)}(z) - \mathbf{r}_j^{(\alpha)}(z)) + \frac{H_{\text{align}}^{(n+1)}}{N}, \end{aligned} \quad (\text{C.21})$$

in terms of that field. In exactly the same manner as in Eqs. (B.5-B.7), we can rewrite the cross-linking term as:

$$\boxed{\sum_{i,j=1}^N \exp\left(-\frac{(\hat{r}_i(z) - \hat{r}_j(z))^2}{2a^2}\right) = \phi^{n+1} N^2 \sum_{\hat{q}} |\text{O}(\hat{q}, z)|^2 \Delta(\hat{q})}, \quad (\text{C.22})$$

with

$$\Delta(\hat{q}) = \exp\left(-\frac{\hat{q}^2 a^2}{2}\right). \quad (\text{C.23})$$

For the second (excluded volume) term in Eq. (C.21), we first define the density field for each replica separately:

$$\text{O}^{(\alpha)}(\mathbf{x}, z) = \frac{1}{N} \sum_{j=1}^N \delta(\mathbf{x} - \mathbf{r}_j^{(\alpha)}(z)), \quad (\text{C.24a})$$

$$\text{O}^{(\alpha)}(\mathbf{q}, z) = \frac{1}{N} \sum_{j=1}^N \exp(i\mathbf{q}\mathbf{r}_j^{(\alpha)}(z)). \quad (\text{C.24b})$$

With that, also the excluded volume Hamiltonian can be rewritten in analogy to the RLP model, see Eq. (B.9), we just have to add the integral over z :

$$\begin{aligned} & \sum_{\alpha=0}^n \sum_{i,j=1}^N \int_0^L dz U(\mathbf{r}_i^{(\alpha)}(z) - \mathbf{r}_j^{(\alpha)}(z)) \\ & = \frac{N^2}{A} \sum_{\alpha=0}^n \sum_{\mathbf{q}^{(\alpha)}} \int_0^L dz |\text{O}^{(\alpha)}(\mathbf{q}^{(\alpha)}, z)|^2 U(\mathbf{q}^{(\alpha)}). \end{aligned} \quad (\text{C.25})$$

With that and by plugging in $\frac{\tilde{\mu}^2}{LN} = \frac{\mu^2}{2LN^2\phi}$ (see Eq. (3.22c)), we can write the replica free energy, Eq. (C.21), as:

$$\begin{aligned} \tilde{f}_{n+1}\{\hat{r}_j\} = & -\phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q}} |\text{O}(\hat{q}, z)|^2 \Delta(\hat{q}) \\ & + \frac{\lambda n_0}{2} \int_0^L dz \sum_{\alpha=0}^n \sum_{\mathbf{q}^{(\alpha)}} |\text{O}^{(\alpha)}(\mathbf{q}^{(\alpha)}, z)|^2 U(\mathbf{q}^{(\alpha)}) + \frac{H_{\text{align}}^{(n+1)}\{\hat{r}_j\}}{N}, \end{aligned} \quad (\text{C.26})$$

where we defined the mean chain density

$$n_0 = \frac{N}{A} \quad (\text{C.27})$$

as number of chains per (hyper-)area.

We now separate the two terms involving fields in Eq. (C.26) into one replica sector (1RS), the set of replicated \hat{q} -vectors with exactly one of the $n+1$ components non-zero, and higher replica sector (HRS), the set of replicated \hat{q} -vectors with at least two non-zero components, as introduced in Sec. 2.4.3. This will allow us to perform the Hubbard-Stratonovich transformation later.

$$\begin{aligned} \tilde{f}_{n+1}\{\hat{r}_j\} &= \overbrace{-\phi^n \frac{\mu^2}{2} + (n+1) \frac{\lambda n_0 L}{2}}{=:f_0} - \phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |\text{O}(\hat{q}, z)|^2 \Delta(\hat{q}) \\ &+ \frac{1}{2} \int_0^L dz \sum_{\hat{q} \in \text{1RS}} \underbrace{\left(\lambda n_0 U(\hat{q}) - \phi^n \frac{\mu^2}{L} \Delta(\hat{q}) \right)}{=: \tilde{\lambda}(\hat{q})} |\text{O}(\hat{q}, z)|^2 + \frac{H_{\text{align}}^{(n+1)}\{\hat{r}_j\}}{N} \end{aligned} \quad (\text{C.28})$$

For the terms involving $\hat{q} = \hat{0}$, which are neither part of the 1RS, nor of the HRS, we used $\text{O}(\hat{q} = \hat{0}) = 1$, $\Delta(\hat{q} = \hat{0}) = 1$ and $U(\mathbf{q} = \mathbf{0}) = 1$.

C.3.3 The Hubbard-Stratonovich Transformation

Here we want to apply the Hubbard-Stratonovich transformation to the replica free energy Eq. (3.31a) to linearize it with respect to the field $\text{O}(\hat{q}, z)$. The appropriate variable for the transformation is $\text{O}(\hat{q}, z)$, similar to the RLP model. A transformation with respect to $\int dz \text{O}(\hat{q}, z)$, similar to what has been done in [Goldbart *et al.*, 1996, Chapter V], would require a different structure, a form like $\int dz_1 dz_2 \text{O}(\hat{q}, z_1) \text{O}(-\hat{q}, z_2)$ in the replica free energy. Hence, we apply Eqs. (B.12) successively for all combinations of \hat{q} and z .

Before performing the transformation, it is important to note that the replica density obeys the relation $\text{O}(-\hat{q}, z) = \text{O}^*(\hat{q}, z)$ for any height z , see definition (3.29b), in particular $|\text{O}(\hat{q}, z)|^2 = |\text{O}(-\hat{q}, z)|^2$; hence, for any \hat{q} , we automatically transform the $|\text{O}(\hat{q}, z)|^2$ -term along the $|\text{O}(-\hat{q}, z)|^2$ -term. In order to carry out the Hubbard-Stratonovich transformation, we restrict the sums over \hat{q} to a half-space with $\hat{q} \cdot \hat{e} > 0$ with $\hat{e} = (\mathbf{e}, \dots, \mathbf{e})$ for arbitrary $\mathbf{e} \neq \mathbf{0}$ (as it was done in Sec. B.2.1 for the RLP model). The terms for \hat{q} and $-\hat{q}$ yield the same contribution, since $\Delta(\hat{q}) = \Delta(-\hat{q})$ and $\tilde{\lambda}(\hat{q}) = \tilde{\lambda}(-\hat{q})$.

With that we get for replica partition function \mathcal{Z}_{n+1} , Eq. (3.27b):

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} &= \int \left(\prod_{j=1}^N \mathcal{D}\hat{r}_j(z) \right) \\ &\exp \left(N\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} |\text{O}(\hat{q}, z)|^2 \Delta(\hat{q}) - N \int_0^L dz \sum_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} |\text{O}(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) - H_{\text{align}}^{(n+1)} \right) \end{aligned} \quad (\text{C.29})$$

Now we apply the Hubbard-Stratonovich transformation with $w = \text{O}(\hat{q}, z)$: For any $\hat{q} \in \text{1RS}$, $\hat{q} \cdot \hat{e} > 0$ and any $z \in [0, L]$ we use Eq. (B.12b) with $c = N\tilde{\lambda}(\hat{q})$:

$$\begin{aligned} &\exp \left(-N |\text{O}(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) \right) \quad (\text{C.30}) \\ &= \int \frac{N\tilde{\lambda}(\hat{q}) d\Omega(\hat{q}, z)}{\pi} \exp \left(-N |\Omega(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) + 2iN\tilde{\lambda}(\hat{q}) \text{Re}(\Omega(\hat{q}, z)\text{O}(-\hat{q}, z)) \right) \end{aligned}$$

and for $\hat{q} \in \text{HRS}$, $\hat{q} \cdot \hat{e} > 0$ and $z \in [0, L]$ we use Eq. (B.12a) with $c = N\phi^n \frac{\mu^2}{L} \Delta(\hat{q})$:

$$\begin{aligned} \exp \left(N\phi^n \frac{\mu^2}{L} |\text{O}(\hat{q}, z)|^2 \Delta(\hat{q}) \right) &= \int \frac{N\phi^n \mu^2 \Delta(\hat{q}) d\Omega(\hat{q}, z)}{\pi L} \quad (\text{C.31}) \\ &\exp \left(-N\phi^n \frac{\mu^2}{L} \Delta(\hat{q}) |\Omega(\hat{q}, z)|^2 + 2N\phi^n \frac{\mu^2}{L} \Delta(\hat{q}) \text{Re}(\Omega(\hat{q}, z)\text{O}(-\hat{q}, z)) \right) \end{aligned}$$

With that Eq. (C.29) becomes:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} &= \\ &\int \mathcal{D}\Omega \exp \left(-N\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} |\Omega(\hat{q}, z)|^2 \Delta(\hat{q}) - N \int_0^L dz \sum_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} |\Omega(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) \right) \\ &\times \int \left(\prod_{j=1}^N \mathcal{D}\hat{r}_j(z) \right) \exp \left(2N\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} \Delta(\hat{q}) \text{Re}(\Omega(\hat{q}, z)\text{O}(-\hat{q}, z)) \right. \\ &\quad \left. + 2iN \int_0^L dz \sum_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} \tilde{\lambda}(\hat{q}) \text{Re}(\Omega(\hat{q}, z)\text{O}(-\hat{q}, z)) - H_{\text{align}}^{(n+1)}(\{\hat{r}_j\}) \right), \end{aligned} \quad (\text{C.32})$$

where we introduced the measure

$$\mathcal{D}\Omega = \prod_z \left(\prod_{\substack{\hat{q} \in \text{1RS} \\ \hat{q} \cdot \hat{e} > 0}} \frac{N\tilde{\lambda}(\hat{q}) d\Omega(\hat{q}, z)}{\pi} \cdot \prod_{\substack{\hat{q} \in \text{HRS} \\ \hat{q} \cdot \hat{e} > 0}} \frac{N\phi^n \mu^2 \Delta(\hat{q}) d\Omega(\hat{q}, z)}{\pi L} \right). \quad (\text{C.33})$$

Here, the product over z has to be carried out in the spirit of the discretization of the z -direction into Z_L segments, as seen in Sec. 3.3 and Fig. 3.5. It implies that we have to integrate over $\Omega(\hat{q}, z)$ for all \hat{q} and at every height z .

Now we abolish the constraint $\hat{q} \cdot \hat{e} > 0$ in the sums. Therefore define the field $\Omega(\hat{q}, z)$ also for $\hat{q} \cdot \hat{e} < 0$ by

$$\Omega(-\hat{q}, z) = \Omega^*(\hat{q}, z). \quad (\text{C.34})$$

and plug in the definition of $O(\hat{q}, z)$, Eq. (3.29b). Furthermore we recall that $H_{\text{align}}^{(n+1)}$ acts on all chains in the same way (see Eqs. (3.17), (3.19)):

$$H_{\text{align}}^{(n+1)} = \sum_{j=1}^N H_{\text{align},1}^{(n+1)}\{\hat{r}_j\} \quad (\text{C.35})$$

Hence the partition function (C.32) becomes:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} = & \int \mathcal{D}\Omega \exp\left(-N\phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |\Omega(\hat{q}, z)|^2 \Delta(\hat{q}) - \frac{N}{2} \int_0^L dz \sum_{\hat{q} \in \text{1RS}} |\Omega(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q})\right) \\ & \times \int \left(\prod_{j=1}^N \mathcal{D}\hat{r}_j(z) \right) \exp\left(\phi^n \frac{\mu^2}{L} \sum_{j=1}^N \int_0^L dz \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}_j(z)} \right. \\ & \left. + i \sum_{j=1}^N \int_0^L dz \sum_{\hat{q} \in \text{1RS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}_j(z)} - \sum_{j=1}^N H_{\text{align},1}^{(n+1)}\{\hat{r}_j(z)\} \right) \end{aligned} \quad (\text{C.36})$$

As in the RLP model before (see Sec. 2.4.4 and B.2.1), the expression is now linear in the density and therefore only generates “linear” sums \sum_j over all particles. We also see, why the term $H_{\text{align}}^{(n+1)}$ was so easy to take along and did not require a Hubbard-Stratonovich transformation: it is already linear in the sum \sum_j . Now, the sums in the exponential and the difficult integral over all chains can be written as a power:

$$\begin{aligned} \frac{\mathcal{Z}_{n+1}}{\exp(-Nf_0)} = & \int \mathcal{D}\Omega \exp\left(-N\phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |\Omega(\hat{q}, z)|^2 \Delta(\hat{q}) - \frac{N}{2} \int_0^L dz \sum_{\hat{q} \in \text{1RS}} |\Omega(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q})\right) \\ & \times \left\{ \int \mathcal{D}\hat{r}(z) \exp\left(\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} \right. \right. \\ & \left. \left. + i \int_0^L dz \sum_{\hat{q} \in \text{1RS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} - H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\} \right) \right\}^N \end{aligned} \quad (\text{C.37})$$

And that form can be written in a simplified way:

$$\mathcal{Z}_{n+1} = \exp(-Nf_0) \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\}) \quad (\text{C.38a})$$

with the replica free energy

$$f_{n+1}\{\Omega\} = \phi^n \frac{\mu^2}{2L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} |\Omega(\hat{q}, z)|^2 \Delta(\hat{q}) + \frac{1}{2} \int_0^L dz \sum_{\hat{q} \in \text{IRS}} |\Omega(\hat{q}, z)|^2 \tilde{\lambda}(\hat{q}) - \ln \mathfrak{z}, \quad (\text{C.38b})$$

and an effective one-particle partition function

$$\mathfrak{z} = \int \mathcal{D}\hat{r}(z) \exp \left(\phi^n \frac{\mu^2}{L} \int_0^L dz \sum_{\hat{q} \in \text{HRS}} \Delta(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} + i \int_0^L dz \sum_{\hat{q} \in \text{IRS}} \tilde{\lambda}(\hat{q}) \Omega(\hat{q}, z) e^{-i\hat{q}\hat{r}(z)} - H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\} \right) \quad (\text{C.38c})$$

C.4 The Saddle-Point Equation with Ansatz

C.4.1 Expansion in Q to Infinite Order

Here we calculate the term $I(\hat{q}_0, z_0)$, Eq. (3.50). With the single chain alignment interaction:

$$H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\} \equiv \frac{\sigma}{2} \int_0^L dz \dot{\hat{r}}^2(z), \quad (\text{C.39})$$

we get:

$$I(\hat{q}_0, z_0) = \int \mathcal{D}\hat{r}(z) e^{-i\hat{q}_0\hat{r}(z_0)} \times \exp \left(\phi^n \frac{\mu^2 Q}{L} \int_0^L dz \sum_{\hat{q}} e^{-i\hat{q}\hat{r}(z)} \delta_{\mathbf{q}\parallel, \mathbf{0}} \int_{\xi^2, z} \exp \left(-\frac{\hat{q}^2 \tilde{\xi}^2}{2} \right) - \frac{\sigma}{2} \int_0^L dz \dot{\hat{r}}^2(z) \right). \quad (\text{C.40})$$

To proceed in the calculation, we would like to perform the functional integral over all chain positions $\hat{r}(z)$. Therefore we expand the integrand of $I(\hat{q}_0, z_0)$, Eq. (C.40), in powers of Q :

$$\begin{aligned} I(\hat{q}_0, z_0) &= \int \mathcal{D}\hat{r}(z) e^{-i\hat{q}_0\hat{r}(z_0)} \exp \left(-\frac{\sigma}{2} \int_0^L dz \dot{\hat{r}}^2(z) \right) \\ &\times \left(1 + \sum_{r=1}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} \left\{ \frac{1}{L} \int_0^L dz \sum_{\hat{q}} e^{-i\hat{q}\hat{r}(z)} \delta_{\mathbf{q}\parallel, \mathbf{0}} \int_{\xi^2, z} \exp \left(-\frac{\hat{q}^2 \tilde{\xi}^2}{2} \right) \right\}^r \right) \\ &= \int \mathcal{D}\hat{r}(z) e^{-i\hat{q}_0\hat{r}(z_0)} \exp \left(-\frac{\sigma}{2} \int_0^L dz \dot{\hat{r}}^2(z) \right) \\ &\times \left(1 + \sum_{r=1}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r! L^r} \int_0^L dz_1 \cdots dz_r \sum_{\hat{q}_1, \dots, \hat{q}_r} \delta_{\mathbf{q}_{1\parallel}, \mathbf{0}} \cdots \delta_{\mathbf{q}_{r\parallel}, \mathbf{0}} e^{-i(\hat{q}_1 \hat{r}(z_1) + \cdots + \hat{q}_r \hat{r}(z_r))} \right. \\ &\quad \left. \times \int_{\xi_1^2, z_1} \cdots \int_{\xi_r^2, z_r} \exp \left(-\frac{\hat{q}_1^2 \tilde{\xi}_1^2 + \cdots + \hat{q}_r^2 \tilde{\xi}_r^2}{2} \right) \right) \quad (\text{C.41}) \end{aligned}$$

All $\hat{r}(z)$ -dependent terms, which are needed to perform the functional integral, are marked in blue. This integral is calculated in Appendix C.6 and the result is given by Eq. (C.88):

$$\begin{aligned} & \int \mathcal{D}\hat{r}(z) e^{-i(\hat{q}_0\hat{r}(z_0)+\hat{q}_1\hat{r}(z_1)+\dots+\hat{q}_r\hat{r}(z_r))} \exp\left(-\frac{\sigma}{2} \int_0^L dz \hat{r}^2(z)\right) \\ &= \delta_{\hat{q}_0+\hat{q}_1+\dots+\hat{q}_r, \hat{0}} \exp\left(\frac{1}{2\sigma} \sum_{0 \leq \gamma < \delta \leq r} \hat{q}_\gamma \hat{q}_\delta |z_\gamma - z_\delta|\right), \end{aligned} \quad (\text{C.42})$$

and therefore:

$$\begin{aligned} I(\hat{q}_0, z_0) &= \delta_{\hat{q}_0, \hat{0}} + \sum_{r=1}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r! L^r} \int_0^L dz_1 \cdots dz_r \sum_{\hat{q}_1, \dots, \hat{q}_r} \delta_{\mathbf{q}_{1\parallel}, \mathbf{0}} \cdots \delta_{\mathbf{q}_{r\parallel}, \mathbf{0}} \delta_{\hat{q}_0+\hat{q}_1+\dots+\hat{q}_r, \hat{0}} \\ &\times \exp\left(\frac{1}{2\sigma} \sum_{0 \leq \gamma < \delta \leq r} \hat{q}_\gamma \hat{q}_\delta |z_\gamma - z_\delta|\right) \int_{\xi_1^2, z_1} \cdots \int_{\xi_r^2, z_r} \exp\left(-\frac{\hat{q}_1^2 \tilde{\xi}_1^2 + \dots + \hat{q}_r^2 \tilde{\xi}_r^2}{2}\right) \end{aligned} \quad (\text{C.43})$$

For this expression it is important to keep in mind that the sum $\sum_{0 \leq \gamma < \delta \leq r}$ in the first exponential also involves the *external* variables \hat{q}_0 and z_0 . We also see at this point that it is useful to normalize the height z by L to make it a dimensionless variable

$$s_\gamma := \frac{z_\gamma}{L} \in [0, 1] \quad \text{and hence} \quad ds_\gamma = \frac{dz_\gamma}{L} \quad (\text{C.44})$$

for $\gamma = 0, \dots, r$ in all orders. As already mentioned in Sec. 3.2.1 it is useful to define

$$\ell^2 := \frac{L}{2\sigma}, \quad (\text{C.45})$$

which essentially is the radius of gyration of the chain, projected along the z -axis. With that we get:

$$\frac{1}{2\sigma} \hat{q}_\gamma \hat{q}_\delta |z_\gamma - z_\delta| = \frac{L}{2\sigma} \hat{q}_\gamma \hat{q}_\delta |s_\gamma - s_\delta| = \ell^2 \hat{q}_\gamma \hat{q}_\delta |s_\gamma - s_\delta| \quad (\text{C.46})$$

Therefore:

$$\begin{aligned} I(\hat{q}_0, s_0) &= \\ &= \delta_{\hat{q}_0, \hat{0}} + \sum_{r=1}^{\infty} \frac{(\phi^n \mu^2 Q)^r}{r!} \int_0^1 ds_1 \cdots ds_r \sum_{\hat{q}_1, \dots, \hat{q}_r} \delta_{\mathbf{q}_{1\parallel}, \mathbf{0}} \cdots \delta_{\mathbf{q}_{r\parallel}, \mathbf{0}} \delta_{\hat{q}_0+\hat{q}_1+\dots+\hat{q}_r, \hat{0}} \\ &\times \exp\left(\ell^2 \sum_{0 \leq \gamma < \delta \leq r} \hat{q}_\gamma \hat{q}_\delta |s_\gamma - s_\delta|\right) \int_{\xi_1^2, s_1} \cdots \int_{\xi_r^2, s_r} \exp\left(-\frac{1}{2} \sum_{\gamma=1}^r \hat{q}_\gamma^2 \tilde{\xi}_\gamma^2\right). \end{aligned} \quad (\text{C.47})$$

Here we needed to define the integral

$$\int_{\xi^2, s} := \int_0^\infty d\xi^2 \mathcal{P}(\xi^2, s), \quad (\text{C.48})$$

where $\mathcal{P}(\xi^2, s)$ depends on the dimensionless height s , but apart from that it is the same as $\mathcal{P}(\xi^2, z)$. Furthermore, as we can see in Eq. (C.47), the only L -dependence remains in the prefactor ℓ^2 in the first exponential.

C.4.2 Expansion to Second Order in Q

Up to this point it was easy to keep $I(\hat{q}_0, s_0)$ up to infinite order in Q . In order to proceed in the calculation, we keep the expansion up to second order (thereby restricting the system to being close to the sol-gel transition) and separate the different orders:

$$\begin{aligned} I(\hat{q}_0, s_0) &= \delta_{\hat{q}_0, \hat{0}} + \\ &+ \phi^n \mu^2 Q \int_0^1 ds_1 \sum_{\hat{q}_1} \delta_{\mathbf{a}_{1\parallel}, \mathbf{0}} \delta_{\hat{q}_0 + \hat{q}_1, \hat{0}} \exp\left(\ell^2 \hat{q}_0 \hat{q}_1 |s_0 - s_1|\right) \int_{\xi_1^2, s_1} \exp\left(-\frac{\hat{q}_1^2 \tilde{\xi}_1^2}{2}\right) \\ &+ \frac{(\phi^n \mu^2 Q)^2}{2} \int_0^1 ds_1 ds_2 \sum_{\hat{q}_1, \hat{q}_2} \delta_{\mathbf{a}_{1\parallel}, \mathbf{0}} \delta_{\mathbf{a}_{2\parallel}, \mathbf{0}} \delta_{\hat{q}_0 + \hat{q}_1 + \hat{q}_2, \hat{0}} \\ &\quad \times \exp\left(\ell^2 \left(\hat{q}_0 \hat{q}_1 |s_0 - s_1| + \hat{q}_0 \hat{q}_2 |s_0 - s_2| + \hat{q}_1 \hat{q}_2 |s_1 - s_2|\right)\right) \\ &\quad \times \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} \exp\left(-\frac{\hat{q}_1^2 \tilde{\xi}_1^2 + \hat{q}_2^2 \tilde{\xi}_2^2}{2}\right) \\ &+ \mathcal{O}(Q^3). \end{aligned} \quad (\text{C.49})$$

In the first order term, the sum over \hat{q}_1 is easy to perform thanks to the $\delta_{\hat{q}_0 + \hat{q}_1, \hat{0}}$ -constraint:

$$\begin{aligned} I_1 &:= \sum_{\hat{q}_1} \delta_{\mathbf{a}_{1\parallel}, \mathbf{0}} \delta_{\hat{q}_0 + \hat{q}_1, \hat{0}} \exp\left(\ell^2 \hat{q}_0 \hat{q}_1 |s_0 - s_1|\right) \exp\left(-\frac{\hat{q}_1^2 \tilde{\xi}_1^2}{2}\right) \\ &= \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \exp\left(-\ell^2 \hat{q}_0^2 |s_0 - s_1| - \frac{\hat{q}_0^2 \tilde{\xi}_1^2}{2}\right) = \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \exp\left(-\frac{q_{0\perp}^2}{2} \left(2\ell^2 |s_0 - s_1| + \tilde{\xi}_1^2\right)\right) \end{aligned} \quad (\text{C.50})$$

In the last line we used the fact that $\hat{q}_0 = q_{0\perp}$, since $\mathbf{a}_{0\parallel} = \mathbf{0}$.

In the second order term, the situation is a little bit more difficult. First get rid of \hat{q}_2 in the same manner:

$$\begin{aligned} I_2 &:= \sum_{\hat{q}_1, \hat{q}_2} \delta_{\mathbf{a}_{1\parallel}, \mathbf{0}} \delta_{\mathbf{a}_{2\parallel}, \mathbf{0}} \delta_{\hat{q}_0 + \hat{q}_1 + \hat{q}_2, \hat{0}} \\ &\quad \times \exp\left(\ell^2 \left(\hat{q}_0 \hat{q}_1 |s_0 - s_1| + \hat{q}_0 \hat{q}_2 |s_0 - s_2| + \hat{q}_1 \hat{q}_2 |s_1 - s_2|\right)\right) \exp\left(-\frac{\hat{q}_1^2 \tilde{\xi}_1^2 + \hat{q}_2^2 \tilde{\xi}_2^2}{2}\right) \\ &= \sum_{\hat{q}_1} \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \delta_{\mathbf{a}_{1\parallel}, \mathbf{0}} \end{aligned}$$

$$\begin{aligned}
& \times \exp\left(\ell^2\left(\hat{q}_0\hat{q}_1|s_0 - s_1| - \hat{q}_0(\hat{q}_0 + \hat{q}_1)|s_0 - s_2| - \hat{q}_1(\hat{q}_0 + \hat{q}_1)|s_1 - s_2|\right)\right) \\
& \times \exp\left(-\frac{\hat{q}_1^2\tilde{\xi}_1^2 + (\hat{q}_0 + \hat{q}_1)^2\tilde{\xi}_2^2}{2}\right) \\
= & \delta_{\mathbf{q}_{0\parallel},\mathbf{0}} \exp\left(-\hat{q}_0^2\ell^2|s_0 - s_2| - \frac{\hat{q}_0^2\tilde{\xi}_2^2}{2}\right) \sum_{\hat{q}_1} \delta_{\mathbf{q}_{1\parallel},\mathbf{0}} \exp\left(-\hat{q}_1^2\left(\ell^2|s_1 - s_2| + \frac{\tilde{\xi}_1^2 + \tilde{\xi}_2^2}{2}\right)\right) \\
& \times \exp\left(\hat{q}_0\hat{q}_1\left(\ell^2(|s_0 - s_1| - |s_0 - s_2| - |s_1 - s_2|) - \tilde{\xi}_2^2\right)\right) \quad (\text{C.51})
\end{aligned}$$

Now let us look at the sum $\sum_{\hat{q}_1}$: it runs over elements $\hat{q}_1 = \frac{2\pi}{A^{1/D}} \hat{n}$ with $\hat{n} \in \mathbb{Z}^{D(n+1)}$. As already noted in Sec. B.5.3 on page 138, it approaches a Riemann sum in the thermodynamic limit $A \rightarrow \infty$. There we changed

$$\sum_{\hat{q}_1}(\dots) \longrightarrow \frac{A^{n+1}}{(2\pi)^{D(n+1)}} \int d\hat{q}(\dots) . \quad (\text{C.52})$$

Here, we additionally have to deal with the $\delta_{\mathbf{q}_{1\parallel},\mathbf{0}}$ -constraint. To include it into the transformation, we can write it as an integral over an exponential. For any function $\mathcal{A}(\hat{q})$:

$$\begin{aligned}
\sum_{\hat{q}} \delta_{\mathbf{q}_{1\parallel},\mathbf{0}} \mathcal{A}(\hat{q}) &= \frac{1}{A} \int_A d\mathbf{x} \sum_{\hat{q}} \mathcal{A}(\hat{q}) e^{i(\mathbf{q}^{(0)} + \dots + \mathbf{q}^{(n)})\mathbf{x}} \\
&\longrightarrow \frac{1}{A} \frac{A^{n+1}}{(2\pi)^{D(n+1)}} \int_{\mathbb{R}^D} d\mathbf{x} \int d\hat{q} \mathcal{A}(\hat{q}) e^{i(\mathbf{q}^{(0)} + \dots + \mathbf{q}^{(n)})\mathbf{x}} \\
&= \frac{A^n}{(2\pi)^{Dn}} \int d\hat{q} \delta(\mathbf{q}^{(0)} + \dots + \mathbf{q}^{(n)}) \mathcal{A}(\hat{q}) = \frac{A^n}{(2\pi)^{Dn}} \int dq_{\perp} \mathcal{A}(q_{\perp}) \quad (\text{C.53})
\end{aligned}$$

Applying this relation to the sum over \hat{q}_1 in Eq. (C.51), we just have to deal with

a simple Gaussian integral:

$$\begin{aligned}
I_2 &= \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \exp\left(-\hat{q}_0^2 \ell^2 |s_0 - s_2| - \frac{\hat{q}_0^2 \tilde{\xi}_2^2}{2}\right) \\
&\quad \times \frac{A^n}{(2\pi)^{Dn}} \int dq_{1\perp} \exp\left(-q_{1\perp}^2 \left(\ell^2 |s_1 - s_2| + \frac{\tilde{\xi}_1^2 + \tilde{\xi}_2^2}{2}\right)\right) \\
&\quad \times \exp\left(q_{0\perp} q_{1\perp} \left(\ell^2 (|s_0 - s_1| - |s_0 - s_2| - |s_1 - s_2|) - \tilde{\xi}_2^2\right)\right) \\
&= \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \exp\left(-\hat{q}_0^2 \ell^2 |s_0 - s_2| - \frac{\hat{q}_0^2 \tilde{\xi}_2^2}{2}\right) \frac{A^n}{(2\pi)^{Dn}} \left(\frac{2\pi}{\tilde{\xi}_1^2 + \tilde{\xi}_2^2 + 2\ell^2 |s_1 - s_2|}\right)^{Dn/2} \\
&\quad \times \exp\left(\frac{q_{0\perp}^2 \left(\ell^2 (|s_0 - s_1| - |s_0 - s_2| - |s_1 - s_2|) - \tilde{\xi}_2^2\right)^2}{2(\tilde{\xi}_1^2 + \tilde{\xi}_2^2 + 2\ell^2 |s_1 - s_2|)}\right) \\
&= \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \left(\frac{A}{(2\pi)^D} \left(\frac{2\pi}{\tilde{\xi}_1^2 + \tilde{\xi}_2^2 + 2\ell^2 |s_1 - s_2|}\right)^{D/2}\right)^n \\
&\quad \times \exp\left(\frac{q_{0\perp}^2}{2} \left\{ \frac{(\ell^2 g(\mathbf{s}) - \tilde{\xi}_2^2)^2}{\tilde{\xi}_1^2 + \tilde{\xi}_2^2 + 2\ell^2 |s_1 - s_2|} - 2\ell^2 |s_0 - s_2| - \tilde{\xi}_2^2 \right\}\right), \quad (\text{C.54})
\end{aligned}$$

where we made the simplifying definition:

$$g(\mathbf{s}) \equiv g(s_0, s_1, s_2) := |s_0 - s_1| - |s_0 - s_2| - |s_1 - s_2| \quad (\text{C.55})$$

Now we can recompose $I(\hat{q}_0, s_0)$, Eq. (C.49), using the expressions (C.50) and (C.54). We also neglect all terms of the form $(\dots)^n = 1 + \mathcal{O}(n)$, and hence only keep the lowest order in n :

$$\begin{aligned}
I(\hat{q}_0, s_0) &= \delta_{\hat{q}_0, \hat{0}} + \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \mu^2 Q \int_0^1 ds_1 \int_{\xi_1^2, s_1} \exp\left(-\frac{q_{0\perp}^2}{2} \left(2\ell^2 |s_0 - s_1| + \tilde{\xi}_1^2\right)\right) \\
&\quad + \delta_{\mathbf{a}_{0\parallel}, \mathbf{0}} \frac{(\mu^2 Q)^2}{2} \int_0^1 ds_1 ds_2 \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} \\
&\quad \exp\left(\frac{q_{0\perp}^2}{2} \left\{ \frac{(\ell^2 g(\mathbf{s}) - \tilde{\xi}_2^2)^2}{\tilde{\xi}_1^2 + \tilde{\xi}_2^2 + 2\ell^2 |s_1 - s_2|} - 2\ell^2 |s_0 - s_2| - \tilde{\xi}_2^2 \right\}\right) + \mathcal{O}(Q^3) \quad (\text{C.56})
\end{aligned}$$

and with that:

$$\begin{aligned}
I(\hat{0}, 0) &= 1 + \mu^2 Q \int_0^1 ds_1 \int_{\xi_1^2, s_1} 1 + \frac{(\mu^2 Q)^2}{2} \int_0^1 ds_1 ds_2 \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} 1 + \mathcal{O}(Q^3) \\
&= 1 + \mu^2 Q + \frac{(\mu^2 Q)^2}{2} + \mathcal{O}(Q^3) \quad (\text{C.57})
\end{aligned}$$

Now we can plug these two results for $I(\hat{q}_0, s_0)$ and $I(\hat{0}, 0)$ back into the original

saddle point equation (3.46):

$$\begin{aligned}
& \left((1-Q)\delta_{\hat{q}_0, \hat{0}} + Q\delta_{\mathbf{q}_{0\parallel}, \mathbf{0}} \int_{\xi^2, s_0} \exp\left(-\frac{\hat{q}_0^2 \xi^2}{2}\right) \right) \left(1 + \mu^2 Q + \frac{(\mu^2 Q)^2}{2} \right) \\
&= \delta_{\hat{q}_0, \hat{0}} + \delta_{\mathbf{q}_{0\parallel}, \mathbf{0}} \mu^2 Q \int_0^1 ds_1 \int_{\xi_1^2, s_1} \exp\left(-\frac{q_{0\perp}^2}{2} (2\ell^2 |s_0 - s_1| + \tilde{\xi}_1^2)\right) \\
&+ \delta_{\mathbf{q}_{0\parallel}, \mathbf{0}} \frac{(\mu^2 Q)^2}{2} \int_0^1 ds_1 ds_2 \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} \\
&\exp\left(\frac{q_{0\perp}^2}{2} \left\{ \frac{(\ell^2 g(\mathbf{s}) - \tilde{\xi}_2^2)^2}{\tilde{\xi}_1^2 + \tilde{\xi}_2^2 + 2\ell^2 |s_1 - s_2|} - 2\ell^2 |s_0 - s_2| - \tilde{\xi}_2^2 \right\}\right) + \mathcal{O}(Q^3). \quad (\text{C.58})
\end{aligned}$$

This equation must be valid for any choice of \hat{q}_0 and s_0 . Let us look at different cases:

For $\hat{q}_0 = \hat{0}$, we get the trivial relation

$$\begin{aligned}
(1-Q) + Q &= 1 \\
1 &= 1 \quad (\text{C.59})
\end{aligned}$$

as one can already see at the saddle point equation (3.46). For $\hat{q}_0 \neq \hat{0}$ and $\mathbf{q}_{0\parallel} \neq \mathbf{0}$, all terms in Eq. (C.58) vanish due to the δ -constraints and we also get the trivial result $0 = 0$.

Hence, from here, we restrict ourselves to the more interesting case $\hat{q}_0 \neq \hat{0}$ and $\mathbf{q}_{0\parallel} = \mathbf{0}$. Furthermore, to be consistent with the terms on the left hand side, we also insert the substitution $\tilde{\xi}^2 = \xi^2 + a^2$ (see Eq. (3.49)) on the left hand side of Eq. (C.58), again:

$$\begin{aligned}
& Q \int_{\xi^2, s_0} \exp\left(-\frac{q_{0\perp}^2 \xi^2}{2}\right) \left(1 + \mu^2 Q + \frac{(\mu^2 Q)^2}{2} \right) \\
&= \mu^2 Q \int_0^1 ds_1 \int_{\xi_1^2, s_1} \exp\left(-\frac{q_{0\perp}^2}{2} (2\ell^2 |s_0 - s_1| + a^2 + \xi_1^2)\right) \\
&+ \frac{(\mu^2 Q)^2}{2} \int_0^1 ds_1 ds_2 \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} \\
&\exp\left(\frac{q_{0\perp}^2}{2} \left\{ \frac{(\ell^2 g(\mathbf{s}) - a^2 - \xi_2^2)^2}{\xi_1^2 + \xi_2^2 + 2a^2 + 2\ell^2 |s_1 - s_2|} - 2\ell^2 |s_0 - s_2| - a^2 - \xi_2^2 \right\}\right) \\
&+ \mathcal{O}(Q^3). \quad (\text{C.60})
\end{aligned}$$

The canceled term in the first line is of the order Q^3 and thus can be neglected.

C.5 Obtaining the Equation for the Localization Length

C.5.1 Normalization of Length Scales

As anticipated in Sec. 3.7.1, we can simplify the saddle point equation by measuring all lengths in units of ℓ , *i.e.* define:

$$\xi_\ell^2 := \xi^2 / \ell^2 \quad (\text{C.61a})$$

$$a_\ell^2 := a^2 / \ell^2 \quad (\text{C.61b})$$

$$q_{0\ell\perp}^2 := \ell^2 q_{0\perp}^2 \quad (\text{C.61c})$$

With the new variables, Eq. (3.51) does not exhibit an explicit dependence on ℓ :

$$\begin{aligned} & (1 + \mu^2 Q) \int_{\xi^2, s_0} \exp\left(-\frac{q_{0\ell\perp}^2 \xi_\ell^2}{2}\right) \\ &= \mu^2 \int_0^1 ds_1 \int_{\xi_1^2, s_1} \exp\left(-\frac{q_{0\ell\perp}^2}{2} (2|s_0 - s_1| + a_\ell^2 + \xi_{1\ell}^2)\right) \\ & \quad + \frac{\mu^4 Q}{2} \int_0^1 ds_1 ds_2 \int_{\xi_1^2, s_1} \int_{\xi_2^2, s_2} \\ & \quad \exp\left(\frac{q_{0\ell\perp}^2}{2} \left\{ \frac{(g(\mathbf{s}) - a_\ell^2 - \xi_{2\ell}^2)^2}{\xi_{1\ell}^2 + \xi_{2\ell}^2 + 2a_\ell^2 + 2|s_1 - s_2|} - 2|s_0 - s_2| - a_\ell^2 - \xi_{2\ell}^2 \right\}\right) \\ & \quad + \mathcal{O}(Q^2). \end{aligned} \quad (\text{C.62})$$

Now we normalize the localization length with ε and therefore, introduce the new variable

$$\theta := \frac{f}{\varepsilon \xi_\ell^2} = \frac{f \ell^2}{(\mu^2 - 1) \xi^2} \quad \Longleftrightarrow \quad \xi_\ell^2 = \frac{f}{\varepsilon \theta}, \quad (\text{C.63a})$$

$$\text{with } f = \frac{2}{3} + a_\ell^2 = \frac{2}{3} + \frac{a^2}{\ell^2} \quad (\text{C.63b})$$

which – as we will see – remains of the order $\mathcal{O}(1)$, when we approach the transition point. Since the localization lengths are very large, we also normalize $q_{0\ell\perp}^2$ with ε , so that we have to deal with variables of the order $\mathcal{O}(1)$. Hence we define:

$$t := \frac{f q_{0\ell\perp}^2}{2\varepsilon} \quad \Longleftrightarrow \quad \frac{q_{0\ell\perp}^2}{2} = \frac{t\varepsilon}{f} \quad (\text{C.64})$$

With that Eq. (C.62) becomes:

$$\begin{aligned}
& (1 + \mu^2 Q) \int_{\theta, s_0} \exp\left(-\frac{t}{\theta}\right) \\
&= \mu^2 \int_0^1 ds_1 \int_{\theta_1, s_1} \exp\left(-t \left\{ \frac{1}{\theta_1} + \frac{2\varepsilon}{f} |s_0 - s_1| + \frac{\varepsilon}{f} a_\ell^2 \right\}\right) \\
&\quad + \frac{\mu^4 Q}{2} \int_0^1 ds_1 ds_2 \int_{\theta_1, s_1} \int_{\theta_2, s_2} \\
&\quad \exp\left(\frac{t}{f} \left\{ \frac{(\varepsilon g(\mathbf{s}) - \varepsilon a_\ell^2 - f\theta_2^{-1})^2}{f\theta_1^{-1} + f\theta_2^{-1} + 2\varepsilon a_\ell^2 + 2\varepsilon |s_1 - s_2|} - 2\varepsilon |s_0 - s_2| - \varepsilon a_\ell^2 - f\theta_2^{-1} \right\}\right) \\
&\quad + \mathcal{O}(Q^2). \tag{C.65}
\end{aligned}$$

Here we also turned the probability density $\mathcal{P}(\xi^2, s)$ for ξ^2 into the probability density $\pi(\theta, s)$ for θ :

$$\begin{aligned}
\int_{\xi^2, s} &= \int_0^\infty d\xi^2 \mathcal{P}(\xi^2, s) \longrightarrow \int_{\theta, s} = \int_0^\infty d\theta \pi(\theta, s). \tag{C.66} \\
\text{with } d\xi^2 \mathcal{P}(\xi^2, s) &= d\theta \pi(\theta, s)
\end{aligned}$$

Now, we can sort the saddle point equation (C.65) in orders of ε , and have to keep in mind that $Q = \mathcal{O}(\varepsilon)$, as found in Eq. (3.58).

For the three exponentials, we can expand all expressions which are of the order $\mathcal{O}(\varepsilon)$: In doing so, we note that the last term on the right hand side of Eq. (C.65) has a prefactor Q and thus already *is* of the order $\mathcal{O}(\varepsilon)$. Hence, we only need to expand the exponential up to zeroth order, *i.e.* neglect all terms of the order $\mathcal{O}(\varepsilon)$:

$$\begin{aligned}
& (1 + \mu^2 Q) \int_{\theta, s_0} \exp\left(-\frac{t}{\theta}\right) \tag{C.67} \\
&= \mu^2 \int_0^1 ds_1 \int_{\theta_1, s_1} \exp\left(-\frac{t}{\theta_1}\right) \left(1 - t \left\{ \frac{2\varepsilon}{f} |s_0 - s_1| + \frac{\varepsilon}{f} a_\ell^2 \right\}\right) \\
&\quad + \frac{\mu^4 Q}{2} \int_0^1 ds_1 ds_2 \int_{\theta_1, s_1} \int_{\theta_2, s_2} \exp\left(\frac{t}{f} \left\{ \frac{(f\theta_2^{-1})^2}{f\theta_1^{-1} + f\theta_2^{-1}} - f\theta_2^{-1} \right\}\right) + \mathcal{O}(\varepsilon^2).
\end{aligned}$$

Here, the argument of the last exponential can be simplified:

$$\begin{aligned}
\frac{t}{f} \left\{ \frac{(f\theta_2^{-1})^2}{f\theta_1^{-1} + f\theta_2^{-1}} - f\theta_2^{-1} \right\} &= t \left\{ \frac{(\theta_2^{-1})^2}{\theta_1^{-1} + \theta_2^{-1}} - \theta_2^{-1} \right\} = t \frac{\cancel{(\theta_2^{-1})^2} - \theta_2^{-1}\theta_1^{-1} - \cancel{(\theta_2^{-1})^2}}{\theta_1^{-1} + \theta_2^{-1}} \\
&= -t \frac{1}{\theta_1 + \theta_2}. \tag{C.68}
\end{aligned}$$

With that the saddle point equation (C.67) becomes:

$$\begin{aligned}
& (1 + \mu^2 Q) \int_{\theta_1, s_0} \exp\left(-\frac{t}{\theta_1}\right) \\
&= \mu^2 \int_0^1 ds_1 \int_{\theta_1, s_1} \exp\left(-\frac{t}{\theta_1}\right) \left(1 - 2t \frac{\varepsilon}{f} \left\{|s_0 - s_1| + \frac{1}{2} a_\ell^2\right\}\right) \\
& \quad + \frac{\mu^4 Q}{2} \int_0^1 ds_1 ds_2 \int_{\theta_1, s_1} \int_{\theta_2, s_2} \exp\left(-\frac{t}{\theta_1 + \theta_2}\right) + \mathcal{O}(\varepsilon^2). \tag{C.69}
\end{aligned}$$

C.5.2 Laplace-Transformation of the Saddle Point Equation

Eq. (C.69) could in principle be used to determine the distribution $\pi(\theta)$. When calculating the actual shape of this distribution, however, is much more convenient, if it has the form of a differential (or integro-differential) equation, and does not involve an external variable t . Therefore, as a next step, we try to convert the three exponential functions to delta functions with the respective θ as argument. This is done by a Laplace transformation: We multiply the whole equation with $\sin(at) \exp(t\theta^{-1})$, integrate over t and take the limit $a \rightarrow 0$. Then we can use the following relations, which are valid for any $\Theta > 0$:

$$\begin{aligned}
& \lim_{a \rightarrow 0} \int_0^\infty dt \sin(at) \exp\left(\frac{t}{\theta}\right) \exp\left(-\frac{t}{\Theta}\right) = \lim_{a \rightarrow 0} \frac{a}{a^2 + (\Theta^{-1} - \theta^{-1})^2} \\
&= \delta(\Theta^{-1} - \theta^{-1}) \tag{C.70a}
\end{aligned}$$

and

$$\begin{aligned}
& - \lim_{a \rightarrow 0} \int_0^\infty dt t \cdot \sin(at) \exp\left(\frac{t}{\theta}\right) \exp\left(-\frac{t}{\Theta}\right) \\
&= \lim_{a \rightarrow 0} \frac{\partial}{\partial \Theta^{-1}} \int_0^\infty dt \sin(at) \exp\left(\frac{t}{\theta}\right) \exp\left(-\frac{t}{\Theta}\right) \\
&= \lim_{a \rightarrow 0} \frac{\partial}{\partial \Theta^{-1}} \frac{a}{a^2 + (\Theta^{-1} - \theta^{-1})^2} \\
&= \delta'(\Theta^{-1} - \theta^{-1}) \tag{C.70b}
\end{aligned}$$

Now, we apply those two relations to Eq. (C.69). In doing so, $\Theta = \theta_1$ in the first two terms, and $\Theta = \theta_1 + \theta_2$ in the last term:

$$\begin{aligned}
& (1 + \mu^2 Q) \int_0^\infty d\theta_1 \pi(\theta_1, s_0) \delta(\theta_1^{-1} - \theta^{-1}) \\
&= \mu^2 \int_0^1 ds_1 \int_0^\infty d\theta_1 \pi(\theta_1, s_1) \left(\delta(\theta_1^{-1} - \theta^{-1}) + \frac{2\varepsilon}{f} \delta'(\theta_1^{-1} - \theta^{-1}) \left\{|s_0 - s_1| + \frac{1}{2} a_\ell^2\right\} \right) \\
& \quad + \frac{\mu^4 Q}{2} \int_0^1 ds_1 ds_2 \int_0^\infty d\theta_1 \pi(\theta_1, s_1) \int_0^\infty d\theta_2 \pi(\theta_2, s_2) \delta\left((\theta_1 + \theta_2)^{-1} - \theta^{-1}\right) \\
& \quad + \mathcal{O}(\varepsilon^2). \tag{C.71}
\end{aligned}$$

Now we want to integrate over θ_1 in the first two terms and over θ_2 in the last term (marked in blue). Therefore we need to use the following relations for δ -functions:

$$\int d\theta_1 \pi(\theta_1) \delta(g(\theta_1)) = \sum_{\nu} \frac{\pi(\theta_{1\nu})}{|g'(\theta_{1\nu})|} \quad (\text{C.72a})$$

$$\int d\theta_1 \pi(\theta_1) \delta'(g(\theta_1)) = \sum_{\nu} \frac{\pi(\theta_{1\nu})g''(\theta_{1\nu}) - \pi'(\theta_{1\nu})g'(\theta_{1\nu})}{|g'(\theta_{1\nu})|^3}, \quad (\text{C.72b})$$

where $\theta_{1\nu}$ are the roots and $g'(\theta_{1\nu})$ and $g''(\theta_{1\nu})$ the derivatives of the function $g(\theta_1)$, which we now find for the presented case:

- **In the first two terms:**

$$\begin{aligned} g(\theta_1) &= \theta_1^{-1} - \theta^{-1} \stackrel{!}{=} 0 \\ \Rightarrow \text{root: } \theta_{1,0} &= \theta \\ \text{derivatives: } g'(\theta_1) &= -\theta_1^{-2} \\ g''(\theta_1) &= 2\theta_1^{-3} \end{aligned} \quad (\text{C.73a})$$

- **In the last term:**

$$\begin{aligned} g(\theta_2) &= (\theta_1 + \theta_2)^{-1} - \theta^{-1} \stackrel{!}{=} 0 \\ \Leftrightarrow \theta &\stackrel{!}{=} \theta_1 + \theta_2 \\ \Rightarrow \text{root: } \begin{cases} \theta_{2,0} = \theta - \theta_1 & \text{for } \theta \geq \theta_1 \\ \text{no root} & \text{for } \theta < \theta_1 \end{cases} \\ \text{derivative: } g'(\theta_2) &= -(\theta_1 + \theta_2)^{-2}, \quad g'(\theta_{2,0}) = -\theta^{-2} \end{aligned} \quad (\text{C.73b})$$

We plug Eqs. (C.72) with the found root and derivatives (C.73) back into the saddle point equation (C.71):

$$\begin{aligned} & (1 + \mu^2 Q) \frac{\pi(\theta, s_0)}{|-\theta^{-2}|} \\ &= \mu^2 \int_0^1 ds_1 \left(\frac{\pi(\theta, s_1)}{|-\theta^{-2}|} + \frac{2\varepsilon}{f} \frac{\pi(\theta, s_1) \cdot 2\theta^{-3} + \partial_\theta \pi(\theta, s_1) \cdot \theta^{-2}}{|-\theta^{-2}|^3} \left\{ |s_0 - s_1| + \frac{1}{2} a_\ell^2 \right\} \right) \\ & \quad + \frac{\mu^4 Q}{2} \int_0^1 ds_1 ds_2 \int_0^\infty d\theta_1 \pi(\theta_1, s_1) \frac{\pi(\theta - \theta_1, s_2)}{|-\theta^{-2}|} + \mathcal{O}(\varepsilon^2), \end{aligned} \quad (\text{C.74})$$

where $\partial_\theta \pi(\theta, s_1)$ is the derivative of $\pi(\theta, s_1)$ with respect to θ . We now multiply this equation with θ^{-2} on both sides, plug in $\mu^2 = 1 + \varepsilon$ and $Q = 2\varepsilon + \mathcal{O}(\varepsilon^2)$, Eq. (3.58), and get:

$$\begin{aligned} & (1 + 2\varepsilon) \pi(\theta, s_0) \\ &= (1 + \varepsilon) \int_0^1 ds_1 \left(\pi(\theta, s_1) + \frac{2\varepsilon}{f} \left(2\theta \pi(\theta, s_1) + \theta^2 \partial_\theta \pi(\theta, s_1) \right) \left\{ |s_0 - s_1| + \frac{1}{2} a_\ell^2 \right\} \right) \\ & \quad + \varepsilon \int_0^1 ds_1 ds_2 \int_0^\infty d\theta_1 \pi(\theta_1, s_1) \pi(\theta - \theta_1, s_2) + \mathcal{O}(\varepsilon^2), \end{aligned} \quad (\text{C.75})$$

and by substituting $s_0 \rightarrow s$:

$$\begin{aligned}
& (1 + 2\varepsilon)\pi(\theta, s) \\
&= (1 + \varepsilon) \int_0^1 ds_1 \pi(\theta, s_1) + \frac{\varepsilon}{f} \int_0^1 ds_1 \partial_\theta \left(\theta^2 \pi(\theta, s_1) \right) \left\{ 2|s - s_1| + a_\ell^2 \right\} \\
&\quad + \varepsilon \int_0^1 ds_1 ds_2 \int_0^\theta d\theta_1 \pi(\theta_1, s_1) \pi(\theta - \theta_1, s_2) + \mathcal{O}(\varepsilon^2) .
\end{aligned}$$

(C.76)

In Sec. 3.7 we examine this integro-differential equation and display the actual shape of $\pi(\theta, s)$.

C.6 Calculation of a Correlator

Throughout chapter 3 about cross-linked directed polymers we need expressions of the form:

$$\begin{aligned}
& \left\langle \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}(z_j) \right) \right\rangle_{n+1}^{H_{\text{align}} + H_{\text{tilt}}} = \int \mathcal{D}\hat{r}(z) e^{-H_{\text{align},1}^{(n+1)} - H_{\text{tilt},1}^{(n+1)}} \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}(z_j) \right) = \\
& \frac{\int d\hat{r}_1 \cdots d\hat{r}_{Z_L} \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}_{Z_j} - \frac{\sigma \Delta z}{2} \sum_{Z=2}^{Z_L} \frac{(\hat{r}_Z - \hat{r}_{Z-1})^2}{\Delta z^2} - \Delta z \sum_{Z=2}^{Z_L} \frac{\hat{h}_Z (\hat{r}_Z - \hat{r}_{Z-1})}{\Delta z} \right)}{\int d\hat{r}_1 \cdots d\hat{r}_{Z_L} \exp \left(-\frac{\sigma \Delta z}{2} \sum_{Z=2}^{Z_L} \frac{(\hat{r}_Z - \hat{r}_{Z-1})^2}{\Delta z^2} \right)}
\end{aligned}$$

(C.77)

First, let us look at the denominator: we can keep \hat{r}_1 , but make the substitution $\hat{t}_Z = \hat{r}_Z - \hat{r}_{Z-1}$ for $Z = 2, \dots, Z_L$, and write:

$$\begin{aligned}
\mathcal{D} &= \int d\hat{r}_1 \cdots d\hat{r}_{Z_L} \exp \left(-\frac{\sigma \Delta z}{2} \sum_{Z=2}^{Z_L} \frac{(\hat{r}_Z - \hat{r}_{Z-1})^2}{\Delta z^2} \right) \\
&= \int d\hat{r}_1 d\hat{t}_2 \cdots d\hat{t}_{Z_L} \exp \left(-\frac{\sigma \Delta z}{2} \sum_{Z=2}^{Z_L} \frac{\hat{t}_Z^2}{\Delta z^2} \right) \\
&= A^{n+1} \left(\int d\hat{t} \exp \left(-\frac{\sigma}{2\Delta z} \hat{t}^2 \right) \right)^{Z_L-1} \\
&= A^{n+1} \left(\frac{2\pi \Delta z}{\sigma} \right)^{D(n+1)(Z_L-1)}
\end{aligned}$$

(C.78)

In the numerator we make the same substitution and get:

$$\begin{aligned}
\mathcal{N} &= \\
&\int d\hat{r}_1 \cdots d\hat{r}_{Z_L} \exp\left(i \sum_{j=1}^J \hat{q}_j \hat{r}_{Z_j} - \frac{\sigma \Delta z}{2} \sum_{Z=2}^{Z_L} \frac{(\hat{r}_Z - \hat{r}_{Z-1})^2}{\Delta z^2} - \Delta z \sum_{Z=2}^{Z_L} \frac{\hat{h}_Z (\hat{r}_Z - \hat{r}_{Z-1})}{\Delta z}\right) \\
&= \int d\hat{r}_1 d\hat{t}_2 \cdots d\hat{t}_{Z_L} \exp\left(i \sum_{j=1}^J \hat{q}_j (\hat{r}_1 + \hat{t}_2 + \dots + \hat{t}_{Z_j}) - \frac{\sigma}{2\Delta z} \sum_{Z=2}^{Z_L} \hat{t}_Z^2 - \sum_{Z=2}^{Z_L} \hat{h}_Z \hat{t}_Z\right) \\
&= \int d\hat{r}_1 \exp\left(i \hat{r}_1 \sum_{j=1}^J \hat{q}_j\right) \\
&\quad \times \int d\hat{t}_2 \cdots d\hat{t}_{Z_L} \exp\left(i \sum_{j=1}^J \hat{q}_j (\hat{t}_2 + \dots + \hat{t}_{Z_j}) - \frac{\sigma}{2\Delta z} \sum_{Z=2}^{Z_L} \hat{t}_Z^2 - \sum_{Z=2}^{Z_L} \hat{h}_Z \hat{t}_Z\right) \quad (\text{C.79})
\end{aligned}$$

In the last two lines, the integral over \hat{r}_1 can easily be performed, yielding $A^{n+1} \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}}$. For the integral over $\hat{t}_2, \dots, \hat{t}_{Z_L}$, we make the definition:

$$\hat{h}'_Z := \hat{h}_Z - i \sum_{j=1}^J \hat{q}_j \theta_{Z_j, Z} \quad (\text{C.80})$$

$$\text{with } \theta_{Z_j, Z} = \begin{cases} 1 & \text{if } Z_j \geq Z \\ 0 & \text{if } Z_j < Z \end{cases} \quad (\text{C.81})$$

With that the numerator \mathcal{N} can be simplified:

$$\begin{aligned}
\mathcal{N} &= A^{n+1} \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \int d\hat{t}_2 \cdots d\hat{t}_{Z_L} \exp\left(-\frac{\sigma}{2\Delta z} \sum_{Z=2}^{Z_L} \hat{t}_Z^2 - \sum_{Z=2}^{Z_L} \hat{h}'_Z \hat{t}_Z\right) \\
&= A^{n+1} \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \prod_{Z=2}^{Z_L} \left\{ \left(\frac{2\pi \Delta z}{\sigma}\right)^{D(n+1)} \exp\left(\frac{\hat{h}'_Z{}^2}{4\sigma/(2\Delta z)}\right) \right\} \\
&= A^{n+1} \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \left(\frac{2\pi \Delta z}{\sigma}\right)^{D(n+1)(Z_L-1)} \exp\left(\frac{\Delta z}{2\sigma} \sum_{Z=2}^{Z_L} \hat{h}'_Z{}^2\right) \quad (\text{C.82})
\end{aligned}$$

With that expression for the numerator and (C.78) for the denominator, we get for the original expression (C.77)

$$\begin{aligned}
& \left\langle \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}(z_j) \right) \right\rangle_{n+1}^{H_{\text{align}} + H_{\text{tilt}}} \\
&= \frac{A^{n+1} \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \left(\frac{2\pi \Delta z}{\sigma} \right)^{D(n+1)(Z_L-1)} \exp \left(\frac{\Delta z}{2\sigma} \sum_{Z=2}^{Z_L} \hat{h}'_Z \right)}{A^{n+1} \left(\frac{2\pi \Delta z}{\sigma} \right)^{D(n+1)(Z_L-1)}} \\
&= \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \exp \left(\frac{\Delta z}{2\sigma} \sum_{Z=2}^{Z_L} \hat{h}'_Z \right) \\
&= \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \exp \left(\frac{\Delta z}{2\sigma} \sum_{Z=2}^{Z_L} \left(\hat{h}_Z - i \sum_{j=1}^J \hat{q}_j \theta_{Z_j, Z} \right)^2 \right) \tag{C.83}
\end{aligned}$$

Changing back to the Riemann-integral by taking $\Delta z = L/Z_L \rightarrow 0$ we get:

$$\begin{aligned}
& \left\langle \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}(z_j) \right) \right\rangle_{n+1}^{H_{\text{align}} + H_{\text{tilt}}} \\
&= \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \exp \left(\frac{1}{2\sigma} \int_0^L dz \left(\hat{h}(z) - i \sum_{j=1}^J \hat{q}_j \theta(z_j - z) \right)^2 \right) \tag{C.84}
\end{aligned}$$

The integral in the exponential can be simplified further. Therefore we will need the following expression (under the restriction $\hat{q}_1 + \dots + \hat{q}_J = \hat{0}$):

$$\begin{aligned}
& \int_0^L dz \left(\sum_{j=1}^J \hat{q}_j \theta(z_j - z) \right)^2 \\
&= \int_0^L dz \left(\sum_{j=1}^J \hat{q}_j^2 \theta(z_j - z) + \sum_{j \neq k} \hat{q}_j \hat{q}_k \theta(z_j - z) \theta(z_k - z) \right) \\
&= \int_0^L dz \left(- \sum_{j \neq k} \hat{q}_j \hat{q}_k \theta(z_j - z) + \sum_{j \neq k} \hat{q}_j \hat{q}_k \theta(z_j - z) \theta(z_k - z) \right) \\
&= \int_0^L dz \left(- \sum_{j \neq k} \hat{q}_j \hat{q}_k \theta(z_j - z) (1 - \theta(z_k - z)) \right) \\
&= - \sum_{j < k} \hat{q}_j \hat{q}_k |z_j - z_k| \tag{C.85}
\end{aligned}$$

Despite the minus-sign, this expression is always ≥ 0 , as it can be seen in the first line. With that expression we can calculate the integral in expression (C.84):

$$\begin{aligned}
& \int_0^L dz \left(\hat{h}(z) - i \sum_{j=1}^J \hat{q}_j \theta(z_j - z) \right)^2 \\
&= \int_0^L dz \hat{h}^2(z) - 2i \sum_{j=1}^J \hat{q}_j \int_0^L dz \hat{h}(z) \theta(z_j - z) - \int_0^L dz \left(\sum_{j=1}^J \hat{q}_j \theta(z_j - z) \right)^2 \\
&= LH^2 - 2i \sum_{j=1}^J \hat{q}_j \hat{H}(z_j) + \sum_{j < k} \hat{q}_j \hat{q}_k |z_j - z_k|, \tag{C.86a}
\end{aligned}$$

where

$$\boxed{\hat{H}(z) := \int_0^z dz' \hat{h}(z')} \quad \text{and} \quad \boxed{H^2 := \frac{1}{L} \int_0^L dz \hat{h}^2(z)} \tag{C.86b}$$

Now, we plug that form back into Eq. (C.84) and get the result:

$$\boxed{
\begin{aligned}
& \left\langle \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}(z_j) \right) \right\rangle_{n+1}^{H_{\text{align}} + H_{\text{tilt}}} \\
&= \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \exp \left(\frac{1}{2\sigma} \left(LH^2 - 2i \sum_{j=1}^J \hat{q}_j \hat{H}(z_j) + \sum_{j < k} \hat{q}_j \hat{q}_k |z_j - z_k| \right) \right)
\end{aligned}
} \tag{C.87}$$

without the tilting interaction, *i.e.* $\hat{h}(z) \equiv \hat{0}$, the result simplifies to:

$$\boxed{
\left\langle \exp \left(i \sum_{j=1}^J \hat{q}_j \hat{r}(z_j) \right) \right\rangle_{n+1}^{H_{\text{align}}} = \delta_{\hat{q}_1 + \dots + \hat{q}_J, \hat{0}} \exp \left(\frac{1}{2\sigma} \sum_{j < k} \hat{q}_j \hat{q}_k |z_j - z_k| \right)
} \tag{C.88}$$

C.7 The Average Cross-Link Density

Now we calculate the disorder-averaged number of cross-links $[M]$ and its variance for the directed polymer model. We calculate expectation values of the form $[M(M-1) \cdots (M-J+1)] =: \mathcal{M}_J$, in analogy to the RLP model, Sec. 2.9.1. The difference is that the disorder average, defined in Eq. (3.22), additionally involves the integrals over the cross-link heights z_e . For simplification we define

$\mathcal{D}\Gamma = \mathcal{D}\mathbf{r}_1(z) \cdots \mathcal{D}\mathbf{r}_N(z)$ and get:

$$\begin{aligned}
\mathcal{M}_J &= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_M=1 \\ j_1, \dots, j_M=1}}^N \frac{(\tilde{\mu}^2)^M}{M! L^M} \int_0^L dz_1 \cdots dz_M Z(\mathcal{C}) \cdot M(M-1) \cdots (M-J+1) \\
&= \frac{1}{\mathcal{Z}_1} \sum_{M=J}^{\infty} \sum_{\substack{i_1, \dots, i_M=1 \\ j_1, \dots, j_M=1}}^N \frac{(\tilde{\mu}^2/L)^M}{M!} \frac{M!}{(M-J)!} \int_0^L dz_1 \cdots dz_M Z(\{i_e, j_e, z_e\}_{e=1}^M) \\
&= \frac{1}{\mathcal{Z}_1} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_{M+J}=1 \\ j_1, \dots, j_{M+J}=1}}^N \frac{(\tilde{\mu}^2/L)^{M+J}}{(M+J-J)!} \int_0^L dz_1 \cdots dz_{M+J} Z(\{i_e, j_e, z_e\}_{e=1}^{M+J}) \\
&= \frac{1}{\mathcal{Z}_1} \int \mathcal{D}\Gamma e^{-H_{\text{ev}} - H_{\text{align}}} \sum_{M=0}^{\infty} \sum_{\substack{i_1, \dots, i_{M+J}=1 \\ j_1, \dots, j_{M+J}=1}}^N \frac{(\tilde{\mu}^2/L)^{M+J}}{M!} \\
&\quad \int_0^L dz_1 \cdots dz_{M+J} \exp\left(-\frac{1}{2a^2} \sum_{e=1}^{M+J} (\mathbf{r}_{i_e}(z_e) - \mathbf{r}_{j_e}(z_e))^2\right) \\
&= \frac{1}{\mathcal{Z}_1} \int \mathcal{D}\Gamma e^{-H_{\text{ev}} - H_{\text{align}}} \sum_{M=0}^{\infty} \frac{1}{M!} \\
&\quad \times \left(\frac{\tilde{\mu}^2}{L} \sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{1}{2a^2} (\mathbf{r}_i(z) - \mathbf{r}_j(z))^2\right) \right)^{M+J} \\
&= \frac{(\tilde{\mu}^2)^J}{\mathcal{Z}_1} \left(\frac{\partial}{\partial \tilde{\mu}^2} \right)^J \int \mathcal{D}\Gamma e^{-H_{\text{ev}} - H_{\text{align}}} \\
&\quad \times \exp\left(\frac{\tilde{\mu}^2}{L} \sum_{i,j=1}^N \int_0^L dz \exp\left(-\frac{1}{2a^2} (\mathbf{r}_i(z) - \mathbf{r}_j(z))^2\right) \right). \tag{C.89}
\end{aligned}$$

As we can see, the expression after the partial derivative is $\lim_{n \rightarrow 0} \mathcal{Z}_{n+1} = \mathcal{Z}_1$. Hence we get an analogous result to Eq. (B.109):

$$\mathcal{M}_J = \frac{(\mu^2)^J}{\mathcal{Z}_1} \left(\frac{\partial}{\partial \mu^2} \right)^J \mathcal{Z}_1. \tag{C.90}$$

The general form of \mathcal{Z}_{n+1} is given in Eq. (3.32a):

$$\mathcal{Z}_{n+1} = \exp(-Nf_0) \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\}) \tag{C.91}$$

Thereby the contribution f_0 , given in Eq. (3.31b), becomes in the limit $n \rightarrow 0$:

$$f_0 \stackrel{n \rightarrow 0}{=} -\frac{\mu^2}{2} + \frac{\lambda n_0 L}{2}. \tag{C.92}$$

The term f_{n+1} is given in Eq. (3.33); in the limit $n \rightarrow 0$ the sums over the HRS vanishes:

$$f_1\{\Omega\} = \lim_{n \rightarrow 0} f_{n+1}\{\Omega\} = 0 - \ln \mathfrak{z} \quad \text{with} \quad \mathfrak{z} = \int \mathcal{D}\hat{r}(z) \exp\left(-H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\}\right) \quad (\text{C.93})$$

The Hamiltonian $H_{\text{align},1}^{(n+1)}\{\hat{r}(z)\}$ should not depend on the parameter μ^2 . Therefore $f_1\{\Omega\}$ is a constant when deriving with respect to μ^2 . With these expressions Eq. (C.91) becomes in the limit $n \rightarrow 0$:

$$\mathcal{Z}_1 = \exp\left(\frac{N\mu^2}{2} + \frac{NL\lambda n_0}{2} + \text{const.}\right). \quad (\text{C.94})$$

This form together with Eq. (C.90) yields the same result for the expectation values \mathcal{M}_J as for the RLP model, Eq. (2.61):

$$\mathcal{M}_J = [M(M-1)\cdots(M-J+1)] = \left(\frac{\mu^2 N}{2}\right)^J. \quad (\text{C.95})$$

In particular, the mean cross-link density and the standard deviation are given by:

$$\boxed{\begin{aligned} \frac{[M]}{N} &= \frac{\mu^2}{2}, \\ \Delta M &= \sqrt{[M^2] - [M]^2} = \sqrt{\frac{\mu^2 N}{2}}, \end{aligned}} \quad (\text{C.96})$$

so that $\Delta M/[M] \propto N^{-1/2}$.

It is noteworthy that for these results we did not use any expressions which are only valid close to the sol-gel transition, so Eq. (C.96) holds for arbitrary μ^2 . If, however, density fluctuations were allowed during cross-linking, contributions from the one replica sector of $f_{n+1}\{\Omega\}$ in Eq. (3.32b) may change the average cross-link density.

Expressions for the Wet Granulates

D.1 Infinite Sums similar to the Exponential Function

$$\sum_{r=0}^{\infty} \frac{x^r}{r!} r = \sum_{r=1}^{\infty} \frac{x^r}{r!} r = x \sum_{r=1}^{\infty} \frac{x^{r-1}}{(r-1)!} = x \sum_{r=0}^{\infty} \frac{x^r}{r!} = xe^x \quad (\text{D.1})$$

$$\sum_{r=0}^{\infty} \frac{x^r}{r!} (r-1) = \sum_{r=0}^{\infty} \frac{x^r}{r!} r - \sum_{r=0}^{\infty} \frac{x^r}{r!} = xe^x - e^x = (x-1)e^x \quad (\text{D.2})$$

D.2 Inverting the Equation $x^{-2}e^x = t$

In section 5.3.2, we derived an equation for the evolution of the inverse dimensionless temperature x :

$$x^{-2}e^x (1 + \mathcal{O}(x^{-1})) = t, \quad (\text{D.3})$$

in the asymptotic limit $x, t \rightarrow \infty$. Here we solve this equation for x . We proceed in analogy to the inversion of a similar equation discussed in [de Bruijn, 1958, Chapter 2.4]. First, we take the logarithm on both sides of Eq. (D.3):

$$\begin{aligned} -2 \ln x + x + \ln(1 + \mathcal{O}(x^{-1})) &= \ln t \\ -2 \ln x + x + \mathcal{O}(x^{-1}) &= \tau, \end{aligned} \quad (\text{D.4})$$

with $\tau := \ln t$. Now, we transform this equation to

$$x \left(1 - 2 \frac{\ln x}{x} + \mathcal{O}(x^{-2}) \right) = \tau, \quad (\text{D.5})$$

which yields two statements:

$$1) \quad \lim_{\tau \rightarrow \infty} \left| \frac{x^{-1}}{\tau^{-1}} \right| = 1 < \infty \quad \Rightarrow \quad x^{-1} = \mathcal{O}(\tau^{-1}) \quad \text{and} \quad (\text{D.6a})$$

$$2) \quad \lim_{\tau \rightarrow \infty} \left| \frac{\ln x}{\ln \tau} \right| = 1 < \infty \quad \Rightarrow \quad \ln x = \mathcal{O}(\ln \tau) \quad (\text{D.6b})$$

Therefore in Eq. (D.4) we can replace $\mathcal{O}(\tau^{-1})$ by $\mathcal{O}(x^{-1})$ using Eq. (D.6a),

$$\boxed{x = \tau + 2 \ln x + \mathcal{O}(\tau^{-1}) .} \quad (\text{D.7})$$

Furthermore we can plug the statement (D.6b) in the main equation (D.7) and successively obtain higher and higher orders of the solution $x(t)$:

- Step 1: (D.6b) in (D.7)

$$x = \tau + \mathcal{O}(\ln \tau) + \cancel{\mathcal{O}(\tau^{-1})} = \tau \left(1 + \mathcal{O}\left(\frac{\ln \tau}{\tau}\right) \right) \quad (\text{D.8a})$$

$$\ln x = \ln \tau + \mathcal{O}\left(\frac{\ln \tau}{\tau}\right) \quad (\text{D.8b})$$

- Step 2: (D.8b) in (D.7)

$$\begin{aligned} x &= \tau + 2 \ln \tau + \mathcal{O}\left(\frac{\ln \tau}{\tau}\right) + \cancel{\mathcal{O}(\tau^{-1})} \\ &= \tau \left(1 + 2 \frac{\ln \tau}{\tau} \right) \left(1 + \mathcal{O}\left(\frac{\ln \tau}{\tau(\tau + 2 \ln \tau)}\right) \right) \end{aligned} \quad (\text{D.9a})$$

$$\begin{aligned} \ln x &= \ln \tau + \ln\left(1 + 2 \frac{\ln \tau}{\tau}\right) + \mathcal{O}\left(\frac{\ln \tau}{\tau(\tau + 2 \ln \tau)}\right) \\ &= \ln \tau + \frac{2 \ln \tau}{\tau} - \frac{1}{2} \left(\frac{2 \ln \tau}{\tau}\right)^2 + \cancel{\mathcal{O}\left(\frac{\ln \tau}{\tau}\right)^3} + \mathcal{O}\left(\frac{\ln \tau}{\tau^2}\right) \end{aligned} \quad (\text{D.9b})$$

- Step 3: (D.9b) in (D.7)

$$x = \tau + 2 \ln \tau + \frac{4 \ln \tau}{\tau} - \left(\frac{2 \ln \tau}{\tau}\right)^2 + \cancel{\mathcal{O}\left(\frac{\ln \tau}{\tau}\right)} + \cancel{\mathcal{O}\left(\frac{\ln \tau}{\tau^2}\right)} + \mathcal{O}(\tau^{-1}) \quad (\text{D.10})$$

Now, it would be possible to find an expression for $\ln x$ and plug it into Eq. (D.7), again. However that would yield additional terms of an order higher than $\mathcal{O}(\tau^{-1})$, and thus could be absorbed by $\mathcal{O}(\tau^{-1})$ in Eq. (D.7). Hence, the result

$$\boxed{x(t) = \ln t + 2 \ln \ln t + \frac{4 \ln \ln t}{\ln t} + \mathcal{O}\left(\frac{1}{\ln t}\right)} \quad (\text{D.11})$$

is the highest order for $x(t)$ that the original equation (D.3) can yield.

D.3 Computation of the radial distribution function $g(r)$

In Sec. 5.4.2.3 we consider the radial distribution function, which is the radial component of Eq. (5.45). Here, some details for the computation for a particle configuration $\{\mathbf{r}_1, \dots, \mathbf{r}_N\}$ are given.

To compute $g(r)$ for small values of r (i.e. $r \leq r^* = 6.4d$), the algorithm chooses one particle $\mathbf{p}_1 \in \{\mathbf{r}_1, \dots, \mathbf{r}_N\}$ and records the distance of *all* particles \mathbf{p}_2 closer than

r^* . Repeating this procedure for *all* particles $\mathbf{p}_1 \in \{\mathbf{r}_1, \dots, \mathbf{r}_N\}$ yields the histogram of distances between pairs of particles, from which $g(r)$ can easily be computed by dividing with $4\pi r^2$.

For increasing r , the number of pairs with distance r increases rapidly and hence not all pairs of particles can be considered anymore; Therefore, for $r > r^*$, the algorithm chooses the particles \mathbf{p}_1 and \mathbf{p}_2 at *random*, with a distance of at least r^* ; in repeating this procedure for about 10^9 initial particles \mathbf{p}_1 , the algorithm approximates the particle-distance histogram in a Monte-Carlo like fashion.

Two remarks have to be made about the system-boundaries: (i) For a pair of particles \mathbf{p}_1 and \mathbf{p}_2 , only the *closest* distance compatible with the periodic boundary conditions (in x - and y -direction) is considered. (ii) If particle \mathbf{p}_1 is close to a hard wall (in z -direction), then the probability to find a close-by particle is lower. Hence, whenever $g(r)$ is shown up to a distance $r = r_{\max}$, then particle \mathbf{p}_1 is chosen at least r_{\max} away from the hard wall, in order to avoid the influence of the boundary.

Further Results from the Spider Silk Model

Contents

| | | |
|------------|--|-----|
| E.1 | Effect of the Continuous Background | 187 |
| E.2 | Relevance of the Coherent Part of the Scattering Function | 187 |
| E.3 | Simplification of the Scattering Amplitude $A(\mathbf{q})$ | 188 |

E.1 Effect of the Continuous Background

In this section we explain why it is necessary to include the continuous background between the crystallites, introduced in section 4.2.4.

Without the background, the system has vast, unphysical density fluctuations on the length scale of the crystallite distances, resulting in a large scattering function $G(\mathbf{q})$ for small q -values. As already explained in section 4.2.4, these density fluctuations are unphysical because the space between the crystallites is filled with the amorphous matrix and water molecules. Fig. E.1 shows the scattering profiles in xy -direction with and without the continuous background. As expected, the system without the background shows a large increase of the scattering function for small q -values. The continuous background, however, acts as a low-pass filter on the scattering density and therefore annihilates the large intensities for small q .

E.2 Relevance of the Coherent Part of the Scattering Function

Here we discuss the influence of the coherent part of the scattering function of Eq. (4.20). Fig. E.2 shows a comparison of the **incoherent part**

$$G_1(\mathbf{q}) = \int \mathcal{D}\underline{D} |A(\underline{D}^T \mathbf{q})|^2, \quad (\text{E.1})$$

which is used to calculate the scattering function throughout chapter 4, and the contribution

$$G'_2(\mathbf{q}) := \left| \int \mathcal{D}\underline{D} A(\underline{D}^T \mathbf{q}) \right|^2 \quad (\text{E.2})$$

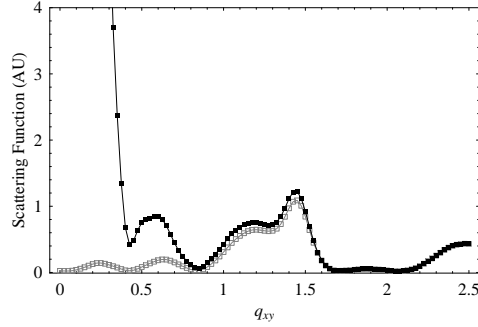


Figure E.1: Scattering function in xy -direction with (\square) and without (\blacksquare) the continuous background. Without background, density fluctuations on large length scales cause an increase of the scattering function for small q -values.

of the **coherent part**, $G_2(\mathbf{q}) = (S(\mathbf{q}) - 1) \left| \int \mathcal{D}\underline{D} A(\underline{D}^T \mathbf{q}) \right|^2$.

Neglecting the coherent part is plausible for two reasons. Firstly, because the contribution of $G'_2(\mathbf{q})$ is small compared to the incoherent part $G_1(\mathbf{q})$, as seen in the figure. And secondly, the length scale for the distances between the crystallites is much larger than atom length scales investigated here. On the length scales we are interested in, we expect $S(\mathbf{q}) \approx 1$, assuming that the crystallite positions have no long range order. Therefore, the prefactor $(S(\mathbf{q}) - 1)$ additionally reduces the contribution of the coherent term.

The (002) peak is special for the coherent scattering term $G'_2(\mathbf{q})$. All peaks except for the (002) peak have a very small contribution in $G'_2(\mathbf{q})$ because of the white average of the crystallites' rotations about the fiber axis. This makes coherent scattering from *different* crystallites very unlikely, no matter how the crystallites are arranged in space. Since there is a preferential alignment of the crystallites in the z -direction, however, contributions of coherent scattering from different crystallites (which are contained in the term $G'_2(\mathbf{q})$) are not completely destroyed; therefore, if the crystallites' distance in the z -direction is a multiple of the unit cell size a_z , causing a large contribution in the prefactor $(S(\mathbf{q}) - 1)$ at the position of the (002) peaks, a contribution of $G_2(\mathbf{q})$ will be present.

E.3 Simplification of the Scattering Amplitude $A(\mathbf{q})$

As defined in Eq. (4.16), the scattering amplitude of a single crystallite is given by

$$A(\mathbf{q}) = \sum_{\mathbf{m}=1}^{\mathbf{M}} \sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q} \cdot \mathbf{r}_{\mathbf{m},k}). \quad (\text{E.3})$$

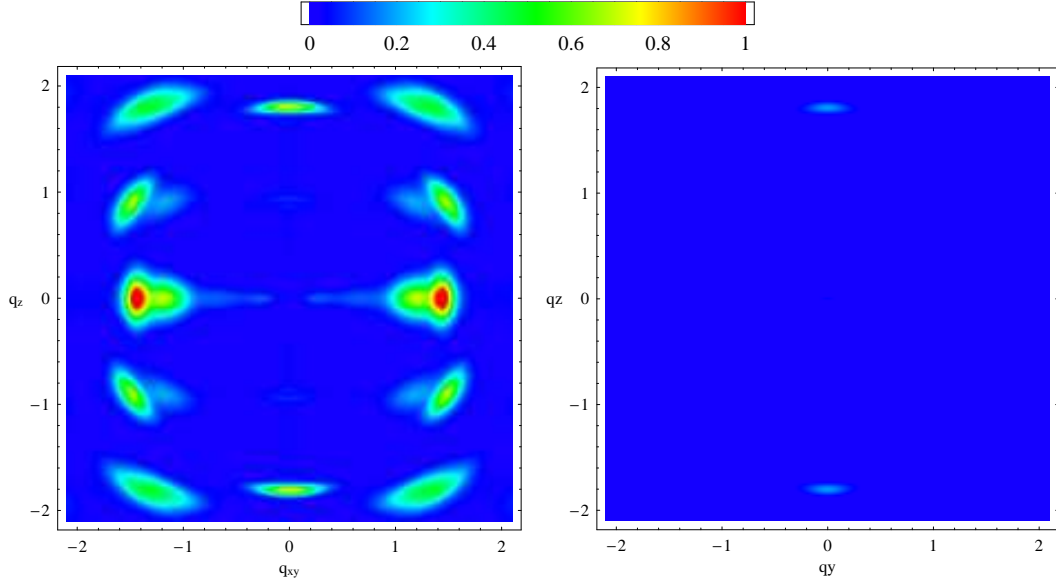


Figure E.2: Left: Calculated scattering images as in Fig. 4.12, but for a Gaussian distribution (rounded to integers) of the crystallite sizes M_x , M_y and M_z . The widths are $\Delta M_x = 2$, $\Delta M_y = 0.75$ and $\Delta M_z = 3$ respectively. Right: *correlated* part $G'_2(\mathbf{q})$ of the scattering function Eq. (4.20) in comparison to the *uncorrelated* part on the left.

The sums over the unit cells $\mathbf{m} = (m_x, m_y, m_z)$ are:

$$\begin{aligned}
 A(\mathbf{q}) &= \sum_{m_x=1}^{M_x} \sum_{m_y=1}^{M_y} \sum_{m_z=1}^{M_z} \sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q}(m_x\mathbf{a}_x + m_y\mathbf{a}_y + m_z\mathbf{a}_z - \mathbf{s}_{\text{cm}} + \mathbf{r}_k)) \\
 &= \sum_{m_x=1}^{M_x} \exp(i\mathbf{q}\mathbf{a}_x m_x) \sum_{m_y=1}^{M_y} \exp(i\mathbf{q}\mathbf{a}_y m_y) \sum_{m_z=1}^{M_z} \exp(i\mathbf{q}\mathbf{a}_z m_z) \\
 &\quad \times \exp\left(-i\mathbf{q} \cdot \frac{(M_x+1)\mathbf{a}_x + (M_y+1)\mathbf{a}_y + (M_z+1)\mathbf{a}_z}{2}\right) \sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q}\mathbf{r}_k). \quad (\text{E.4})
 \end{aligned}$$

For each $\nu = x, y, z$ the sum over m_ν is a geometric progression

$$\sum_{m_\nu=1}^{M_\nu} \exp(i\mathbf{q}\mathbf{a}_\nu m_\nu) = \frac{\exp(i\mathbf{q}\mathbf{a}_\nu(M_\nu+1)) - \exp(i\mathbf{q}\mathbf{a}_\nu)}{\exp(i\mathbf{q}\mathbf{a}_\nu) - 1}, \quad (\text{E.5})$$

and hence

$$\begin{aligned}
 &\sum_{m_\nu=1}^{M_\nu} \exp(i\mathbf{q}\mathbf{a}_\nu m_\nu) \exp\left(-i\mathbf{q} \cdot \frac{(M_\nu+1)\mathbf{a}_\nu}{2}\right) \\
 &= \frac{\exp(i\mathbf{q}\mathbf{a}_\nu(M_\nu+1)) - \exp(i\mathbf{q}\mathbf{a}_\nu)}{\exp(i\mathbf{q}\mathbf{a}_\nu) - 1} \frac{\exp(-i\mathbf{q}\mathbf{a}_\nu(M_\nu+2)/2)}{\exp(-i\mathbf{q}\mathbf{a}_\nu/2)}
 \end{aligned}$$

$$\begin{aligned}
&= \frac{\exp(i\mathbf{q}\mathbf{a}_\nu M_\nu/2) - \exp(-i\mathbf{q}\mathbf{a}_\nu M_\nu/2)}{\exp(i\mathbf{q}\mathbf{a}_\nu/2) - \exp(-i\mathbf{q}\mathbf{a}_\nu/2)} \\
&= \frac{\sin(i\mathbf{q}\mathbf{a}_\nu M_\nu/2)}{\sin(i\mathbf{q}\mathbf{a}_\nu/2)}. \tag{E.6}
\end{aligned}$$

With that, Eq. (E.4) becomes:

$$A(\mathbf{q}) = L_{M_x}(\mathbf{q}\mathbf{a}_x)L_{M_y}(\mathbf{q}\mathbf{a}_y)L_{M_z}(\mathbf{q}\mathbf{a}_z) \sum_{k=1}^K F_k(\mathbf{q}) \exp(i\mathbf{q}\mathbf{r}_k), \tag{E.7}$$

where $L_{M_\nu}(\mathbf{q}\mathbf{a}_\nu) = \frac{\sin(\mathbf{q}\mathbf{a}_\nu M_\nu/2)}{\sin(\mathbf{q}\mathbf{a}_\nu/2)}$ is the Laue function. The result (E.7) is used in Sec. 4.3.3.

Used Symbols and Notation

Contents

| | | |
|------------|---|------------|
| F.1 | Notation | 191 |
| F.2 | Symbols for the Randomly Cross-Linked Particle Model . . | 192 |
| F.3 | Symbols for the Cross-Linked Directed Polymers | 194 |
| F.4 | Symbols for the Spider Silk Model | 197 |
| F.5 | Symbols for the Wet Granular System | 199 |

F.1 Notation

A few matters concerning the notation are addressed here. The following remarks apply to the whole thesis, in the attempt to allow for a simple notation, but still dispel possible confusion:

For **volume integrals** with a D -dimensional variable \mathbf{x} , the simplified notation $\int_V d\mathbf{x} \dots$ or $\int d\mathbf{x} \dots$ is used, instead of $\int_V d^D \mathbf{x} \dots$ which can appear quite overloaded as soon as expressions become more complicated. Confusion with *line integrals*, like $\int \mathbf{F}(\mathbf{x}) d\mathbf{x} := \int \mathbf{F}(\mathbf{x}(t)) \cdot \mathbf{x}'(t) dt$, is not given, because they do not appear in this work. **Fourier and real functions** have the same name; e.g. $U(\mathbf{q})$ is the Fourier transform of the potential $U(\mathbf{r})$ in chapters 2 and 3. The type of the argument – real-space variable \mathbf{r} with units of length, or Fourier-space variable \mathbf{q} with units of inverse length – determines what function is meant. To describe the asymptotic behavior of a function, the **Landau symbol** \mathcal{O} is used. And the **imaginary unit** is always a roman “i” to avoid confusion with a variable “i”.

In the following, the used symbols and variables for the main chapters of this thesis can be found. For most symbols, also a reference to its introduction or definition is listed. “Temporary symbols”, however, which are only used near their definition, are not quoted in order to keep the list short and clear.

F.2 Symbols for the Randomly Cross-Linked Particle Model

| | |
|----------------------------------|---|
| $[\cdot]$ | disorder average, see Sec. 2.3 and Eq. (2.7) |
| $\int_{\xi_1^2, \dots, \xi_r^2}$ | $= \int_0^\infty d\xi_1^2 \mathcal{P}(\xi_1^2) \cdots d\xi_r^2 \mathcal{P}(\xi_r^2)$, introduced in Eq. (B.53) |
| \parallel, \perp | used for the separation of replicated $D(n+1)$ -dimensional vectors into “common” and “relative” components; see Eqs. (2.35)-(2.37) |
| 0RS | zero replica sector, $0\text{RS} = \{\hat{0}\}$, see Sec. 2.4.3 |
| 1RS | one replica sector, set of \hat{q} -values with exactly one replica component non-zero: $1\text{RS} = \{\hat{q} \neq \hat{0} \mid \hat{q} = (\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}^{(\alpha)}, \mathbf{0}, \dots, \mathbf{0})$ with $0 \leq \alpha \leq n\}$, see Sec. 2.4.3 |
| a | typical cross-link distance (mostly appearing as a^2) |
| A | arbitrary and unspecified quantity, used to define <i>general</i> expressions like a thermal average |
| α | specifying the replica and appearing as index $^{(\alpha)}$; $\alpha \in \{0, \dots, n\}$ |
| C_γ | $= (a^2 + \xi_\gamma^2)^{-1}$ defined in Eq. (B.54) |
| C | $= \sum_{\gamma=1}^r C_\gamma$ defined in Eq. (B.54) |
| \mathcal{C} | cross-link configuration, representing the M pairs of connected particles: $\mathcal{C} = \{i_e, j_e\}_{e=1}^M$ |
| D | spatial dimension |
| Dr | $= \text{d}\mathbf{r}/V$ dimensionless measure, introduced in the partition function Eq. (2.6) |
| $\text{D}\mathbf{x}_\parallel$ | $= \text{d}\mathbf{x}_\parallel/V_\parallel$ dimensionless measure for \parallel -vectors, residing in the volume V_\parallel |
| $\text{D}\hat{r}$ | $= \text{d}\hat{r}/V^{n+1}$ dimensionless measure for “hatted” vectors |
| $\mathcal{D}\Omega$ | measure for the integration over the fluctuating field Ω ; $\mathcal{D}\Omega \propto \prod_{\hat{q}} \text{d}\Omega(\hat{q})$, see Eq. (B.16) |
| $\Delta(\hat{q})$ | $= \exp(-\hat{q}^2 a^2/2)$, introduced in Eq. (2.20) |
| ε | distance from the critical cross-link concentration, $\varepsilon = \mu^2 - 1$ |
| f_0 | $= -\phi^n \frac{\mu^2}{2} + (n+1) \frac{\lambda n_0}{2}$, see Eq. (2.19) |
| $\tilde{f}_{n+1}\{\hat{r}_j\}$ | defined such that $\mathcal{Z}_{n+1} = \int (\prod_j \text{d}\hat{r}_j) \exp(-N\tilde{f}_{n+1}\{\hat{r}_j\})$, see Eq. (2.14) |
| $f_{n+1}\{\Omega\}$ | defined such that $\mathcal{Z}_{n+1} = e^{-Nf_0} \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\})$, see Eqs. (2.27) |
| $F_{\mathcal{C}}$ | $= -\ln Z_{\mathcal{C}}$, free energy (all energies in units of $k_B T$) |
| ϕ | $= (2\pi)^{D/2} \cdot a^D/V$ |
| G | shear modulus; see Eqs. (2.53) and Sec. 2.9.3 |
| $H_{\mathcal{C}}$ | $= H_{\text{Xlink}} + H_{\text{ev}}$ |
| H_{ev} | excluded volume Hamiltonian, $H_{\text{ev}}(\{\mathbf{r}_j\}) = \frac{\lambda}{2} \sum_{i,j=1}^N U(\mathbf{r}_i - \mathbf{r}_j)$ |
| $H_{\text{ev}}^{(n+1)}$ | $= \sum_{\alpha=0}^n H_{\text{ev}}(\{\mathbf{r}_j^{(\alpha)}\})$, replicated excluded-volume interaction |
| H_{Xlink} | cross-link Hamiltonian, $H_{\text{Xlink}} = \frac{1}{2a^2} \sum_{e=1}^M (\mathbf{r}_{i_e} - \mathbf{r}_{j_e})^2$, see Eq. (2.1) |
| HRS | higher replica sector, all \hat{q} -vectors with at least two non-zero entries: $\text{HRS} = \{\hat{q} \mid \hat{q} \neq \hat{0} \vee \hat{q} \notin 1\text{RS}\}$, see Sec. 2.4.3 |
| λ | strength of the excluded-volume interaction, see H_{ev} |
| $\tilde{\lambda}(\hat{q})$ | $= n_0 \lambda U(\hat{q}) - \phi^n \mu^2 \Delta(\hat{q})$, introduced in Eq. (2.24b). |

| | |
|---|---|
| M | number of cross-links |
| μ^2 | parameter of the Deam-Edwards distribution, see Eq. (2.7); found to be the average coordination number of the particles, see Sec. 2.9.1 |
| $\tilde{\mu}^2$ | $= \frac{V}{2Na^D} \frac{\mu^2}{(2\pi)^{D/2}} = \frac{\mu^2}{2N\phi}$ |
| N | number of particles |
| n | number of replicas |
| n_0 | $= N/V$, particle density |
| $O(\hat{x})$ | $= \frac{1}{N} \sum_{j=1}^N \delta(\hat{x} - \hat{r}_j)$, replicated real-space density; see Eq. (2.16) |
| $O(\hat{q})$ | $= \frac{1}{N} \sum_{j=1}^N \exp(i\hat{q}\hat{r}_j)$, replicated Fourier-space density; see Eq. (2.17a) |
| $\Omega(\hat{q})$ | order parameter of the theory, see e.g. Eq. (2.27b); in the thermodynamic limit equal to $O(\hat{q})$ |
| $\bar{\Omega}_{\text{sp}}$ | saddle-point solution for the order parameter; see Eqs. (2.33) and (2.48) |
| $\bar{\Omega}_u(\hat{q})$ | Ansatz for the order parameter with displacement field $u_{\perp}(\mathbf{x}_{\parallel})$, see Eq. (2.45) |
| $P(\mathcal{C})$ | probability for the cross-link configuration \mathcal{C} ; see Sec. 2.3 |
| $\mathcal{P}(\xi^2)$ | distribution of localization lengths; see Eq. (2.43) |
| π | $\pi/(2\pi) = 1/2$; simplifying notation |
| Q | gel fraction, i.e. the fraction of particles in the macroscopic cluster; introduction in Sec. 2.6 and discussion in Sec. 2.9.2 |
| $u_{\perp}(\mathbf{x}_{\parallel})$ | spatially dependent deformation field; see Sec. 2.6.2 |
| $\mathbf{u}_{\text{phys}}^{(\alpha)}(\mathbf{x}_{\parallel})$ | physically relevant deformation field (relative to the state during cross-linking); see Sec. 2.6.3 |
| $U(\mathbf{r})$ | potential for the excluded-volume interaction H_{ev} |
| $U(\mathbf{q})$ | Fourier transform of the potential $U(\mathbf{r})$ for the excluded-volume interaction H_{ev} |
| $U(\hat{q})$ | $= U(\mathbf{q}^{(\alpha)})$ for $\hat{q} = (\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}^{(\alpha)}, \mathbf{0}, \dots, \mathbf{0}) \in 1\text{RS}$ |
| \mathbf{r}_j | position of particle j |
| V | D -dimensional volume of the system |
| V_{\parallel} | $= V(n+1)^{D/2}$; rescaled volume of the system for \parallel -vectors, which are scaled by a factor of $\sqrt{n+1}$, compared to normal vectors; see Sec. 2.6.1 |
| ξ | localization length, quantifying the extent of the fluctuations of a localized particle; see Sec. 2.6 |
| Ξ_{r,a^2} | $= \int_{\xi_1^2, \dots, \xi_r^2} \ln \left(\left(\frac{V}{(2\pi)^{D/2}} \right)^{r-1} \left(\frac{\prod_{\gamma=1}^r (a^2 + \xi_{\gamma}^2)^{-1}}{\sum_{\gamma=1}^r (a^2 + \xi_{\gamma}^2)^{-1}} \right)^{D/2} \right)$ |
| $Z_{\mathcal{C}}$ | partition function, depending on the cross-link configuration \mathcal{C} |
| \mathcal{Z}_{n+1} | Made such that $[Z_{\mathcal{C}}^n] = \mathcal{Z}_{n+1}/\mathcal{Z}_1$, actual form given in Eq. (2.14c) |
| \mathfrak{z} | effective one-particle partition function, see Eq. (2.27c) |

F.3 Symbols for the Cross-Linked Directed Polymers

| | |
|----------------------------------|--|
| $[\cdot]$ | disorder average, see Sec. 3.3.2 and Eq. (3.22) |
| $\int_{\xi^2, z}$ | $= \int_0^\infty d\xi^2 \mathcal{P}(\xi^2, z)$, introduced in Eq. (3.45) |
| \parallel, \perp | used for the separation of replicated $D(n+1)$ -dimensional vectors into “common” and “relative” components; see Appendix A |
| $\sum'_{\hat{q} \in \text{HRS}}$ | $\sum'_{\hat{q} \in \text{HRS}} (\dots) := Q + \sum_{\hat{q} \in \text{HRS}} (\dots)$; simplifying definition, see Eq. (3.37) |
| 0RS | zero replica sector, 0RS = $\{\hat{0}\}$, see Sec. 3.4.2 |
| 1RS | one replica sector, set of \hat{q} -values with exactly one replica component non-zero: 1RS = $\{\hat{q} \neq \hat{0} \mid \hat{q} = (\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}^{(\alpha)}, \mathbf{0}, \dots, \mathbf{0})$ with $0 \leq \alpha \leq n\}$, see Sec. 3.4.2 |
| a | typical cross-linking length in the transverse plane; see H_{Xlink} and Eq. (3.15) |
| $a_{\hat{q}}^2$ | $= a^2/\ell^2$, defined in Eq. (3.59b) |
| A | D -dimensional (hyper-)area of the system in the transverse plane; see Sec. 3.3 |
| \mathcal{A} | arbitrary and unspecified quantity, used to define <i>general</i> expressions like a thermal average |
| α | specifying the replica and appearing as index (α) ; $\alpha \in \{0, \dots, n\}$ |
| \mathcal{C} | cross-link configuration, representing the M pairs of chains connected at height z_e ; $\mathcal{C} := \{(i_e, j_e, z_e)\}_{e=1}^M$; see Sec. 3.3 |
| $\mathcal{D}\mathbf{r}(z)$ | measure for the functional integral over all possible conformations of a chain $\mathbf{r}(z)$; see Eq. (3.21) |
| $\Delta(\hat{q})$ | $= \exp(-\hat{q}^2 a^2/2)$, introduced in Eq. (3.31c) |
| ε | distance from the critical cross-link concentration, $\varepsilon = \mu^2 - 1$ |
| f | $\frac{2}{3} + \frac{a^2}{\ell^2}$, see Eq. (3.60b) |
| f_0 | $= -\phi^n \frac{\mu^2}{2} + (n+1) \frac{\lambda n_0 L}{2}$; see Eq. (3.31b) |
| $\tilde{f}_{n+1}\{\hat{r}_j\}$ | defined such that $\mathcal{Z}_{n+1} = \int \left(\prod_{j=1}^N \mathcal{D}\hat{r}_j(z) \right) \exp(-N\tilde{f}_{n+1}\{\hat{r}_j\})$; see Eq. (3.27) |
| $f_{n+1}\{\Omega\}$ | defined such that $\mathcal{Z}_{n+1} = e^{-Nf_0} \int \mathcal{D}\Omega \exp(-Nf_{n+1}\{\Omega\})$, see Eqs. (3.32) |
| ϕ | $= (2\pi)^{D/2} \cdot a^D/A$ |
| H_{align} | $= \frac{\sigma}{2} \sum_{j=1}^N \int_0^L dz \mathbf{r}_j^2(z)$; alignment interaction of the whole system; defined in Eq. (3.17); note that $H_{\text{align}} = \sum_{j=1}^N H_{\text{align},1}\{\mathbf{r}_j\}$; |
| $H_{\text{align},1}$ | $= \frac{\sigma}{2} \int_0^L dz \mathbf{r}^2(z)$; alignment interaction of a single chain $\mathbf{r}(z)$; see Sec. 3.2.1 |
| $H_{\text{align}}^{(n+1)}$ | replicated alignment interaction; $H_{\text{align}}^{(n+1)}\{\hat{r}_j\} = \sum_{\alpha=0}^n H_{\text{align}}\{\mathbf{r}_j^{(\alpha)}\}$ |
| H_{ev} | $= \frac{\lambda}{2} \sum_{i,j=1}^N \int_0^L dz U(\mathbf{r}_i(z) - \mathbf{r}_j(z))$; excluded volume interaction, defined in Eq. (3.16) |
| $H_{\text{ev}}^{(n+1)}$ | replicated excluded volume interaction; $H_{\text{ev}}^{(n+1)}\{\hat{r}_j\} = \sum_{\alpha=0}^n H_{\text{ev}}\{\mathbf{r}_j^{(\alpha)}\}$ |
| H_{Xlink} | $= \frac{1}{2a^2} \sum_{e=1}^M (\mathbf{r}_{i_e}(z_e) - \mathbf{r}_{j_e}(z_e))^2$, defined in Eq. (3.15) |
| HRS | higher replica sector, all \hat{q} -vectors with at least two non-zero entries: |

| | |
|------------------------------|---|
| | HRS = $\{\hat{q} \mid \hat{q} \neq \hat{0} \vee \hat{q} \notin \text{1RS}\}$, see Sec. 3.4.2 |
| $I(\hat{q}_0, z_0)$ | simplifying definition, actual form given in Eq. (3.36) |
| L | height of the system in the direction of preferred alignment; see Sec. 3.3 |
| ℓ^2 | $:= L/(2\sigma) \propto R_g^2$; simplifying definition, see Sec. 3.2.1 and Eq. (3.6) |
| λ | strength of the excluded-volume interaction, see H_{ev} and def. (3.16) |
| $\tilde{\lambda}(\hat{q})$ | $= \lambda n_0 U(\hat{q}) - \phi^n \frac{\mu^2}{L} \Delta(\hat{q})$, introduced in Eq. (3.31d) |
| M | number of cross-links |
| μ^2 | parameter of the Deam-Edwards distribution, see Eq. (3.22); found to be the average coordination number of the chains; see Appendix C.7 |
| $\tilde{\mu}^2$ | $= \frac{\mu^2}{2N\phi} = \frac{A}{2Na^D} \frac{\mu^2}{(2\pi)^{D/2}}$ |
| n | number of replicas |
| N | number of chains |
| $O(\hat{x}, z)$ | $= \frac{1}{N} \sum_{j=1}^N \delta(\hat{x} - \hat{r}_j(z))$, replicated real-space density; see Eq. (3.29a) |
| $O(\hat{q}, z)$ | $= \frac{1}{N} \sum_{j=1}^N \exp(i\hat{q}\hat{r}_j(z))$, replicated Fourier density; see Eq. (3.29b) |
| $\Omega(\hat{q}, z)$ | order parameter of the theory, see e.g. Eq. (3.32); in the thermodynamic limit equal to $O(\hat{q}, z)$ |
| $\tilde{\Omega}(\hat{q}, z)$ | Ansatz for the order parameter $\Omega(\hat{q}, z)$, shown to solve the saddle-point equation; actual form given in Eq. (3.44) |
| $P(\mathcal{C})$ | probability for the cross-link configuration \mathcal{C} ; see Sec. 3.3.2 |
| $\mathcal{P}(\xi^2, z)$ | distribution of localization lengths at a given height z ; see Eq. (3.42) |
| $\mathcal{P}(\xi^2, s)$ | distribution of localization lengths at normalized height $s = z/L$; see Eq. (3.53) |
| $\pi(\theta, s)$ | distribution for θ at normalized system height $s = z/L$; see Eq. (3.60c) |
| $\bar{\pi}(\theta)$ | distribution for θ , averaged over the system height; see Eq. (3.62) |
| $\delta\pi(\theta, s)$ | deviation of $\pi(\theta, s)$ from the mean value $\bar{\pi}(\theta)$, normalized with ε ; see Eq. (3.64) |
| $q_{0\ell\perp}^2$ | $= \ell^2 q_{0\perp}^2$, defined in Eq. (3.59c) |
| Q | gel fraction, i.e. the fraction of chains in the macroscopic cluster; introduction in Sec. 3.5 and discussion in Sec. 3.6.2 |
| R_g | transverse radius of gyration of an uncrosslinked chain; $R_g^2 = \frac{D}{6} \frac{L}{2\sigma}$, see Sec. 3.2.1 |
| θ | $= f/(\varepsilon\xi_\ell^2)$; normalized inverse and squared localization length, see Sec. 3.7.1 and Eq. (3.60a) |
| $U(\mathbf{r})$ | potential for the excluded-volume interaction H_{ev} |
| $U(\mathbf{q})$ | Fourier transform of the potential $U(\mathbf{r})$ for the excluded-volume interaction H_{ev} |
| $U(\hat{q})$ | $= U(\mathbf{q}^{(\alpha)})$ for $\hat{q} = (\mathbf{0}, \dots, \mathbf{0}, \mathbf{q}^{(\alpha)}, \mathbf{0}, \dots, \mathbf{0}) \in \text{1RS}$ |
| ξ | localization length, quantifying the extent of the fluctuations of a localized chain segment; see Sec. 3.5 |
| $\tilde{\xi}^2$ | $= \xi^2 + a^2$, defined in Eq. (3.49) |
| ξ_ℓ^2 | $= \xi^2/\ell^2$, defined in Eq. (3.59a) |
| $Z_{\mathcal{C}}$ | partition function, depending on the cross-link configuration \mathcal{C} , introduced in Eq. (3.20) |

\mathcal{Z}_{n+1} Made such that $[Z_C^n] = \mathcal{Z}_{n+1}/\mathcal{Z}_1$, actual form given in Eq. (3.27)
 \mathfrak{z} effective one-particle partition function, see Eq. (3.32c)

F.4 Symbols for the Spider Silk Model

| | |
|---|--|
| $\langle \rangle$ | average over crystallite positions $\mathbf{R}^{(j)}$ and orientations $\underline{\underline{D}}^{(j)}$; see Sec. 4.2.3 and Eq. (4.6) |
| $\sum_{\mathbf{m}=1}^{\mathbf{M}}$ | $= \sum_{m_x=1}^{M_x} \sum_{m_y=1}^{M_y} \sum_{m_z=1}^{M_z}$, simplifying notation |
| $\mathbf{a}_{x,y,z}$ | primitive vectors of the crystallite, see Sec. 4.2.2 |
| $a_{x,y,z}$ | magnitudes of the primitive vectors \mathbf{a}_x , \mathbf{a}_y and \mathbf{a}_z |
| $A(\mathbf{q})$ | scattering amplitude of a single crystallite; defined in Eq. (4.16), treated in Sec. 4.3.3 |
| C_α | for the alanine amino acid, the C-atom to which the CH_3 group is bound |
| Δy_{12} | amount of the shift of strands 1 and 2 (see Fig. 4.8) in y -direction |
| Δz_{12} | amount of the shift of strands 1 and 2 (see Fig. 4.8) in z -direction |
| Δz_{24} | amount of the shift of strands 2 and 4 (see Fig. 4.8) in z -direction |
| $\underline{\underline{D}}^{(j)}$ | rotation matrix specifying orientation of crystallite j ; comprised by the Euler angles $\phi^{(j)}$, $\theta^{(j)}$, $\psi^{(j)}$; see Sec. 4.2.3 and Fig. 4.5 |
| $\underline{\underline{D}}\underline{\underline{D}}$ | measure for the integration over the orientation matrix $\underline{\underline{D}}$, see Sec. 4.2.3 and Eq. (4.7) |
| f_k | form factor of atom $k \in \{1, \dots, K\}$, see Sec. 4.2.1 |
| \bar{f} | amplitude of the cavities $V(\mathbf{r})$, see Sec. 4.2.4 and Eq. (4.8) |
| $F_k(\mathbf{q})$ | $= f_k - \tilde{V}(\mathbf{q})$, effective form factor of atom k ; see Eq. (4.12) and Sec. 4.3 |
| $G(\mathbf{q})$ | scattering function, general definition in Eq. (4.1), applied to our model in Eq. (4.11) |
| $G_1(\mathbf{q})$ | incoherent contribution of the scattering function $G(\mathbf{q})$, accounting for scattering from a single crystallite; treated in Sec. 4.3.1 |
| $G_2(\mathbf{q})$ | coherent contribution of the scattering function $G(\mathbf{q})$, accounting for scattering from different crystallites; treated in Sec. 4.3.2 |
| K | number of atoms in one unit cell |
| \mathbf{m} | $= (m_x, m_y, m_z)$; vector index specifying a unit cell inside a crystallite, see Sec. 4.2.2 |
| $M_{x,y,z}$ | number of repetitions of the unit cell in x -, y - and z -direction, respectively, in order to build one crystallite; see Sec. 4.2.2 |
| N | number of crystallites |
| \mathcal{P}_{pos} | $\mathcal{P}_{\text{pos}}(\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)})$ is the (correlated) distribution function for the crystallite positions $\mathbf{R}^{(1)}, \dots, \mathbf{R}^{(N)}$; introduced in Sec. 4.2.3 |
| $\mathcal{P}_{\text{angle}}$ | $\mathcal{P}_{\text{angle}}(\phi, \theta, \psi)$ is the probability to find a crystallite orientation specified by the Euler angles ϕ, θ, ψ ; see Sec. 4.2.3 and Fig. 4.5 |
| $\left. \begin{array}{l} \phi^{(j)} \\ \psi^{(j)} \end{array} \right\}$ | see $\underline{\underline{D}}^{(j)}$ |
| q_{xy} | component of the wave vector \mathbf{q} in x - or y -direction; the two directions are indistinguishable due to rotational symmetry of the system about the z -axis |
| $\mathbf{R}^{(j)}$ | position of crystallite j |

| | |
|-----------------------------------|---|
| \mathbf{r}_k | position of atom $k \in \{1, \dots, K\}$ relative to the center of the unit cell |
| $\mathbf{r}_{\mathbf{m},k}$ | position of atom k of unit cell \mathbf{m} , relative to the center of the crystallite; see Eq. (4.4) and Sec. 4.2.2 |
| $\mathbf{r}_{\mathbf{m},k}^{(j)}$ | position of atom k of unit cell \mathbf{m} of crystallite j ; see Eq. (4.5) and Sec. 4.2.3 |
| ϱ_0 | scattering strength of the continuous background between the crystallites, see Sec. 4.2.4 |
| $\mathbf{s}_{\mathbf{m}}$ | vector pointing from the center of an unrotated crystallite to the center of unit cell $\mathbf{m} = (m_x, m_y, m_z)$, see Sec. 4.2.2 |
| $\theta^{(j)}$ | Euler angle specifying the tilt of crystallite j from the fiber axis, see $\underline{D}^{(j)}$ |
| θ_0 | width of the distribution for the tilting angle θ of the crystallites from the fiber axis; see Sec. 4.2.3 |
| $\langle u^2 \rangle$ | mean square displacement of the atoms in any direction, to account for the Debye-Waller factor; see Eq. (4.21) |
| V | volume of the system |
| $V(\mathbf{r})$ | shape of the cavity of the “continuous background” around the atoms that prevents background scattering inside the crystallites; see Sec. 4.2.4 and Eq. (4.8) |
| $\tilde{V}(\mathbf{q})$ | Fourier transform of the cavity $V(\mathbf{r})$ |
| ξ | width of the cavities $V(\mathbf{r})$ |

F.5 Symbols for the Wet Granular System

| | |
|-------------------|--|
| β_2 | kurtosis, quantifying deviations from a Gaussian velocity profile; see Eqs. (5.25)-(5.26) |
| d | particle diameter |
| d_c | bond breaking distance, $d_c = 1.07d$ unless noted otherwise |
| D | spatial dimension |
| D_f | fractal dimension of a cluster |
| ΔE | bond breaking energy, <i>i.e.</i> energy needed for one bond rupture |
| f_{coll} | collision frequency |
| $g(\mathbf{r})$ | pair correlation function, see Sec. 5.4.2.3 and Eq. (5.45) |
| $g(r)$ | radial distribution function, spherical symmetric part of $g(\mathbf{r})$ |
| γ | proportionality factor between collision frequency and temperature in the clustered state, defined in Eq. (5.14) |
| L | linear dimension of the system |
| L_{box} | edge length of the boxes for the box counting algorithm, see Sec. 5.4.2.2 |
| L_{co} | cross-over length from fractal to compact behavior of the asymptotic cluster, see Sec. 5.4.2 |
| m | mass of a particle or a cluster of particles |
| $\bar{m}(t)$ | mean cluster mass, defined in Eq. (5.36) |
| n | $= N/V$, particle density |
| n_{cl} | number of clusters, see Sec. 5.4.1.3 |
| N | number of particles |
| N_{box} | number of boxes of edge length L_{box} needed to cover the cluster, see Sec. 5.4.2.2 |
| $N_m(t)$ | cluster mass distribution, <i>i.e.</i> number of clusters with m particles at time t ; see Sec. 5.4.1.2 |
| \mathcal{N}_i | the i -th of a total of n_{cl} clusters, used in Sec. 5.3.4 |
| $\mathcal{N}(i)$ | partial cluster of the asymptotic cluster, containing all particles that can be reached with i or less neighbor-to-neighbor steps, starting from a random particle, see Sec. 5.4.2.5 |
| ν | $= \frac{2}{3\sqrt{\pi}} \cdot \gamma \Delta E^{1/2}$, see discussion around Eq. (5.15) |
| P_{bb} | probability for a bond rupture, <i>i.e.</i> probability that the kin. energy is large enough to break a bond |
| ϕ | $= \frac{N}{V} \cdot \frac{\pi}{6} d^3$, volume fraction |
| \mathbf{r}_j | position of particle j |
| r_g | radius of gyration of a cluster, see Eq. (5.33) |
| σ | scattering cross section for a pair of particles |
| t | time |
| \tilde{t} | $= \nu t + c$, simplifying definition; see discussion around Eq. (5.22) |
| t_0 | transition time from free cooling to aggregation, see Eqs. (5.7) |
| T | granular temperature, see Sec. 5.3 and Eq. (5.2) |
| T_0 | initial value of the temperature; for all simulations $T_0 = 45\Delta E$ |

| | |
|-----------------|--|
| \mathbf{v}_j | velocity of particle j |
| V | $= L^3$; volume of the system |
| $w(\mathbf{v})$ | velocity distribution, see Sec. 5.3.3 |
| z | coordination number, <i>i.e.</i> number of neighbors of a particle |
| \bar{z} | average coordination number |

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Index

- alanine, 58, 69, 70, 76
- alignment interaction, 37
- amorphous matrix, 58
- amphiphilic block copolymers, 32
- angular distribution function
 - for crystallites, 64
- angular momentum, 99
- Ansatz for the order parameter, 18–20
- antiparallel beta-sheet, 70, 75
- asymptotic cluster
 - coordination number, 118–120
 - fractal dimension
 - box counting method, 110–112
 - correlation dimension, 116
 - from radius of gyration, 108
 - pair correlation function, 114–117
- asymptotic limit of cooling, *see*
 - free cooling, late stage
- average coordination number, 25
 - for the RLP model, 7
- ballistic aggregation models, 106
- beta-sheets, 58, 60, 70
 - classification of Warwicker, 59
- Bombyx mori, 76
- bond-breaking
 - distance d_c , 82, 84, 86, 118
 - energy ΔE , 82–84
 - probability P_{bb} , 86, 90, 97
- boundary effects
 - for cross-linked directed chains, 52–54
- bridge formation, 83
- capillary bridge, *see* liquid bridge
- chain density, *see* mean chain density
- classical theory of rubber elasticity, 28
- cluster (of granular particles)
 - definition, 98
 - number of \sim s, 106
 - partial \sim , 108, 122
- cluster mass distribution $N_m(t)$, 102, 105
 - scaling, 105–106
- clustering instability, 95
- coefficient of restitution, 82, 84
 - effective \sim for wet granulates, 84
- coherent scattering, 67–68, 187
- collision frequency, 86, 87, 91, 107
- continuous background, 64–65, 187
- continuous symmetry, 14, 15
- cooling, *see* free cooling
- coordination number, *see also* cross-link density
 - for the RLP model, 7, 25
 - for wet granulates, 118–120
- correlation dimension, 116
- correlations between crystallites, 67–68, 187
- cross-link
 - configuration, 4, 6, 36
 - density, 24, 25, 150, 180
 - fluctuations, 25
 - for directed polymers, 36–37
 - Hamiltonian, 5, 37
 - in spider silk, 58, *see* crystallites
- cross-linking, 7, 8
- crystallites, 58, 60
 - addition of glycine, 76
 - ensemble, 63–64
 - modeling, 60–61
 - tilting of \sim , 63
 - variations between \sim , 72, 76
- Deam-Edwards distribution, 6–8, 39
- deformation field u_{\perp}
 - expansion in, 140
 - introduction, 19

- delocalized particles, 16, 18, 26, 45
- density fluctuations, 12
- diffusion limited cluster-cluster aggregation, 103
- dimensionless inverse and squared localization length, 50
- dimensionless inverse temperature, 91
- dimensionless system height, 48
- disorder
 - in beta-sheets, 76
 - in replica calculations, *see* cross-link configuration
- disorder average, 6–7, 39–41
 - ~ed free energy, 7, 23–24, 40
 - ~ed replicated density, 10
- distribution function
 - for crystallite orientations, 64
 - for crystallite positions, 63
- distribution of localization length, 19
- dragline fibers, 57

- ensemble of crystallites, 63–64
- Euler angles, 63
 - illustration, 64
- excluded-volume interaction
 - for directed polymers, 37, 43
 - for the RLP model, 5
 - infinitely strong limit, 13, 44

- fiber axis, 63
- field theory
 - introduction, 12–13
- flocculation regime, 102
- flux lines, 32
- form factor, 60
 - effective \sim , 66
- formation of a liquid bridge, 83
- fractal dimension D_f
 - box counting method, 110
 - time evolution, 120
 - correlation dimension, 116
 - from radius of gyration, 104, 108
- free cooling
 - early stage, 87–90
 - late stage, 90–93
- free energy, 7, 40
- gel fraction, 16, 18, 25–27, 45
- gel phase, 4, 9, 14, 17
- gelation transition, 1, 4, 14, 26, 42, 49, 55
- glycine (substitution inside a crystallite), 76
- Goldstone fluctuations, 14–15
 - Ansatz with, 18–20
- Goldstone modes, 15
- granular temperature, 86

- Hausdorff dimension, 110
- high energy tails, 95
- higher replica sector, 11, 42, 136, 151, 164
- Hubbard-Stratonovich transformation, 12, 43, 130, 164

- incoherent scattering, 66–67
- incompressible deformations, *see* shear deformations
- inertia tensor, *see* moment of inertia tensor
- infinite cluster, *see* percolating cluster
- internal temperature, 99–101

- kurtosis, 97

- Laplace transformation, 175
- leptokurtic distribution, 97
- linearization, *see* Hubbard-Stratonovich transformation
- liquid bridge, 82
 - formation, 83
- liquid film
 - on granular particles, 82
- liquid phase, *see* sol phase
- localization length, 10, 16, 45
 - distribution of, 16, 19, 46
 - illustration, 10, 18
 - normalized distribution of, 50
 - final result, 52

- normalized inverse and squared, 50
 - typical, 54–55
- localized particles, 9, 26, 42, 45
- logarithmic time decay, 92
- macroscopic cluster, *see* percolating cluster
- mean chain density, 42, 164
- mean cluster mass, 105, 106
 - scaling, 107
- mean particle density
 - for the RLP model, 11, 130
 - for the wet granular system, 87
- moment of inertia tensor, 99, 122
- neighborhood definition, 98
- nematic field
 - polymers in a \sim , 38–39
- nematic polymers, 31
- normalization of length scales, 49
- normalized inverse and squared localization length, 50
- number of clusters, 106
- oblate object, 122
- one replica sector, 11, 42, 154
- one-particle partition function \mathfrak{z}
 - of cross-linked directed chains
 - introduction, 43, 167
 - with additional interactions, 160
 - of the RLP model
 - expansion, 137
 - introduction, 13, 132
- order parameter Ω
 - for directed polymers
 - Ansatz, 45–47
 - for the RLP model
 - Ansatz, 15–20
 - in the IRS, 154
 - insertion into $f_{n+1}\{\Omega\}$, 145
 - insertion into \mathfrak{z} , 137
 - meaning, 132
- pair correlation function, 114
 - at contact, 87
- parallel beta-sheet, 70, 75
- partial cluster, 108, 122
- particle density, *see* mean particle density
- partition function Z_C
 - of cross-linked directed polymers, 39
 - of the RLP model, 6
- percolating cluster, 4, 9, 16, 42, 101, 102
- percolation theory, 1, 102
- percolation time, 102
- periodic boundary conditions, 86
- permanent cross-links, *see* cross-link
- physical deformation field, 21
- platykurtic distribution, 97
- poly-L-alanine, 58, 76, 79
- polyalanine, *see*
 - alanine *or* poly-L-alanine
- polymer brushes, 31
- polymers in a nematic field, 38–39
- primitive vectors, 60
- principal axes, 122
- principal moments of inertia, 122
- principal vectors, 70
- prolate object, 122
- pure shear deformations, *see* shear deformations
- quenched disorder, *see* cross-link configuration
- radial distribution function, 114
 - Percus-Yevick solution, 114, 115
- radius of gyration, 103, 108
 - of a single polymer chain, 34, 48
- randomly cross-linked particle model, 3
- replica free energy $f_{n+1}\{\Omega\}$
 - of cross-linked directed chains
 - introduction, 43, 167
 - saddle point equation, 44–45
 - of the RLP model
 - introduction, 13, 132
 - result, 22, 149
 - saddle point value, 22
- replica free energy $\tilde{f}_{n+1}\{\hat{r}_j\}$, 8, 41

- replica partition function \mathcal{Z}_{n+1}
 - for cross-linked directed chains, 41
 - for the RLP model, 8
- replica trick, 7, 40
- replicated density field
 - introduction
 - for directed polymers, 41
 - for the RLP model, 9–11
- replicated vectors \hat{r}
 - introduction, 8
 - longitudinal component \mathbf{r}_{\parallel} , 17
 - properties, 17, 125–126
 - transverse component r_{\perp} , 17
- restitution coefficient, 84
- rotational temperature, 99–101

- saddle point equation, 44–45
- saddle point solution
 - family of \sim s, 17–18
 - fluctuations around the \sim , 135
 - for the RLP model, 16–18
- scattering amplitude
 - of a single crystallite $A(\mathbf{q})$, 67–69
 - peak shift, 69
- scattering cross section, 87, 89
- scattering function $G(\mathbf{q})$, 60, 65
 - effect of the parameters, 74
 - experimental, 73
 - from the model, 75–78, *see also*
 - Eqs. (4.11), (4.20)
 - peak shift, 69
 - side maxima, 76, 78
- shear deformations, 18–20
 - physically relevant \sim , 20
- shear modulus G , 22, 24, 27–29, 152
- single fiber diffraction, 58, 73–74
- single polymer chain
 - clamped in space, 34–36
 - radius of gyration, 33–34
- sol phase, 4, 9, 14
- sol-gel transition, *see* gelation transition
- spontaneous symmetry breaking, 15, 17–18
- statistical parameters of the spider silk
 - model, 74–75
- structural parameters of the spider silk
 - model, 74–75
- system spanning cluster, *see* percolating cluster

- tenacity, 58
- tensile strength, 57
- thick film model, 85
- thin film model, 85
 - comparison to thick film model, 88–90, 119–120
- time scale
 - for free cooling t_0 , 87
- translational invariance, 17
- translational temperature, 99–101
- tussah silk, 76
- type-II superconductors, 32

- unit cell
 - atomic configuration, 69–72
 - shifts, 72
 - unshifted, 69
 - variations between \sim s, 72, 76
- universality
 - of the RLP model, 13
- velocity distribution $w(\mathbf{v})$, 90, 95–98
- volume fraction, 86
- WAXS (wide angle X-ray scattering), 73
- worm-like chain, 38

- XY-model, 15

- zero replica sector, 12, 164

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