# Development of a Rhesus macaque engineered heart muscle model from pluripotent stem cells

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# **Table of Contents**

Acknowledgements	I
Abstract	III
List of Figures	IV
List of Tables	V
Abbreviations	VI
1. Introduction	1
1.1 The human heart	2
1.2 Heart failure	3
1.3 Heart failure treatments	4
1.4 Stem cells	6
1.4.1 Embryonic stem cells	6
1.4.2 Induced pluripotent stem cell	6
1.4.3 Parthenogenetic stem cells	7
1.4.3.1 Parthenote zygosity	8
1.5 Directed cardiomyocyte differentiation	11
1.6 Alginate encapsulation	12
1.7 Engineering of heart muscle	13
1.8 Animal models	14
1.9 Dissertation Objectives	17
2. Materials and Methods	18
2.1 Animal experiments	18
2.1.1 Superovulation of Rhesus Macaque (Macaca mulatta)	18
2.1.2 Collection of oocytes via follicle aspiration	19
2.1.3 Collection of oocytes via ovariectomy	21
2.1.4 Collection of oocytes via necropsy	21
2.1.5 Teratoma assay	21
2.2 Cell models	22
2.2.1 Human foreskin fibroblasts	22
2.2.2 Rhesus fibroblasts	22
2.2.3 Fibroblast cell culture	23
2.2.4 Preparation of fibroblast feeder layers	23
2.2.5 Preparation of Rhesus fibroblasts for engineered heart muscle	24

	2.2.6 In vitro maturation of Rhesus oocytes	. 25
	2.3 Stem cells	. 25
	2.3.1 Rhesus macaque parthenogenetic stem cells	. 25
	2.3.2 Rhesus Induced Pluripotent Stem Cell line C1	. 28
	2.3.3 Rhesus Induced Pluripotent Stem Cell line 43110-4	. 28
	2.3.4 Stem cell culture	. 28
	2.3.5 Rhesus Mamu sequencing	. 29
	2.4 Cardiomyocyte differentiation	. 34
	2.4.1 Directed differentiation	. 34
	2.4.2 Alginate encapsulation	. 35
	2.5 EHM Generation	. 35
	2.5.1 Construction of EHM casting molds and dynamic stretchers	. 35
	2.5.2 Casting of EHM	. 37
	2.5.3 Culturing of EHM	. 38
	2.5.4 Dissociation of EHM	. 39
	2.6 Measurements and imaging	. 39
	2.6.1 Cell count and size measurements	. 39
	2.6.2 Isometric contraction measurements	. 39
	2.6.3 Flow cytometry	. 42
	2.6.4 Immunofluorescence imaging	. 44
	2.6.5 Whole mount imaging	. 44
	2.7 Statistical analysis	. 45
3	. Results	. 46
	3.1 Encapsulation of Lenti-iPS cells in an alginate:gelatin hydrogel	. 46
	3.2 Directed differentiation of alginate encapsulated Rhesus Lenti-iPS cells in	1
	suspension culture	. 48
	3.3 Directed differentiation of Rhesus Sendai iPS cells in monolayer culture	. 50
	3.4 Mamu screening	. 52
	3.5 Teratoma assay	. 54
	3.6 EHM Isometric force measurement	. 55
	3.7 EHM cardiomyocyte content	. 57
	3.8 EHM post transplantation gene expression	. 59
	3.9 Superovulation and oocyte retrieval of Rhesus macaque oocytes	. 61
	3.10 Activation of retrieved oocytes	. 63

# **Table of Contents**

4. Discussion	69
4.1 RhiPS cells can differentiate into spontaneously beating cardiomyocytes	
4.2 EHM can be constructed from RhiPS cell-derived cardiomyocytes	72
4.3 Rhesus macaque EHM appear less mature than human EHM	73
4.4 iPS reprogramming by integrating vectors increases the likelihood of oncogene	
expression	74
4.5 Encapsulation enhanced directed differentiation in Lenti-iPS cells	75
4.6 RhpSC can be derived from unfertilized oocytes after superovulation	77
5.0 Summary and outlook	80
Bibliography	81
Appendix1	10

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This work is dedicated to my father who passed away during my PhD candidature.
This work is dedicated to my father who passed away during my PhD candidature.  "Problems are not stop signs; they are guidelines." – Robert Schuller

## **Abstract**

Myocardial infarction is the leading cause of death globally. Remuscularization of the heart with stem cell based treatments is one potential approach to heart repair which is presently being tested. Proof-of-concept studies, mostly in rodent models, have shown promising results. Nevertheless, to be therapeutically relevant it is essential to test evolving therapeutic strategies in large animal models with high predictive value as to clinical outcome. Unfortunately, a lack of pluripotent stem cell models available in most animal species used for late stage preclinical testing creates a major challenge. Non-human primate models, and here in particular macaque models, appear to be an exception with the documentation of successful derivation of stable pluripotent stem cell models by several groups. In this dissertation the following hypotheses were tested: (1) induced pluripotent stem (iPS) cells from Rhesus macaque (Macaca mulatta) can be used to derive cardiomyocytes, (2) Rhesus macaque iPS-derived cardiomyocytes can be used to construct engineered heart muscle (EHM) with contractile properties similar to observations in human EHM, (3) parthenogenetic pluripotent stem cells can be derived from Rhesus macaque unfertilized oocytes as an alternative source of pluripotent stem cells with distinct advantages regarding immunological matching. Two available Rhesus iPS cell lines, either generated by lentiviral or Sendai viral transduction of reprogramming factors, were subjected to directed differentiation using a similar protocol as recently established for human pluripotent stem cells. High cardiomyocyte yield and purity could be achieved. Cardiomyocytes differentiated from both iPS cell lines could be used to generate EHM with similar contractile properties. Evidence for successful induction of parthenogenesis was obtained. Collectively, this dissertation provides the methodological groundwork for the implementation of Rhesus macaque pluripotent stem cell derived EHMs for pivotal late stage preclinical studies of tissue engineered heart failure repair.

# **List of Figures**

Figure 1: Heart failure caused by myocardial infarction	3
Figure 2: Reprogramming of somatic cells to iPS cells	7
Figure 3: Parthenogenic activation of unfertilized MII oocytes	9
Figure 4: Possible chromatid arrangements in an oocyte	10
Figure 5: Directed differentiation of cardiomyocytes for EHM generation	12
Figure 6: Polysaccharide residues of alginate	13
Figure 7: Evolutionary distance of mammalian test models	15
Figure 8: Reusable EHM mold and stretcher	36
Figure 9: Rhesus iPS EHM	38
Figure 10: EHM in organ bath	42
Figure 11: Gating strategy for α–actinin positive cardiomyocytes	43
Figure 12: Encapsulation of Lenti-iPS cells:	47
Figure 13: Increase in cell number following encapsulation	48
Figure 14: Cardiomyocyte purity following differentiation in alginate beads	49
Figure 15: Cardiomyocyte purity following differentiation of Sendai iPS cells in	
monolayer culture	51
Figure 16: Fluorescence image of RhiPS derived cardiomyocytes	52
Figure 17: Teratoma assays confirmed pluripotency in Sendai RhiPS cells	54
Figure 18: Calcium concentration response in the two tested iPS cell-derived EH	IM.
	55
Figure 19: Positive inotropic responses to isoprenaline in Rhesus EHM	56
Figure 20: Negative force-frequency response in Rhesus EHM	57
Figure 21: Cardiomyocyte content of Rhesus EHM	58
Figure 22: Whole mount staining of RhEHM	59
Figure 23: Ultrasound examination of hormone stimulated Rhesus ovaries	61
Figure 24: Excised ovaries from super-ovulated Rhesus macaques	62
Figure 25: Oocytes collected from excised ovaries	63
Figure 26: Cellular division of activated Rhesus oocytes	64
Figure 27: Rhesus ICM plating	65
Figure 28: Summary of all Rhesus macaque oocyte activations	68

# **List of Tables**

Table 1: Heart transplants in 2016 in the Eurotransplant zone5
Table 2: Comparative heart anatomy and physiological data between human and
common heart disease animal models16
Table 3: Female Rhesus macaques used for superovulation and oocyte harvest 18
Table 4: TALP-HEPES medium for follicle aspiration
Table 5: HECM-9 medium for oocyte activation and culture
Table 6: Amino acid/Pantothenate solution for oocyte activation and culture 27
Table 7A: Oligonucleotide sequences for Rhesus MHC class I PCR amplification 30
Table 7B: Amplification program for MHC Class I
Table 8A: Oligonucleotide sequences for Rhesus MHC class II PCR amplification . 34
Table 8B: Amplification program for MHC Class II
Table 10: Tyrode's solutions for isometric force measurements
Table 11: Rhesus macaque MHC Class I (Mamu) antigen screening 53
Table 12: Comparative RNA expression of pluripotency and cardiomyocyte marker
genes in Lenti-EHM explants and Sendai-cardiomyocytes 60
Table 13: Oocyte activations of Rhesus macaques from ovulation cycle 1 65
Table 14: Oocyte activations of Rhesus macaques from ovulation cycle 2 66
Table 15: Oocyte activations of Rhesus macaques from ovulation cycle 3 67
Table 16: Oocyte activation of Rhesus #237967

## **Abbreviations**

Asc Ascorbic acid

ANOVA Analysis of variance

BMP4 Bone morphogenetic factor 4

BSA Bovine serum albumin

C Celsius/Centigrade

CHIR Wnt signaling activator CHIR99021

CO<sub>2</sub> Carbon dioxide

CVD Cardiovascular disease
Cx43 Connexin 43 (*aka* GJA1)

DAPI 4', 6-diamidino-2-phenylindole

DMEM Dulbecco's modified Eagle's medium

DMSO Dimethyl sulfoxide

Dpn Days postnatal

EC<sub>50</sub> Half maximum effective concentration

ECM Extracellular matrix

EDTA Ethylenediaminetetraacetic acid

EHM Engineered heart muscle

ESC Embryonic stem cell

EtOH Ethanol

FBS Fetal bovine serum

FGF-2 Basic fibroblast growth factor

FITC-A Fluorescein isothiocyanate Area

FOC Force of contraction
FSC-A Forward scatter area

g Gravitational force

H<sub>2</sub>O Water

hCG Recombinant human chorionic gonadotropin

HECM-9 Hamster embryo culture medium 9

HEPES 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid

hES Human embryonic stem cell
hFF Human foreskin fibroblasts

hFSH Recombinant human follicle stimulating hormone

IgG Immunoglobulin G
IM Intramuscularly

IMDM Iscove's modified Dulbecco's medium

iPS Induced pluripotent stem cell

IU International units

IVF In vitro fertilization

IVM In vitro maturation

IWP4 Inhibitor of Wnt production 4

KO Knockout

L Liter

Lenti Lentiviral transduced RhIPS line C1

LIF Leukemia inhibitory factor

mESC Mouse embryonic stem cell

α-MHC Myosin heavy chain, alpha isoform (aka MYH6)
 β-MHC Myosin heavy chain, beta isoform (aka MYH7)

MII Meiosis II
ml Milliliter
mN Millinewton

MYC Transcriptional gene encoding for c-MYC

NaCl Sodium chloride
NaOH Sodium hydroxide

NEAA Non-essential amino acids

NHP Non-human primate

PBS Phosphate buffered saline

POU5F1 Transcriptional gene encoding for OCT4

PP Polypropylene

P/S Penicillin/Streptomycin

pSC Parthenogenetic stem cell

Rh Rhesus macaque (Macaca mulatta)

RhGFB Rhesus gingival fibroblasts

RhiPS Rhesus macaque induced pluripotent stem cell

RhpSC Rhesus parthenogenetic stem cell

RI ROCK inhibitor Y-27632

RPMI Roswell Park Memorial Institute

# Abbreviations

RT Room temperature

SEM Standard error of the mean SFBM Serum free basal medium

SFMM Serum free maturation medium

SSC-A Side scatter area

TALP Tyrode's albumin lactate pyruvate
TGF-β Transforming growth factor beta

U Unit V Volts

# 1. Introduction

Cardiovascular diseases (CVD) cause 31% of all deaths and thus are considered the number one cause of death globally (WHO 2016). The associated long term consequences of CVD place an increasing health and economic burden on society. As the global population continues to increase and live longer than any previous generation, the demands on the scientific and medical communities to create more specialized and personalized medicine is greater than ever before. Stem cells can be derived from individual patients and thus appear well suited as personalized therapeutics, including cell-based therapeutics for heart repair. Remuscularization of the heart is one potential approach to heart repair which is presently tested either by the implantation of pluripotent stem cell derived cardiomyocytes (Chong et al. 2014, Klug et al. 1996, Shiba et al. 2012, Shiba et al. 2016) or engineered heart muscle constructs (Zimmermann et al. 2006, Didie et al. 2013, Riegler et al. 2015, Weinberger et al. 2016). After proof-of-concept in mostly rodent models, it is absolutely essential to test the evolving therapeutic strategies in large animal models with high predictive value as to clinical outcome. Large animal models with widespread use in late preclinical studies include dogs, sheep, pig and non-human primate (NHP). However, the lack of pluripotent stem cell models in most of these species constitutes a major challenge. An exception is the NHP model. Several groups have demonstrated that robust pluripotent stem cell models can be derived from various NHPs, including macaque species - Macaca mulatta (Chan et al. 2010, Liu et al. 2008), Macaca fascicularis (Deleidi et al. 2011, Shiba et al. 2016), Macaca nemestrina (Kiem et al. 2004, Sourisseau et al. 2013) - as well as the common marmoset, Callithrix jacchus (Tomioka et al. 2010, Wu et al. 2010) - and the olive baboon - Papio anubis (Navara et al. 2013). These models utilize embryonic stem cells (ESC; Debowski et al. 2016, Thomson et al. 1995), induced pluripotent stem cells (iPSCs; Debowski et al. 2015, Liu et al. 2008, Zhong et al. 2011), and parthenogenetic stem cells (pSCs; Mitalipov et al. 2001, Vrana et al. 2003). Moreover, the close phylogenetic distance between macaque and human was used as a rationale for studies testing the electromechanical integration of human embryonic stem cell-derived cardiomyocytes in macaque models (Chong et al. 2014). Immunologically more relevant allograft studies have recently been performed in Rhesus macaque (Shiba et al. 2016) with similar results

as in the aforementioned xenograft study. The importance of NHP models is further exemplified by earlier studies in guinea pig with opposite results as to the side effects elicited by the cardiomyocyte grafts: namely, anti-arrhythmic effects of human cardiomyocyte grafts in the guinea pig model (Shiba et al. 2012) and pro-arrhythmic effects in the macaque model (Shiba et al. 2016).

Accurate and reproducible animal models of human diseases are invaluable in not only their utilization for pre-clinical treatment evaluations, but also in their ability to provide a deeper and better understanding of disease mechanisms. Rats and mice represent the most common preclinical animal models, mainly because of relatively low costs, easy breeding with short lifecycles (1-2 years), and high reproduction rates as well as the possibility for genetic manipulations in genetically defined backgrounds. However, differences in physiology, and in particular cardiovascular physiology, limit their predictive value for clinical outcome. Animals such as sheep and pigs are more closely related in physiology to humans and have been previously used in myocardial infarction and stem cell studies (Dayan et al. 2016, McCall et al. 2012). However, with regards to their use in stem cell studies, their limitation is in the high level of difficulty present in generating species-specific stem cells (Harding et al. 2013).

## 1.1 The human heart

The heart is a complex, multicellular, four chambered pump designed to propel blood through more than 60,000 miles of vasculature (Rajendran et al. 2013) in the adult human. Starting at day 20 of embryogenesis, the human heart begins to beat (Moorman et al. 2003). From this point on, the heart is continuously transporting blood throughout the body (Moorman et al. 2003), making it the longest continuously functioning organ in the body. In a normal human heart, cardiomyocytes comprise 75% of the total volume but only account for 30-40% of the total cell number (Vliegen et al. 1991). The non-myocyte fraction consists of fibroblasts, smooth muscle cells, endothelial cells (Camelliti et al. 2005), and transient immune response cells (lymphocytes, macrophages, and mast cells). These cells must work in precise harmony to maintain proper function and homeostasis of the heart. The specific morphological transition and intramyocardial arrangement of cardiomyocytes from an elongated, ovoidal embryonic phenotype to a rod-shaped adult phenotype enables the cardio-

myocytes to maintain electrically and mechanically networked connectivity and anisotropy ensuring the highly-synchronized contractility of the adult heart. This highly specialized nature of adult cardiomyocytes comes at the cost of a having very limited, if any, regenerative capacity (Laflamme and Murry 2011, Lundy et al. 2013).

#### 1.2 Heart failure

The chance of developing heart failure has been calculated at one in five over the course of a human lifetime (Bui et al. 2011). Myocardial infarction, affecting typically the left ventricle, is one of the main reasons for heart failure development. The left ventricle of young adults (17-30) consists of approximately  $6 \times 10^9 \pm 1.8 \times 10^9$  cardiomyocytes and decreases up to 33% to  $4 \times 10^9 \pm 1.3 \times 10^9$  between ages 65-90 (Olivetti et al. 1991). Approximately 1 billion cardiomyocytes are usually lost in the left ventricle of the human heart following a myocardial infarction (Gepstein 2002, Laflamme and Murry 2005). In response to this loss of myocardium, a remodeling cascade, mediated by fibroblasts and myofibroblasts, is initiated to prevent ventricular wall rupture (Shinde and Frangogiannis 2014, van den Borne et al. 2010). This remodeling cascade results in both functional and morphological changes to the myocardium in the form of fibrosis. Over time, the chronic stresses inherent from maladaptive remodeling lead to more severe CVD states including; ventricular dilation, cardiac fibrosis, cardiomyocyte hypertrophy, and chronic inflammation (Figure 1).

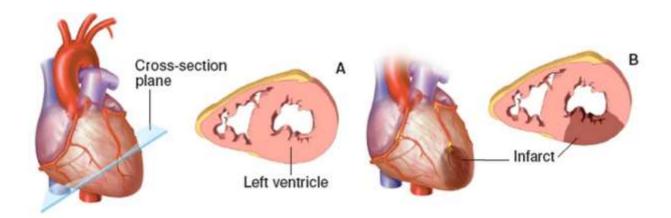


Figure 1: Heart failure caused by myocardial infarction

Myocardial infarction causes the death of approximately 1 billion cardiomyocytes, usually in the left ventricle of the heart. Due to the limited capacity of the heart to re-

place the cardiomyocytes, tissue scarring occurs and results in decreased left ventricular output. (image modified from Goldthwaite 2006)

#### 1.3 Heart failure treatments

Current treatment modalities following myocardial infarction consist of surgical repair procedures, pharmaceutical drug interventions, mechanical assistance device implantations, and organ transplantations. Despite advances in these treatments, patients suffering from heart failure still face substantial morbidity and mortality within the 5 years following a cardiac incident (Bui et al. 2011). Full organ transplantations are currently the only course of action for patients with chronically scarred myocardium and end-stage heart failure. In 2016, 702 people were on a waiting list for a heart only transplant, with nearly half already having waited 24 months or more (Table 1A). This relatively low number of patients on the heart transplant waiting list, in light of the huge number of patients affected by heart failure, is a result of very stringent selection criteria, e.g., patients with an age above 65 are typically not considered. Finally, the total numbers of patients with end stage heart failure in need of heart transplantation in Germany are estimated to be approximately 8,000. In 2016, only 297 heart transplants (287 heart only, 10 heart + other organ) were performed in Germany (Table 1B). In light of changing patient demographics, this number and thus the need for advanced heart failure therapeutics will further increase.

As an alternative to full organ replacement, stem cell based treatments have gained intense interest as an option for physical replacement of damaged tissues in patients with end stage heart failure (Agnetti et al. 2015, Nicolini et al. 2015). Stem cell based treatment options consist primarily of direct injection (reviewed in Terrovitis et al. 2010) or application of cell patches (reviewed in Ye et al. 2013). Both treatments are being investigated with either non-cardiac cells (Farahmand et al. 2008, Mazo et al. 2010) or stem cell derived cardiomyocytes (Chong et al. 2014, Riegler et al. 2015, Weinberger et al. 2016)

ths waiting (on WL)	Α	В	D	H	HR	NL.	SLO	Total
0-5	21	28	136	18	18	20	16	257
6-11	12	34	115	10	11	22	12	216
12-23	8	42	138	5	3	33	15	244
24+	10	9	313	15	2	24	7	380
Total	51	113	702	48	34	99	50	1097
Age	Α	В	D	Н	HR	NL	\$LO	Total
0-15	2		34		1	3		40
16-55	25	70	356	30	16	60	26	583
56-64	16	32	251	17	14	29	19	378
65+	8	11	61	_ 1	3	7	5	96
Total	51	113	702	48	34	99	50	1097
Urgency	Α	В	D	Н	HR	NL	SLO	Total
Elective	48	111	614	48	28	96	42	987
High Urgency	3	2	88		6	3	8	110
Total	51	113	702	48	34	99	50	1097
eurotransplant.org : 3013P_20	16_heart : 1	2.01.2017	excludes p	atients also w	raiting for oth			
Heart transplants								
ased donor transplants	А	В	D	н	IR N	L SLO	Non-ET	Tota
ased donor transplants heart		B 68	D 287				Non-ET	573
ased donor transplants heart heart + lungs	А	В	D 287 7	н	IR N	L SLO	Non-ET	57
ased donor transplants heart heart + lungs heart + liver	A 56	B 68	D 287 7 1	н	IR N	L SLO	Non-ET	57
ased donor transplants heart heart + lungs	А	B 68	D 287 7	н	IR N	L SLO	Non-ET 0 4	57

Table 1: Heart transplants in 2016 in the Eurotransplant zone

A) In 2016, over 700 people were actively waiting for a heart transplant in Germany, with almost 45% of patients waiting for a suitable donor for 24 months or more. B) Due to a lack of suitable heart transplants only 297 heart transplants were performed in Germany during 2016. (A= Austria, B = Belgium, D= Germany, H = Hungary, HR = Croatia, NL = Netherlands, SLO = Slovenia. Images modified from http://statistics.eurotransplant.org/, accessed 28 February 2017.)

#### 1.4 Stem cells

Stem cells are defined as cells that can self-renew and proliferate in an undifferentiated state indefinitely. These cells give rise to every cell type in the body. Stem cells are classified into four categories dependent on their differentiation potential: totipotent, pluripotent, multipotent, or unipotent. Blastomeres produced by the first zygotic divisions can differentiate into all cell types and thus are considered totipotent. Pluripotent stem cells can differentiate into almost all cell types and include blastocoel derived embryonic stem cells (ESC), induced pluripotent stem cells (iPS), and parthenogenetic stem cells (pSC). Multipotent and unipotent stem cells are of adult origin and give rise to cells within their related family – such as adult hematopoietic stem cells – or cells that possess the ability to self-renew, but are only capable of producing cells of their own type, such as muscle stem cells.

# 1.4.1 Embryonic stem cells

Derived from the inner cell mass of blastocyst stage preimplantation embryos, ESCs were first generated from mice by Martin Evans at Cardiff University (Evans and Kaufman 1981). Over the course of the following decades, ESCs were generated from other animals including hamsters (Doetschman et al. 1988), sheep (Notarianni et al. 1991), rabbits (Graves and Moreadith 1993), non-human primates (Thomson et al. 1995), humans (Thomson et al. 1998), canines (Hatoya et al. 2006), and rats (Li et al. 2008). ESC-like cell lines have also been generated from cattle (Saito et al. 1992) and pigs (Hou et al. 2016). ESCs possess the ability to proliferate in an undifferentiated state indefinitely, giving rise to every cell type in the body. Of note, although a vast amount of ESCs from various species has been reported, there is a huge discrepancy as to the actual acceptance as bona fide pluripotent stem cells in the field. Especially, stable long term maintenance of a pluripotent state and susceptibility to directed differentiation has been limited to mouse, macaque and human ESCs. The discovery of human ESCs has however brought with it legal and ethical issues primarily associated with the destruction of the blastocyst stage embryos used to isolate the inner cell mass, from which ESCs are derived (Hyun 2010).

#### 1.4.2 Induced pluripotent stem cell

In 2006, it was reported that mouse fibroblasts could be reprogrammed back to a pluripotent state using four reprogramming factors delivered via retroviruses

(Takahashi and Yamanaka 2006). A year later, human iPS cells were generated making it possible to transform a patient's own somatic cells back into stem cells and then differentiate those cells into practically any cell required for a therapeutic treatment (Takahashi et al. 2007). Much like their predecessors before them, iPS cells have also been generated from a variety of species, including Rhesus macaque (Liu et al. 2008), rats (Liao et al. 2009), rabbits (Honda et al. 2010), and canines (Koh and Piedrahita 2015). With iPS cells, it is possible to generate stem cell lines from multiple somatic cells sources without the destruction of blastocyst stage embryos (Egusa et al. 2010).

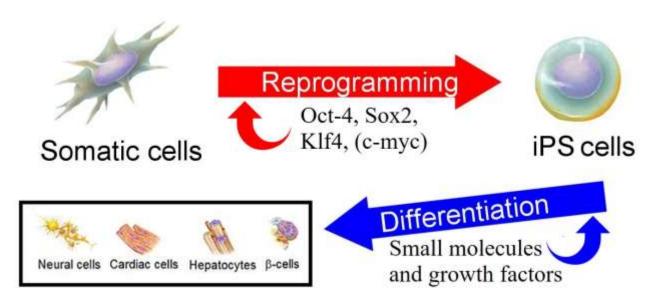


Figure 2: Reprogramming of somatic cells to iPS cells

Generation of iPS cells from somatic cells requires the viral transduction of four transient pluripotency genes: Oct-4, Sox2, Klf4, and c-Myc. Once transduced, iPS cells can be terminally differentiated into any cell in the body.

#### 1.4.3 Parthenogenetic stem cells

Similar to ESC, parthenogenetic stem cells (pSC) are generated by isolation of the inner cell mass of a blastocyst (Revazova et al. 2007). The difference is, however, that these are non-embryonic blastocysts and thus the term "parthenogenetic embryonic stem cells" used by some authors is a clear misnomer. To date, parthenote blastocysts have been generated in human (de Fried et al. 2008, Revazova et al. 2007), non-human primates (Mitalipov et al. 2001, Vrana et al. 2003), rabbit (Naturil-

Alfonso et al. 2015), goat (Kharche et al. 2014), mouse (Cui et al. 2007, Didie et al. 2013), and pig (Margues et al. 2011). Stable pSC lines have so far only been generated from human (Harness et al. 2011, Revazova et al. 2008, Turovets et al. 2011), non-human primate (Dighe et al. 2008, Mitalipov et al. 2001, Vrana et al. 2003, Yang et al. 2013), and mouse (Chen et al. 2009, Didie et al. 2013). To generate an inner cell mass for the derivation of pSCs, metaphase II (MII) oocytes are activated through exposure to exogenous calcium (Ca<sup>2+</sup>), causing an increase of intracellular Ca<sup>2+</sup>. The exposure of an unfertilized oocyte to Ca<sup>2+</sup> mimics the release of phospholipase C-zeta (PLC- $\zeta$ ) – otherwise known as the "sperm factor" – from the sperm head during fertilization. This binds to the inositol 1,4,5-trisphospahte receptors (IP3R) of the endoplasmic reticulum and initiates Ca<sup>2+</sup> oscillations in a positive Ca<sup>2+</sup> feedback loop, resulting in the resumption of meiosis II (Saunders et al. 2002, Swann and Yu 2008). PLC-ζ has been found to be associated with the induction of Ca<sup>2+</sup> oscillations of fertilized eggs in chickens (Coward et al. 2005), bovine (Ross et al. 2008), as well as human and non-human primates (Cox et al. 2002, Rogers et al. 2004). Since there is no paternal DNA present in the activated oocyte, the chromosomal zygosity of the resulting pSC is dependent upon the activation protocol used (Revazova et al. 2008).

# 1.4.3.1 Parthenote zygosity

pSCs can be derived from activated meiosis II oocytes either before or after extrusion of the 2<sup>nd</sup> polar body, resulting in partly heterozygous or strictly homozygous genomes, respectively (reviewed in Daughtry and Mitalipov 2014).

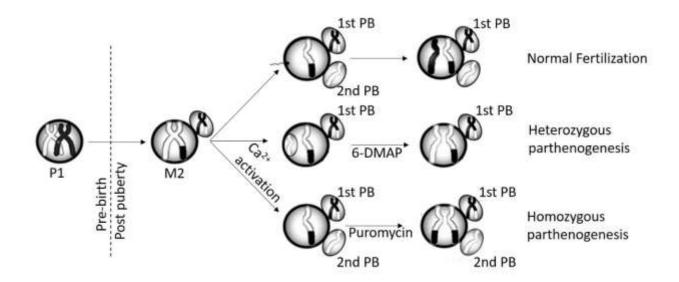


Figure 3: Parthenogenic activation of unfertilized MII oocytes

During normal fertilization, resumption of meiosis II occurs when a sperm penetrates the MII oocyte, triggering a calcium release. Immediately following this reactivation, the sister chromatids are segregated. One chromatid is packaged into the second polar body and is extruded. The other chromatid is supplemented with the paternal chromatid supplied by the sperm which results in the formation of a diploid zygote. Parthenogenetic activation of an arrested MII oocyte is achieved through exposure to exogenous calcium. Immediately following activation, incubation with 6-DMAP causes the retention of the second polar body within the oocyte. Incubation with puromycin does not inhibit second polar body extrusion, resulting in a transient haploid state. DNA-synthesis typically sets in to support the formation of a homozygous diploid state (Image modified from Daughtry and Mitalipov 2014).

Heterozygosity results from crossing over events during meiosis I with exchange of genome information from one of the two alleles within a pair of matching parental chromosomes (Figure 4). In metaphase II oocytes, derived after blockade of the extrusion of the 2<sup>nd</sup> polar body, diploid pSCs exhibit a large degree of heterozygosity in genes encoded distant from the centromere; conversely, genomic sequences close the centromere are typically homozygous (Didie et al. 2013, Kim et al. 2007b).

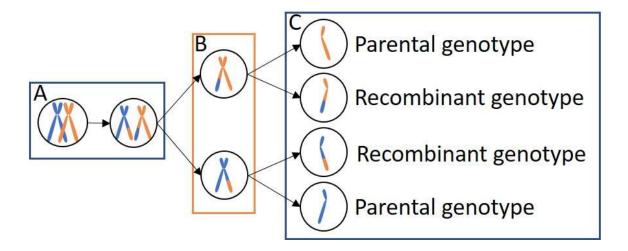


Figure 4: Possible chromatid arrangements in an oocyte

Chromosomes are comprised of two sister chromatids bound together by a centromere. A) Prior to birth, during prophase (PI) of meiosis I in oogenesis, chromosomal crossing occurs when matching regions of two sister chromatids from separate homologous chromosomes break apart and recombine on opposite chromatid strands that are near each other. This chromosomal crossing over results in recombinant chromosomes that are genetically diverse from the parental chromosomes. Following this crossing over event, oogenesis is halted. B) When puberty is reached meiosis I is resumed and the first polar body is formed and extruded. The first polar body contains one full half of the original oocyte's diploid chromosome set. The observance of an extruded first polar body within the zona pellucida of the oocyte signifies the advancement of the oocyte to the metaphase II (MI) stage of meiosis II. At this point, oogenesis is halted until activation of the MII oocyte either by fertilization or parthenogenetic activation. C) Schematic display of four genotypic possibilities for a haploid oocyte dependent upon on how the chromatids recombine (crossing over) and separate during meiosis I and II, respectively. Image modified from http://www.mun.ca/biology/desmid/brian/BIOL2060/BIOL2060-20/20\_16.jpg

Extrusion of the 2<sup>nd</sup> polar body is typically achieved by blockade of mitotic spindle activity either with cytochalasin B (Didie et al. 2013, Kim et al. 2007a) or treatment with 6-dimethylaminopurine (6-DMAP; Revazova et al. 2007, Turovets et al. 2011), a potent non-specific protein kinase inhibitor and cyclin-dependent kinase inhibitor, which disrupts proper mitotic spindle formation and thus results in the retention of the

now separated set of sister chromatids. An alternative method is by the treatment with puromycin, an antibiotic protein synthesis inhibitor, which does not inhibit the extrusion of the 2<sup>nd</sup> polar body (Sagi et al. 2016a, Sagi et al. 2016b). The resulting haploid pSCs can be maintained for some passages in a haploid state, but typically convert to diploid pSCs with a completely homozygous genome (Figure 3). In both cases, heterozygote and homozygote pSC, major histocompatibility complex (MHC) information, which is essential for the detection of self and non-self antigens, is typically homozygous because it is encoded in close distance to the centromere on the short arm of chromosome 6 in humans (reviewed in Horton et al. 2004), and chromosome 17 in mouse (Didie et al. 2013). The Rhesus macaque MHC homolog, Mamu, is located similarly as the human HLAs on chromosome 6. In contrast to the human chromosome 6, the Rhesus chromosome 6 is extended by genomic duplications (Huber et al. 2003). Because of the wide-spread encoding of the MHC information on chromosome 6 (telomere → centromere: HLA-A/MamuA → HLA-B/MamuB → HLA-DR/MamuDR; (Daza-Vamenta et al. 2004) creating fully homozygous pSC may be advantageous. MHC homozygosity results in reduced variability of MHC protein combinations and thus is considered an advantage for immune matching (Didie et al. 2013, Nakatsuji et al. 2008, Taylor et al. 2005, Taylor et al. 2012).

# 1.5 Directed cardiomyocyte differentiation

After the isolation and culture of human embryonic stem cells (hES) and the demonstration of their pluripotency, including the propensity to give rise to cardiomyocytes in 1998 (Thomson et al. 1998), suspension cultures of hES aggregates were introduced to study cardiomyogenesis in more detail (Kehat et al. 2001). Directed differentiation for increased cardiomyocyte yields were first established in monolayer cultures (Burridge et al. 2007, Burridge et al. 2011, Burridge and Zambidis 2013, Kattman et al. 2011, Laflamme et al. 2007, Lian et al. 2012, Lian et al. 2013, Xu et al. 2011) and further optimized for higher yields in stirring bioreactors (Chen et al. 2015, Kempf et al. 2014, Olmer et al. 2010, Riegler et al. 2015). All more recent protocols use defined factors and small molecules to modulate the Wnt-signal cascade (Figure 5) for optimal results (Hudson and Zimmermann 2011). Metabolic selection via glucose starvation may be used to further enhance cardiomyocyte purity (Sharma et al. 2015, Tohyama et al. 2013).

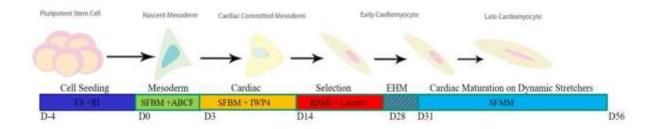


Figure 5: Directed differentiation of cardiomyocytes for EHM generation

Stem cells are seeded in feeder free culture and sequentially treated with small molecules for 14 days, followed by metabolic selection for cardiomyocytes. Following differentiation, cardiomyocytes are combined with fibroblasts in a collagen matrix to form EHMs and cultured on dynamic stretchers to facilitate auxotonic contractions (Tiburcy et al. 2017). A: activin-A, B: BMP4, C: CHIR99021, E8: mTeSR E8 medium, F: FGF-2, IWP4: inhibitor of Wnt production 4, RI: Rock Inhibitor Y27632, RPMI: Rosewell Park Memorial Institute medium, SFBM: Serum Free Basal Medium, SFMM: Serum Free Maturation Medium, d: days).

#### 1.6 Alginate encapsulation

The application of bioreactor cultures is essential for enhanced cardiomyocyte yields (Chen et al. 2015, Riegler et al. 2015). To control the cell growth and differentiation in stirred bioreactors, encapsulation has been applied (Jing et al. 2010, Serra et al. 2011). Sodium alginate is an anionic polysaccharide extracted from the cell walls of brown seaweed which can be used for controlled encapsulation of cells (Sidhu et al. 2012, Wilson et al. 2014). It is comprised of (1-4)-linked  $\beta$ -D-mannuronate(M) and  $\alpha$ -L-guluronate (G) residues (Figure 6) covalently linked as linear monomers of consecutive residues: M-blocks, G-blocks, or MG-Blocks. When alginate is exposed to a solution containing multivalent cations, such as calcium, the sodium ions are replaced, and the polymers become ionically cross linked and form a hydrogel (Figure 6). The encapsulation of stem cells in alginate creates a three-dimensional developmental niche like that provided during embryogenesis (Sidhu et al. 2012, Wilson et al. 2014). Alginate encapsulation has been used in a variety of applications such as: the culture of stem cells (Agarwal et al. 2013, Siti-Ismail et al. 2008), differentiation of stem cells (Dixon et al. 2014, Wilson et al. 2014), as a tissue engineering scaffold (Dahlmann et al. 2013, Rosellini et al. 2009, Ruvinov and Cohen 2016), as a delivery system for cell based therapy (Levit et al. 2013, Luca et al. 2014), and therapeutic drug delivery (Giovagnoli et al. 2014).

Figure 6: Polysaccharide residues of alginate

Alginate is a polysaccharide extracted from cell walls of brown seaweed comprised of  $\alpha$ -L-guluronate (G) and (1-4)-linked  $\beta$ -D-mannuronate (M) monomer residues (top). When exposed to a solution containing divalent ions, the sodium ions are replaced and the monomers become crosslinked as a hydrogel (bottom; images modified from Paredes Juarez et al. 2014).

#### 1.7 Engineering of heart muscle

The *in vitro* generation of heart tissue was developed as a three dimensional surrogate cardiac tissue for the investigation of various important themes in cardiac research: assessment of pharmaceutical interventions, biophysical development of cardiac organogenesis *in vitro*, and the feasibility of a engineered heart construct for the repair of damaged myocardium post myocardial infarction (Eschenhagen et al. 1997, Naito et al. 2006, Zimmermann et al. 2006, Zimmermann et al. 2000, Zimmermann et al. 2002, Zimmermann and Eschenhagen 2003). At first, engineered heart tissue was constructed using embryonic chicken cardiomyocytes in a collagen matrix (Eschenhagen et al. 1997) and neonatal rat cardiomyocytes in a colla-

gen/matrigel™ matrix (Zimmermann et al. 2000). These constructs were used to directly measure isometric contractile forces in response to increases in mechanical strain, electrical stimulation, and pharmacological stimuli. Further optimization of the tissue engineering protocols resulted in the successful generation of engineered heart muscle (EHM) from human ESC (Soong et al. 2012, Tiburcy and Zimmermann 2014), mouse pSC (Didie et al. 2013), and human iPS cells (Streckfuss-Bomeke et al. 2013). In the most recent protocol, EHM are constructed under serum-free conditions from defined mixtures of cardiomyocytes derived from ESC or iPS cells and fibroblasts in a collagen matrix (Tiburcy et al. 2017).

EHM culture resulted in a so far unsurpassed degree of maturation in ESC- and iPS cell-derived cardiomyocytes (Tiburcy et al. 2017). This includes the presence of Frank-Starling and Bowditch responses as well as responsiveness to pharmacological interventions similar to what can be observed in *bone fide* myocardium. An unbiased RNAseq experiment confirmed the advanced degree of maturation, but also demonstrates that maturation remains low compared to adult myocardium. Despite this caveat, application in heart repair (Qin et al. 2016, Riegler et al. 2015) and disease modelling (Tiburcy et al. 2017) has been attempted successfully.

#### 1.8 Animal models

Multiple animal models have been utilized in the study of heart failure. Canines were originally used in the 1970's to study cell death due to myocardial ischemia (Reimer et al. 1977, Reimer and Jennings 1979) and later for the studies of reperfusion (Jugdutt and Menon 2004, Przyklenk et al. 1986). Major drawbacks of the canine as a surrogate human heart failure model include its myocardial collateral circulation (White et al. 1986) and the inability to easily and accurately produce a myocardial infarction phenotype (Sabbah et al. 1991). Pigs and sheep have become reasonable alternatives to canines for myocardial studies due to their lack of myocardial collateral vasculature and similar gross cardiac anatomy to humans (Weaver et al. 1986), and the ability to create predictable and reproducible myocardial infarctions (Gorman et al. 1998, Jackson et al. 2002, Markovitz et al. 1989). Pigs and sheep have been used as human surrogates for testing of non-pharmaceutical therapies (Blom et al. 2005, Sabbah et al. 2003), pharmacological therapies (Lu et al. 2007), heart valve

replacement (Flanagan et al. 2009, Honge et al. 2010), and cell injection therapies (Zeng et al. 2007, Zhang et al. 2007).

Evolutionarily, non-human primates are the most closely related large animals to humans used as surrogates in scientific research (reviewed in Phillips et al. 2014). Humans belong to the family Hominidae or "Great apes" along with chimpanzees, gorillas, and orangutans. NHPs from the great apes have not been used in research in Europe since 1999 (Prescott 2010). Old World and New World monkeys are the most predominately used NHPs in scientific research respectively (Carlsson et al. 2004).

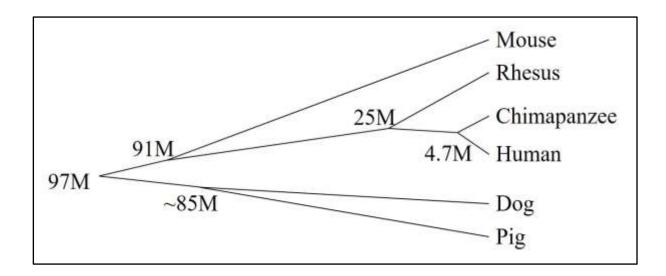


Figure 7: Evolutionary distance of mammalian test models

Humans and chimpanzees belong to the Family Hominidae with the other Great Apes and share the closet genetic similarities. Since 1999 Great apes have not been used in Europe for scientific research. Therefore, the evolutionary distance of 25 million years between the Rhesus macaque and human make them the most closely related mammalian species available for large animal model testing.

Macaques (*M. mulatta* [rhesus macaque], *M. fascicularis* [crab-eating macaque], *M. nemestrina* [pig-tailed macaque]) are considered Old World monkeys and are the most widely used NHP for late preclinical studies (reviewed in Phillips et al. 2014). The Rhesus macaque is the best characterized macaque species with similar cardi-

ovascular physiology compared to the human (Table 2) and shares 97% of its genome sequence with the human (Rogers and Gibbs 2014). Macaques have been used extensively as an alternative human model for infectious diseases (reviewed in Gardner and Luciw 2008), gene therapy (Lozier et al. 1999), cancer research (Brewer et al. 2001), reproduction (Lee et al. 2011, Nichols et al. 2010, VandeVoort et al. 2015), neurosciences (Bjugstad et al. 2008, Morin et al. 2014), and heart research (Bel et al. 2010, Chong et al. 2014, Kawamura et al. 2016, Shiba et al. 2016).

	Heart weight (grams)	Heart beats/minute	Systolic pressure
Human	360-480	60-90	60-120 mmHg
Mouse	0.14-0.15	500-600	80-160 mmHg
Rabbit	9-11	120-300	70-170 mmHg
Dog	160-420	60-120	120-150 mmHg
Pig	400-500	65-75	70-130 mmHg
Sheep	240-360	70-80	80-120 mmHg
Rhesus (Small)	30-40	125-240	122-194 mmHg
Rhesus (Large)	36-48	95-235	68-172 mmHg

Table 2: Comparative heart anatomy and physiological data between human and common heart disease animal models

Rhesus data from Max Planck Institute for Biological Cybernetics: http://hirnforschung.kyb.mpg.de/fileadmin/uploads/files/Methoden/Bet%C3%A4ubun gsverfahren\_und\_Chirurgie/SOP\_About\_Monkeys.pdf (table modified from Gandolfi et al. 2011)

## 1.9 Dissertation Objectives

Human stem cells have been successfully used as a source to generate cardiomyocytes for tissue engineering applications (Tiburcy et al. 2017). Due to the significant similarities between Rhesus macaque and humans, it is postulated that Rhesus induced pluripotent stem (RhiPS) cells can be directly differentiated into cardiomyocytes for future use in preclinical testing of myocardial infarction repair strategies. As an alternative and potentially immunologically more compatible stem cell source for allografting, methods to derived parthenogenetic stem cells were explored.

The following hypotheses were tested:

- Cardiomyocytes can be differentiated effectively from Rhesus macaque iPS cells.
- 2. EHM with functional properties of native myocardium and similar properties to human EHM can be constructed from Rhesus macaque iPS cells.
- 3. Pluripotent parthenogenetic stem cells can be generated from unfertilized, superovulated Rhesus macaque oocytes.

#### 2. Materials and Methods

Materials used in this research that are not listed under each section can be found in the Appendix.

#### 2.1 Animal experiments

Animal experiments – oocyte harvest from Rhesus macaque at the German Primate Center (DPZ) - were approved by the Niedersächsisches Landesamt für Verbraucherschutz und Lebensmittelsicherheit (LAVES – 33.9-42502-04-13/1363). Animal care was provided the veterinarians (Dr. med. vet. Charis Drummer and Sophie Mißbach) and animal care takers (Daniel Aschoff, Philip Rose and Carina Hunger) at the DPZ. Dr. med. vet. Charis Drummer and Sophie Mißbach performed all surgeries.

#### 2.1.1 Superovulation of Rhesus Macaque (Macaca mulatta)

Five female rhesus macaques (Table 3) were caged individually or in pairs in the same room with a 0600 – 1800 light cycle and maintained at a temperature of 25–27 °C at the DPZ animal housing facility.

Animal number	Years of age	Body weight (kg)		
2483	6.5	4.9-5.5		
2616	4.6	5.1-5.6		
2500	5.8	6.5-6.9		
2506	5.8	7.0-7.5		
2551	5.0	5.3-5.6		

Table 3: Female Rhesus macaques used for superovulation and oocyte harvest.

The cages were positioned to enable socialization with each other. Animals were provided food and water *ad libitum*. Seasonal produce, seeds, and cereal were offered as supplements for environmental enrichment. Females were observed daily for signs of vaginal bleeding and individual menstrual cycles were recorded by the animal care takers at the DPZ to identify the optimal time-point for superovulation

treatment. The first sign of bleeding was assigned as "menses cycle day 1". Hormone injection (injection day 0) was done as outlined below on menses cycle day 1, 2, 3 or 4 depending upon scheduling for the oocyte aspiration procedure. During superovulation cycle 1 all macagues received 37.5 International Units (IU) of recombinant human FSH (hFSH; Gonal F, 75 IU, Merck Serono) was administered intramuscularly (IM) twice daily for 6-8 days starting on injection day 0. For super ovulation cycles 2 and 3, Rhesus macaques greater than 6 kilograms received 37.5 International Units (IU) of recombinant human FSH (hFSH; Gonal F, 75 IU, Merck Serono) IM twice daily for 6-8 days starting on injection day 0. Rhesus macaques less than 6 kilograms were administered 18.75 IU of recombinant hFSH IM, twice daily for 6-8 days starting on injection day 0. Optimal stimulation of ovaries was identified, via ultrasound observation, as a total of 10 follicles with size measurements greater than 6 mm/follicle (Trounson et al. 2001). Once optimal stimulation was established and regardless of weight, an injection of 1,000 IU of recombinant human chorionic gonadotropin (hCG; Ovitrelle 250mg, Merck Serono) was administered IM, 24-36 hours before scheduled ovarian follicle aspiration (2 cycles) or ovariectomy (3<sup>rd</sup> cycle). Ultrasound (General electric-"Logiq-e" with 8 MHZ probe) guided transabdominal aspiration of the follicles to retrieve oocytes and ovariectomy were both performed by primate veterinarians in a surgical operating room while the animals were under general anesthesia. Anesthesia plans were designed for each animal individually by DPZ veterinarians (Appendix A7).

# 2.1.2 Collection of oocytes via follicle aspiration

Between 24-36 hours after hCG injection, macaques were anesthetized by intramuscular ketamine injection based on individual body weight (Appendix A7) and transported by large animal transporter (Petmate Vari Kennel) to the operating suite. Ovaries were located, photographed, and measured by ultrasound. Follicle aspiration was performed as previously described by Vandevoort et al (2003). Briefly, ovarian follicles were punctured using a single lumen IVF aspiration needle (Gynetics, Belgium) and oocytes were manually aspirated by vacuum suction with a 50 ml syringe attached to the needle. Aspirated oocytes were collected in a 15 ml polypropylene (PP) tube containing warm Tyrode's Albumin Lactate Pyruvate-4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (TALP-HEPES; Table 4) and maintained at >35° C in a heated specimen transport incubator (Cell-Trans 4016, Labotect). For parthenoge-

netic induction and culture, oocytes were transferred to the Department of Degenerative Diseases at the DPZ.

Component	MW	Final [mmol/L]	g/50 ml	g/100 ml
NaCl (Sigma S-9888)	58.44	127	0.3710	0.7420
KCL (Sigma P-3911)	74.55	3.16	0.0118	0.0236
CaCL <sub>2</sub> *2H <sub>2</sub> O Sigma C-7902)	147.02	2.00	0.0147	0.0294
MgCl <sub>2</sub> *6H <sub>2</sub> O (Sigma M-	203.30	0.60	0.0051	0.0102
2393)				
Glucose (Sigma G-8270)	180.16	5.00	0.0451	0.0901
NaH <sub>2</sub> PO <sub>4</sub> *H <sub>2</sub> O (Fisher	137.99	0.35	0.0024	0.0048
BP330-500)				
NaHCO <sub>3</sub> (Sigma S-8875)	84.01	2.00	0.0084	0.0168
HEPES – Na Salt (Sigma H-	260.30	5.00	0.0651	0.1302
3784)				
HEPES – Free Acid (Sigma	238.30	5.00	0.0596	0.1192
H-4034)				
Penicillin G (Sigma P-7794)			0.0030	0.0060
Streptomycin Sulfate (Sigma			0.0030	0.0060
S-9137)				
Na Lactate (60% syrup)	112.10	13	0.0926	0.1852
(Sigma L-1375)				
PVA (Sigma P-8136)	30,000-	0.1 mg/ml	0.0050	0.0100
	70,000			

Table 4: TALP-HEPES medium for follicle aspiration.

Combine all components in 90% of total volume in embryo tested water. Osmolarity was measured on an osmometer (Advanced Instruments Model 3320). Osmolarity was adjusted 290 mOsm by adding embryo tested water calculated by [(current osmolarity/290) x current volume] = total volume. Total volume – current volume = volume of embryo tested water to be added. The pH was measured using a pH meter (WTW model pH 323) and adjusted to  $7.4 \pm 0.05$  with the addition of 1 N Sodium

Chloride (Roth K025.1) or 1 N Sodium Hydroxide (Roth K021.1). Medium was passed through a 0.22 µm syringe filter. Modified from Boatman (1987).

# 2.1.3 Collection of oocytes via ovariectomy

Between 24-36 hours after hCG injection, macaques were anesthetized by ketamine injection (Appendix A6) and transported by animal transporter (Petmate Vari Kennel) to the operating suite. Ovaries were removed via midline incision and placed in a 15 ml PP tube containing warm TALP-HEPES and maintained at ≥35° C in a heated specimen transport incubator (Cell-Trans 4016, Labotect, Rosdorf, Germany). For parthenogenetic induction and culture, oocytes were transferred to the Institute of Pharmacology and Toxicology at the University Medical Center Göttingen. Ovaries were washed with warm TALP-HEPES and cumulus oocyte complexes (COCs) were manually dissected from the ovarian follicles with a scalpel under a laminar flow cell culture hood. COCs were collected in 6 cm culture dish containing TALP-HEPES and prepared for parthenogenetic activation (section 2.3.1).

# 2.1.4 Collection of oocytes via necropsy

Unstimulated ovaries were provided by the DPZ following euthanasia of a sixth female (#2379) for medical reasons as determined by the veterinarians at the DPZ. Ovaries were collected in a 15 ml PP tube containing warm phosphate buffered saline and maintained at ≥35° C in a heated specimen transport incubator (Cell-Trans 4016, Labotect, Rosdorf, Germany) and transferred to the Institute of Pharmacology and Toxicology at the University Medical Center Göttingen. Ovaries were washed with warm TALP-HEPES and COCs were manually dissected from the ovarian follicles with a scalpel under a laminar flow cell culture hood. COCs were collected in a 6 cm culture dish containing TALP-HEPES and prepared for *in vitro* maturation (IVM; Section 2.2.6).

#### 2.1.5 Teratoma assay

Non-Obese Diabetic Severe Combined Immune Deficiency (NOD SCID) mice (n = 4) housed at the Institute for Human Genetics (Göttingen) were injected with 5 x  $10^6$  Sendai iPS cells in suspension in hES medium (Appendix A1) into the left flank of individual mice by Prof. Dr. Ralf Dressel. The hES medium was comprised of Dulbecco's modified eagle medium (DMEM; Invitrogen) supplemented with 20% Knock-

out serum replacement (KOSR; Invitrogen) 100 U/ml Penicillin and 100 µg/ml Streptomycin and FGF-2 (10 ng/ml). Mice were monitored for the formation of tumors. Following formation of palpable tumors, mice were euthanized and tumors resected, weighed, and histological analysis was performed.

## 2.2 Cell models

Human and Rhesus macaque cells (fibroblasts and oocyte derived cells) were utilized. Human foreskin fibroblasts (hFF) were purchased from the American Type Culture Collection (SCRC-1041; ATCC). Rhesus embryonic stem cells (E366.4) were purchased from WiCell. Rhesus dermal (RhDFB) and gingiva fibroblasts (RhGFB) were obtained from the DPZ (Prof. R. Behr). Rhesus iPS cells were generated at the Institute of Pharmacology and Toxicology (Dr. M. Tiburcy) from a skin biopsy provided by the DPZ or at Stanford University (Dr. M. Wang/Prof. J. Wu) using retroviral or Sendai virus transduction of Yamanaka factors, respectively (Takahashi and Yamanaka 2006).

## 2.2.1 Human foreskin fibroblasts

hFFs were maintained in hFF medium consisting of Dulbecco's modified eagle medium (DMEM; Gibco) with 150  $\mu$ l/ml of fetal bovine serum (FBS; PAA) and 100 U/ml Penicillin and 100  $\mu$ g/ml Streptomycin (P/S; Invitrogen) and supplemented with 20 ng/ml of fibroblast growth factor (FGF-2; Miltenyi Biotech). Multiple passages not exceeding passage 30 were used for all subsequent experiments.

# 2.2.2 Rhesus fibroblasts

Rhesus gingival fibroblasts (RhGFB) used for EHM production were isolated from biopsy of the buccal mucosa from a 15-year-old euthanized male from the DPZ. Additional gingival biopsies were obtained from five female Rhesus macaques undergoing surgical operations under anesthesia. Biopsies were transported from the surgery suite to the research facility in warm PBS with 100 U/ml Penicillin and 100 µg/ml Streptomycin and 2.5 µg/ml Amphotericin B. Biopsies were placed on a Synthemax (Corning) coated dish and cut into 1-2 mm pieces with a sharp scalpel and spread around dish to facilitate cellular outgrowth. Pieces were covered with serum and incubated for 30 minutes at 37 °C and 5% CO<sub>2</sub> to ensure attachment of tissue to plate. Tissues were further incubated in hFF medium with 2.5 µg/ml Amphotericin B at 37

°C and 5% CO<sub>2</sub> for approximately 14 days with medium changes every other day. Once outgrowths from the primary biopsies began to appear, mechanical dissociation was performed and cells were propagate as described below.

#### 2.2.3 Fibroblast cell culture

Primary fibroblasts (hFF, RhDFB or RhGFB) were cultured in hFF medium (section 2.2.1). Cultures were maintained in a humidified incubator at 37 °C and 5% CO<sub>2</sub>. Routine passaging of fibroblasts was performed when confluency reached a maximum of 90% based on visual inspection. Cells were rinsed once with PBS (warmed to 37 °C). Cellular dissociation was performed using pre-warmed Tryple™ Express Enzymatic Dissociation Solution (ThermoFisher Scientific) for 2-5 minutes at 37 °C. Visual inspection of cell detachment was monitored with a light microscope during dissociation. Following the dissociation of the cells form the culture flask 2x volumes of hFF medium were added and gently pipetted to ensure cell clumps were broken up into a single cell suspension. Dissociated cells were pelleted through centrifugation at 300 g for 5 minutes and resuspended in fresh medium. Cells were counted (section 2.6.1) and seeded onto new culture dishes at the required density, or used for EHM generation (section 2.5.2). Aliquots of fibroblasts were frozen in cryovials for storage purposes at concentrations of 3 x 10<sup>6</sup> in freezing medium (90:10%) FBS:DMSO). Cell suspensions were transferred into cryovials (1 ml cell suspension/vial), and immediately placed into an insulated freezing container (Mr. Frosty, ThermoFisher Scientific) filled with isopropanol and placed at -80 °C, to freeze cells slowly enough to maintain viability. Previously frozen cells to be used for experiments were thawed rapidly in warm water bath followed by the addition of 1 ml of fresh hFF medium (warmed to 37 °C) into the cryovial to resuspend the cells. Thawed cells were immediately transferred into 15 ml PP tubes with additional warmed hFF medium and centrifuged at 300 g for 5 minutes at RT and resuspended in hFF medium, to remove DMSO. Cell suspensions were then plated accordingly to be used for subsequent experiments requiring fibroblast feeder layers for culturing.

## 2.2.4 Preparation of fibroblast feeder layers

Mitotically inactivated hFFs were used as feeder layers for standard culturing of undifferentiated Rhesus iPS cells. Culturing the stem cells on an inactivated fibroblast feeder layer provides a supportive environment for the Rhesus stem cells without

dividing and diluting the cultivation of undifferentiated stem cells (Fuegemann et al. 2010, Soong et al. 2012, Wobus et al. 1991). Prior to use as a cell culture feeder layer hFFs were mitotically inactivated by y-irradiation (30 Gy). To produce large stocks of inactivated fibroblasts, hFFs were cultured in T175 culture flasks in hFF medium until a maximum of 90% confluency was reached. Fibroblasts were rinsed once with PBS (warmed to 37 °C) in a volume appropriate for the size of the culture flask. Cellular dissociation was performed using pre-warmed Tryple™ for 2-5 minutes at 37 °C. Visual inspection of cell detachment was monitored with a light microscope during dissociation. Following the dissociation of the cells from the culture flask, 2x volumes of hEF medium were added to stop the enzymatic reaction. Dissociated fibroblasts were transferred into a 50 ml conical PP tube and centrifuged at 300 g for 5 minutes. After centrifugation cells were resuspended in fresh hFF medium and counted following the procedure found in section 2.6.1. Appropriate viable cell numbers (2 x 10<sup>6</sup>/ml), based on counting results, were transferred into new 15 ml PP tubes prior to irradiation. Fibroblasts were exposed to one cycle of 30 Gy for 15 minutes. After irradiation, cell suspensions were counted again (section 2.6.1) and frozen down into cryovial stocks (8 x 10<sup>6</sup> cells/ml) in freezing medium until further use or immediately plated at a seeding density of 40,000/cm<sup>2</sup> in hEF medium.

# 2.2.5 Preparation of Rhesus fibroblasts for engineered heart muscle

Culturing of RhGFB was performed in T175 or T225 (Nunc) culture flasks to ensure sufficient numbers of fibroblasts for EHM construction. Culture flasks were coated with 10-15 ml of 0.1% gelatin to facilitate cell attachment. Flasks were incubated for 20 minutes at 37 °C with 5% CO<sub>2</sub>. Following solidification of gelatin, the remaining fluid component was aspirated. A cryovial of previously frozen cells was thawed by adding prewarmed hFF medium to the frozen pellet and gently pipetted. The medium and cells were transferred into a 15 ml PP tube and centrifuged at 300 g for 5 minutes. Following centrifugation media was aspirated and the pellet was resuspended in warmed hFF media. Cells were counted using a CASY counter (section 2.6.1). An appropriate volume of cell suspension, based on viable cell content obtained from counting, was then pipetted into each flask, containing an appropriate volume of warmed hFF media plus FGF-2, at a density of 17,000 cells/cm<sup>2</sup>.

# 2.2.6 In vitro maturation of Rhesus oocytes

Rhesus oocytes were obtained from the DPZ following a routine necropsy. After manual dissection from the ovarian follicles COCs were incubated at 37  $^{\circ}$ C in a humidified incubator with 5% CO<sub>2</sub> and 5% O<sub>2</sub> for 24 hours in IVM medium (Appendix A8). After IVM COCs were parthenogenetically activated (Section 2.3.1).

#### 2.3 Stem cells

# 2.3.1 Rhesus macaque parthenogenetic stem cells

Rhesus parthenogenetic stem cell lines 2551 (RhpSC-2551) and 2616 (RhpSC-2616) were generated via parthenogenetic activation of MII phase oocytes. Following aspiration of follicles (ovulation cycles 1 and 2), ovariectomy (ovulation cycle 3), or IVM (ovaries obtained from necropsy), oocytes were first exposed to 0.5 mg/ml hyaluronidase (Type IV; Sigma Aldrich) for 5-10 minutes to remove cumulus cells followed by three washes with TALP-Hepes medium. After cumulus cell removal, all oocytes were subjected to a five-minute incubation with a calcium ionophore, lonomycin (5 mmol/L, Sigma), followed by either 4-hour incubation with 6-DMAP (1 mmol/L; Sigma Aldrich; ovulation cycle 1) or a 5-hour incubation with puromycin (5 μg/ml; Gibco; ovulation cycles 2 and 3) at 37 °C in a humidified incubator with 5% CO<sub>2</sub> and 5% O<sub>2</sub>. All activation chemicals were prepared in Hamster Embryo Culture Medium 9 (HECM-9) base medium (Table 5). After activation oocytes from ovulation cycles 1 and 2 were transferred into drops of HECM-9 medium covered by mineral oil (or liquid paraffin) and incubated at 37 °C in a humidified incubator with 5% CO<sub>2</sub> and 5% O<sub>2</sub>. Oocytes were transferred to a fresh HECM-9 medium drop after the first 18-20-hours post activation and then every 48 hours until blastocyst formation was observed. Prior to transferring, photos were taken using Zeiss Discovery V8 and Octax Eyeware software. Following activation, oocytes from ovulation cycle 3 were incubated using a Vitrolife Embryoscope™. Oocytes remained in the original drop of HECM-9 medium for the duration of culture and images were automatically obtained via the embryoscope software program every 30-45 minutes until morula/blastocyst stage was achieved. Mature metaphase II (MII) oocytes that were successfully activated and developed to morula stage or further were removed from culture and plated on feeder layers.

Component	MW	Final [mmol/L]	g/50 ml	g/100 ml
NaCl (Sigma S-9888)	58.44	127	0.3710	0.7420
KCL (Sigma P-3911)	74.55	3.16	0.0118	0.0236
CaCL <sub>2</sub> *2H <sub>2</sub> O Sigma C-7902)	147.02	2.00	0.0147	0.0294
MgCl <sub>2</sub> *6H <sub>2</sub> O (Sigma M-	203.30	0.60	0.0051	0.0102
2393)				
NaHCO <sub>3</sub> (Sigma S-8875)	84.01	2.00	0.0084	0.0168
Na Lactate (60% syrup)	112.10	13	0.0926	0.1852
(Sigma L-1375)				
PVA (Sigma P-8136)	30,000-	0.1 mg/ml	0.0050	0.0100
	70,000			
1 N HCL solution	1.4 µl/ml		70 µl	140 µl

Table 5: HECM-9 medium for oocyte activation and culture.

Combine all components in 90% of total volume in embryo tested water. Osmolarity was measured on an osmometer (Advanced Instruments Model 3320). Osmolarity was adjusted 290 mOsm by adding embryo tested water calculated by [(current osmolarity/290) x current volume] = total volume. The pH was measured using a pH meter (WTW model pH 323) and adjusted to  $7.4 \pm 0.05$  with the addition of 1 N Sodium Chloride (Roth K025.1) or 1 N Sodium Hydroxide (Roth K021.1). Combine with 100X AA/P solution before use (Table 6).

Component	Sigma Catalog #	MW	Final concentration	g/10 ml
			[mmol/L]	
Taurine	T-8691	125.1	0.50	0.0626
Asparagine	A-4159	132.1	0.01	0.0013
Cysteine	C-6852	175.6	0.01	0.0018
Histidine	H-5659	209.6	0.01	0.0021
Lysine	L-8662	182.6	0.01	0.0018
Proline	P-5607	115.1	0.01	0.0012
Serine	S-4311	105.1	0.01	0.0011
Aspartic Acid	A-4534	133.1	0.01	0.0013
Glycine	G-8790	75.07	0.01	0.0008
Glutamic Acid	G-5889	169.1	0.01	0.0017
Glutamine	G-8540	146.1	0.20	0.0292
Pantothenic	P-5155	238.3	0.003	0.0007
Acid				

Table 6: Amino acid/Pantothenate solution for oocyte activation and culture Combine with 10 milliliters of embryo tested water. Add 10 µl per 1 ml of HECM-9 medium.

Successfully activated Rhesus oocytes that progressed to the morula stage or further, were transferred onto fibroblast feeder cultures in attempt to propagate a cell line. To do so, morula stage cells were briefly exposed to 0.5% pronase to degrade the zona pellucida. Degradation was directly observed under a dissecting microscope and morulae were gently pipetted until the central aggregation of blastomeres was released from the degrading zona pellucida. The blastomeres were then immediately transferred onto fibroblast feeder cultures in hES medium supplemented with 10% human umbilical cord blood serum, 4 ng/ml FGF-2, and 5 ng/ml leukocyte inhibiting factor (LIF; cycle 1; Revazova et al. 2007) and incubated at 37°C and 5% O<sub>2</sub>. The blastomere culture medium was amended with the increase of FGF-2 to 10 ng/ml for cycles 2 and 3. Following transfer onto mitotically inactivated human fibro-

blast feeder cells, RhpSC 2551 (cycle 3) blastomere outgrowths began to spontaneously differentiate. Therefore, reprogramming of the cells was performed using the Cytotune 2.0 Sendai virus reprogramming kit (ThermoFisher Scientific). Blastomere outgrowths from RhpSC 2616 (cycle 3) proceeded normally. No blastomeres from cycles 1 or 2 generated successful outgrowths on feeder cultures.

# 2.3.2 Rhesus Induced Pluripotent Stem Cell line C1

Rhesus induced pluripotent stem cell line C1 (C1) was generated by Dr. Malte Tiburcy at the Institute of Pharmacology and Toxicology, University Medical Center Göttingen, using six factor lentiviral reprogramming with OCT4, SOX2, KLF4, c-MYC, LIN28 and NANOG (=OSKMLN; unpublished). OSKM were encoded on a STEMCAA vector (Somers et al. 2010). Nanog and Lin 28 were contained on two additional lentiviral vectors, pSin-EF2-Nanog-Pur and pSin-EF2-Lin28-Pur (Yu et al. 2007), respectively.

# 2.3.3 Rhesus Induced Pluripotent Stem Cell line 43110-4

Rhesus induced pluripotent stem cell line 43110-4 (Sendai) was generated in the laboratory of Dr. Joseph Wu at Stanford University using Cytotune 2.0 (Thermo Fisher) Sendai viral reprogramming kit and generously provided for use in this project.

#### 2.3.4 Stem cell culture

Rhesus stem cell cultures were cultured on inactivated hFF cell feeder layers in hES medium. Cultures were maintained in a humidified incubator at 37 °C and 5% CO<sub>2</sub>. Routine passaging of stem cells was performed when confluency reached a maximum of 90% based on visual inspection. Cells were rinsed once with PBS (warmed to 37 °C) in a volume appropriate for the size of the culture dish. Cellular dissociation was performed using pre-warmed Tryple™ for 2-5 minutes at 37 °C. Visual inspection of cell detachment was monitored with a light microscope during dissociation. Following the dissociation of the cells from the culture flask 2x volumes of hES medium were gently added ensuring that stem cells and fibroblasts were removed in an intact sheet. The cell sheet was gently agitated in the medium to free any loose stem cells without too many fibroblasts in cell suspension. Medium with stem cells was gently pipetted into PP tubes, making sure to leave the fibroblast cell sheet behind.

Dissociated cells were pelleted through centrifugation at 300 g for 5 minutes and resuspended in fresh hES medium. Cells were counted using a CASY counter (section 2.6.1) and seeded onto new fibroblast feeders at the required density or used for EHM generation (section 2.5.2). Aliquots of dissociated stem cells were frozen in cryovials and stored until further use. Cells were re-suspended at a concentration of 2 x 10<sup>6</sup> cells/ml in freezing medium and transferred into cryovials (1 ml/vial) and immediately placed in an insulated freezing container (Mr. Frosty, ThermoFisher Scientific) and placed at -80 °C, to freeze cells slowly enough to maintain viability. Frozen cells to be used for experiments were thawed rapidly at 37 °C and washed in 1 ml of fresh hES medium (warmed to 37 °C). Thawed cells were immediately transferred into 15 ml PP tubes with additional warmed medium and centrifuged at 300 g for 5 minutes at RT to remove DMSO. Supernatant was removed and the cell pellet was resuspended in hES medium. Cell suspensions were then plated onto hFF feeders and cultured in hES medium with FGF-2 (10 ng/ml) in a humidified incubator at 37 °C and 5% CO<sub>2</sub>.

# 2.3.5 Rhesus Mamu sequencing

Cell cultures of Lenti- and Sendai-iPS lines were provided to Prof. Dr. Lutz Walter at the DPZ Department of Primate Genetics for MHC Class I and Class II Mamu screening. Sendai iPS cell-derived cardiomyocytes were treated with 100 units/ml recombinant human interferon gamma (IFNy, peprotech; Suarez-Alvarez et al. 2010) or 50 µg/ml lipopolysaccharides (LPS; Sigma-Aldrich) for 24 hours at 37 °C in a humidified incubator with 5% CO<sub>2</sub> to elicit MHC Class I Mamu overexpression. RNA was isolated from the cell cultures and amplified by PCR by Nico Westphal from the Walter Lab at the DPZ and sequenced at the Transcriptome and Genome Analysis Laboratory (TAL Göttingen). Total RNA was extracted from cells using the following Trizol protocol. Cells were removed from culture flasks using a cell scraper and collected in a 1.5 ml tube. 360 µl of TRIzol Reagent was added to the tube cells were pipetted up and down to homogenize them. Homogenized samples were incubated for 5 minutes at RT to dissociate the cells and then centrifuged at 12,000 g for 10 minutes at 4 °C. Supernatant was transferred to a fresh 1.5 ml tube and 72 µl of chloroform was added to the tube and vortexed for 30 seconds. Samples were centrifuged at 12,000 g for 15 minutes at 4 °C. Following centrifugation, the aqueous phase (upper phase) containing the RNA was transferred into a fresh 1.5 ml tube.

Precipitation of the RNA from the aqueous phase was achieved by adding 180  $\mu$ l of isopropyl-alcohol (isopropanol) and 1  $\mu$ l Glyco blue to the sample. Samples were vortexed and precipitated overnight at -20 °C. Following overnight precipitation samples were centrifuged at 12,000 g for 45-55 minutes at 4 °C. After centrifugation, supernatant was removed with a pipette and the pellet was washed with 0.5 ml of 75% EtOH and centrifuged at 12,000 g for 10 minutes at 4 °C. After centrifugation, the supernatant was removed with a pipette and the pellet was washed with 0.5 ml of 75% EtOH and centrifuged at 12,000 g for 10 minutes at 4 °C. After centrifugation, the supernatant was removed with a pipette and the pellet dried at 37 °C. The dried RNA pellet was dissolved in 10-20  $\mu$ l of RNase free water and mixed by pipetting and stored at -80 °C. A 5  $\mu$ l sample was checked for quality and quantity.

Amplification of the MHC Class I RNA was performed with cDNA generated using a Transcriptor high fidelity cDNA synthesis kit from Roche according to the manufacturer's instructions. PCR reaction used the primers listed in Table 7A for MHC Class I and primers from Table 8A for MHC Class II with FastStart™ High Fidelity PCR System polymerase (Roche). Amplification of MHC Class I was performed according to Table 7B. Amplification of MHC Calls II was performed according to table 8B. Following amplification of the cDNA libraries, PCR products were purified with the MinElute PCR purification kit (Qiagen) and combined into pooled samples with concentrations of 10 nmol/L and sequenced on a Miseq sequencer (Illumina) at the TAL Göttingen.

Gene transcripts	Forward primer	Reverse primer	Base Pairs
MHC I	5'-GCT ACG TGG ACG	5'-TCC CAC TTS CGC	F = 17
IVITO	ACA CG -3'	TGG GT-3'	R = 17

Table 7A: Oligonucleotide sequences for Rhesus MHC class I PCR amplification

Cycle Step	Temperature	Time	Cycles
Initial Denaturation	95 °C	5 minutes	

Denaturation	95 °C	15 seconds	
Annealing	52 °C	45 seconds	35 X
Extension	72 °C	45 seconds	
Final extension	72 °C	5 minutes	1X

Table 7B: Amplification program for MHC Class I

Gene	Forward primer	Reverse primer	Base Pairs
transcripts			
	5'- CGT ATC GCC TCC		
	CTC GCG CCA TCA	CCA GCC CGC TCA	F = 57
DPA-1	GAC GAG TGC GTA	GAC GAG TGC GTC	R = 50
	TGC GCC CTG AAG	CCC TGG GCM CGG	K = 00
	ACA GAA TGT -3'	GG -3'	
	5'- CGT ATC GCC TCC	5'- CTA TGC GCC TTG	
	CTC GCG CCA TCA	CCA GCC CGC TCA	F
DPA-2	GAC GCT CGA CAA	GAC GCT CGA CAC	F = 57
	TGC GCC CTG AAG	CCC TGG GCM CGG	R = 50
	ACA GAA TGT -3'	GG -3'	
	5'-CGT ATC GCC TCC	5'-CTA TGC GCC TTG	
	CTC GCG CCA TCA	CCA GCC CGC TCA	Г 57
DPA-3	GAG ACG CAC TCA	GAG ACG CAC TCC	F = 57
	TGC GCC CTG AAG	CCC TGG GCM-3'	R = 50
	ACA GAA TGT -3'	CGG GG -3'	
	5'- CGT ATC GCC TCC	5'- CTA TGC GCC TTG	
	CTC GCG CCA TCA	CCA GCC CGC TCA	F
DPA-4	GAG CAC TGT AGA	GAG CAC TGT AGC	F = 57
	TGC GCC CTG AAG	CCC TGG GCM CGG	R = 50
	ACA GAA TGT -3',	GG -3'	
	5'- CGT ATC GCC TCC	5'- CTA TGC GCC TTG	
DPB-1	CTC GCG CCA TCA	CCA GCC CGC TCA	F = 57
ו-סרט	GAC GAG TGC GTA	GAC GAG TGC GTT	R = 60
	TGA TGG TTC TGS	TAT GCA GAT CCT	

	AGG TWT CTG -3'	CGT TGA ACT TTC -3'	
DPB-2	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GCT CGA CAA TGA TGG TTC TGS AGG TWT CTG -3'	CCA GCC CGC TCA GAC GCT CGA CAT TAT GCA GAT CCT CGT TGA ACT TTC -3'	F = 57 R = 60
DPB-3	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG ACG CAC TCA TGA TGG TTC TGS AGG TWT CTG -3'	CCA GCC CGC TCA	F = 57 R = 60
DPB-4	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG CAC TGT AGA TGA TGG TTC TGS AGG TWT CTG -3'		F = 57 R = 60
DQA-1	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GAG TGC GTA TGA TCC TAA ACA AAG CTC TG -3',		F = 56 R = 57
DQA-2	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GCT CGA CAA TGA TCC TAA ACA AAG CTC TG -3'		F = 56 R = 57
DQA-3	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG ACG CAC TCA TGA TCC TAA ACA AAG CTC TG -3'	CCA GCC CGC TCA GAG ACG CAC TCT	F = 56 R = 57
DQA-4	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG CAC TGT AGA TGA TCC TAA ACA AAG CTC TG -3'	CCA GCC CGC TCA	F = 56 R = 57
DQB-1	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GAG TGC GTG AAR AAG KCT TTG CGG ATY CC -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAC GAG TGC GTC CCC AGC ACR ARG CCT CC -3'	F = 56 R = 53
DQB-2	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GCT CGA CAG AAR AAG KCT TTG CGG ATY CC -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAC GCT CGA CAC CCC AGC ACR ARG CCT CC -3'	F = 56 R = 53
DQB-3	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG ACG CAC TCG	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAG ACG CAC TCC	F = 56 R = 53

	AAR AAG KCT TTG CGG ATY CC -3'	CCC AGC ACR ARG CCT CC -3'	
DQB-4	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG CAC TGT AGG AAR AAG KCT TTG CGG ATY CC -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAG CAC TGT AGC CCC AGC ACR ARG CCT CC -3'	F = 55 R = 53
DRA-1	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GAG TGC GTA TGG CCA TAA GTG GAG TCC C -3'		F = 55 R = 55
DRA-2	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GCT CGA CAA TGG CCA TAA GTG GAG TCC C -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAC GCT CGA CAT TAC AGA GGC CCC CTG CGT T -3'	F = 55 R = 55
DRA-3	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG ACG CAC TCA TGG CCA TAA GTG GAG TCC C -3'		F = 55 R = 55
DRA-4	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG CAC TGT AGA TGG CCA TAA GTG GAG TCC C -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAG CAC TGT AGT TAC AGA GGC CCC CTG CGT T -3'	F = 55 R = 55
DRB-1	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GAG TGC GTG TGC TGA GCY CCC SAC TRG -3'	5'- CTA TGC GCC TTG	F = 54 R = 56
DRB-2	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAC GCT CGA CAG TGC TGA GCY CCC SAC TRG -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAC GCT CGA CAC AAG GAA GAG CAG GCC CAG CA -3'	F = 54 R = 56
DRB-3	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG ACG CAC TCG TGC TGA GCY CCC SAC TRG -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAG ACG CAC TCC AAG GAA GAG CAG GCC CAG CA -3'	F = 54 R = 56
DRB-4	5'- CGT ATC GCC TCC CTC GCG CCA TCA GAG CAC TGT AGG TGC TGA GCY CCC SAC TRG -3'	5'- CTA TGC GCC TTG CCA GCC CGC TCA GAG CAC TGT AGC AAG GAA GAG CAG GCC CAG CA -3'	F = 54 R = 56

Table 8A: Oligonucleotide sequences for Rhesus MHC class II PCR amplification

Cycle Step	Temperature	Time	Cycles
Initial Denaturation	95 °C	5 minutes	
Denaturation	95 °C	30 seconds	
Annealing	55 °C	30 seconds	35 X
Extension	72 °C	60 seconds	
Final extension	72 °C	7 minutes	1X

Table 8B: Amplification program for MHC Class II

# 2.4 Cardiomyocyte differentiation

## 2.4.1 Directed differentiation

iPS cells were enzymatically dissociated with Tryple™ (section 2.3.4) and plated at a seeding density of 25,000 cells/cm<sup>2</sup> on Matrigel™ coated culture flasks (T25, Corning) for the Sendai RhiPS cells or encapsulated at a density of 2 x 10<sup>6</sup> in alginate for the Lenti RhIPS cells (section 2.4.2) with approximately 25 free floating beads per T25 culture flask, in mTeSR™ E8 medium (E8, Stemcell Technologies) supplemented with 10 µmol/L Rock Inhibitor (RI; Y-27632); this starting time-point was recorded as differentiation day -4. Exchange with fresh E8 Medium with 5 µmol/L RI was performed after 48 hours (differentiation day -2). After four days of recovery and cellular expansion, directed differentiation under defined medium supplements was started (differentiation day 0). To initiate the mesodermal stage of differentiation, E8 medium was replaced by SFBM (serum-free basal medium; Appendix A2) supplemented with small molecule growth factors ABCF: Activin-A (R&D systems; 9 ng/ml), BMP4 (R&D systems; 5 ng/ml), CHIR99021 (Stemgent; 1 µmol/L) and FGF-2 (5 ng/ml). Medium was exchanged daily until day 3, at which point the cardiac specification stage was induced for the next 10 days. Small molecules ABCF were replaced in the SFBM medium by the addition of an inhibitor of Wnt production (IWP4; Stemgent; 5 µmol/L). Fresh medium was exchanged every 2-3 days until day 13. On day 14 the

differentiated cardiomyocytes underwent lactate selection, SFBM was replaced by a lactate selection medium (RPMI without glucose, 100 U/mI Penicillin, 100  $\mu$ g/mI Streptomycin, 2.2 mmol/L sodium lactate and 0.1 mmol/L 2-mercaptoethanol) and changed daily for 4-7 days dependent upon visual inspection for loss of non-myocytes. Following lactate selection cardiomyocyte cultures were returned to SFBM medium until differentiation day 28 when EHMs were generated (refer to section 2.5.2).

# 2.4.2 Alginate encapsulation

Undifferentiated Lenti-iPS cells were passaged with Tryple™ (section 2.3.4), counted using a CASY counter (section 2.6.1), and manually encapsulated in an alginate hydrogel. Briefly, a 1% alginate-gelatin encapsulation solution was prepared by mixing alginic acid sodium salt (MP Biomedicals) with 0.1% gelatin overnight at RT on an orbital shaker (Certomat® R; B. Braun) at 150 revolutions per minute. After the alginate salt dissolved completely in gelatin, 9% NaCl was added to the alginate-gelatin mixture and vortexed thoroughly for 3 minutes. For encapsulation, previously dissociated stem cells (2 x 10<sup>6</sup>/ml) were centrifuged at 300 g for 5 minutes. Supernatant was removed from the pellet and 1% alginate-gelatin solution plus 10 µmol/L RI was added to the pellet at a volume dependent on cell quantity. Cells and alginate were pipetted to ensure equal distribution of cells in suspension. Encapsulated beads were generated by pipetting 25 µl of the cell:alginate suspension dropwise into a CaCl<sub>2</sub> solution (50 mmol/L) in a swirled 50 ml PP tube. Beads stayed in the bath for five minutes. The contents of the PP tube were then decanted through a 70 µm cell strainer (BD Biosciences) to separate the alginate beads from the CaCl<sub>2</sub>. Beads were washed three times by dispensing 5 ml of CaCl<sub>2</sub> over the beads in the filter. The beads were then transferred into culture flasks (25 beads/T25; adapted from Sidhu et al. 2012). Beads were cultured according to section 2.4.1.

## 2.5 EHM Generation

## 2.5.1 Construction of EHM casting molds and dynamic stretchers

For the generation of EHM, circular molds (diameter: 10 mm) with a central silicone pole (diameter: 2 mm, length: 10 mm) were fabricated using a silicone mixture (Sylgard<sup>®</sup> 185 silicone mixture) in a multistep process (Figure 8A). Hollow silicone tubing

(diameter: 4 mm) was placed over the central poles of each mold to create a removable spacer in which the EHM condenses around. A more detailed protocol for the preparation of the casting molds was previously published (Soong et al. 2012, Tiburcy et al. 2014). Additionally, flexible silicone stretchers (length: 5 mm, thickness: 1.5 mm, inner/outer distance: 3/6 mm) with a circular silicone base (diameter: 12 mm, thickness: 2 mm) were fabricated for facilitation of auxotonic contractions of EHM (Fig. 8B). The fully assembled casting molds and the flexible stretchers can be autoclaved and reused.

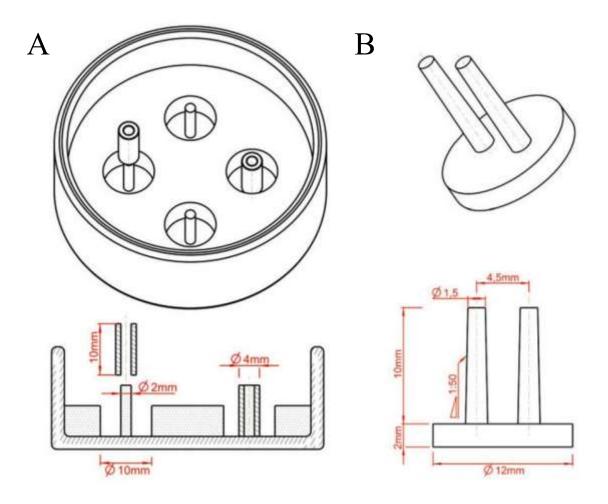


Figure 8: Reusable EHM mold and stretcher

Reusable silicone casting molds with silicone poles for EHM generation and elastic stretchers for mechanical loading and EHM maturation. (A) Glass culture dish filled with transparent silicone creating 4 circular recesses (diameter: 10 mm) with central poles (diameter: 2 mm, length: 10 mm). Each central pole has silicone tubing (outer diameter: 4 mm) placed over it to provide an inner diameter of 4 mm during EHM

generation. (B) Dynamic silicone poles (length: 10 mm, thickness: 1.5 mm, inner/outer distance: 3/6 mm) with silicone base (diameter: 12 mm, thickness: 2 mm). Images modified from Tiburcy et al., 2014.

# 2.5.2 Casting of EHM

A master mix for EHM (Table 9) generation was made by combining ice cold type I bovine collagen (0.4 mg/EHM), 2x RPMI, and pH neutralized with NaOH (0.1 mol/L) in a prechilled 50 ml PP tube (Table 9). Cardiomyocytes in single cell suspension (1.015 x 10<sup>6</sup>/EHM) and primary fibroblasts (0.435 x 10<sup>6</sup>/EHM) were combined in appropriate quantities depending on the desired number of EHMs in a 15 ml PP tube and centrifuged at 300 g for 5 minutes. Supernatant was removed from the pellet and the pellet was resuspended in the appropriate volume of pre-warmed Serum Free Maturation Medium (SFMM; Appendix A4) according to Table 9 making a cardiomyocyte:fibroblast cell suspension. The cell suspension was combined with the EHM master mix and gently pipetted until cells were equally distributed through the master mix, ensuring that no bubbles were formed. 450 µl of the resulting EHM mixture was gently pipetted into individual circular recesses of the casting molds (Figure 8A) and placed in a 37 °C humidified incubator with 5% CO<sub>2</sub> for 1 hour. Following solidification of the collagen matrix, 6 ml of pre-warmed SFMM with TGFβ-1 (1 ng/ml) was added to the dishes ensuring complete coverage of each EHM with medium. Fresh SFMM with TGFβ-1 (1 ng/ml) was exchanged at 24 and 72 hours after casting.

Number of EHM	4x	8x
Bovine Collagen Type I (6.6 mg/ml)	283 μΙ	566 µl
2X DMEM	283 µl	566 µl
NaOH 0.1 mol/L	56 µl	112 µl
Cell Suspension	1,479 µl	2,958µl

(1.45 x 10 <sup>6</sup> cells/EHM)		
TOTAL VOLUME	2,106 µl	4,212 µl

**Table 9: EHM reconstitution mixture** 

# 2.5.3 Culturing of EHM

Freshly cast EHMs were incubated in SFMM with TGFβ-1 (5 ng/ml) for the first 3 days of culture in which the EHMs progressively condensed. On culture day 31 EHMs were transferred onto dynamic silicon stretchers (Figure 9A). First, autoclaved stretchers were placed into a tissue culture plate filled with SFMM. Next, to transfer the EHMs from the casting mold onto stretchers the removable silicone tubing was gently removed from the stationary central support pole with a pair of curved forceps. The poles of the stretchers were pinched together with a second set of forceps and the silicon tube was placed over the poles of the stretcher and the EHM was gentle slid off the silicone tube onto the stretcher poles with a pipette tip. The contracting EHM will bend the silicone poles inducing dynamic load for the EHM to work against. Medium was exchanged every other day over the next 25 days of EHM maturation. EHMs were ready for end-point analyses after 28 days in EHM culture (Figure 9B).



Figure 9: Rhesus iPS EHM

After casting of EHMs they are placed on auxotonic silicon stretchers (A). Prior to isometric force measurements EHM are removed from the stretchers and inspected for integrity and spontaneous beating frequency (B). Scale bar: 1mm

## 2.5.4 Dissociation of EHM

Following isometric measurement of EHM functionality they were enzymatically dissociated for FACS analysis. To do so, collagenase (Sigma, 0.2%) was prepared in PBS (with Calcium/Magnesium) with 20% FBS and added to each EHM in a 12 well plate (1 ml/well/EHM) and incubated for 60-90 minutes at 37 °C. After incubation, supernatant from each well was collected in individual PP tubes (15 ml). An Accutase® solution (Millipore) supplemented with 0.025% Trypsin (2.5%, Gibco) and 2% DNase I (Lab stock at 1 mg/ml, Calbiochem; Appendix A1) was added to each well (1 ml/well) to digest any remaining large pieces of EHM and incubated for an additional 30 minutes at room temperature. To stop Accutase® digestion, 5% FBS in PBS (without Calcium/Magnesium; 1 ml/well) was added and gently pipetted 2-3 times. All contents of each well were transferred to the corresponding PP tube containing the collagenase supernatant. The contents of each PP tube were then filtered with a 40 µm cell strainer cell strainer to remove any undigested clumps and leave a single cell suspension. Single cell suspensions obtained from the digested EHM were fixed in ethanol (70% Lab stock) for staining of intracellular markers via flow cytometry.

#### 2.6 Measurements and imaging

#### 2.6.1 Cell count and size measurements

Cell number, size and viability measurements from single cell suspensions were assessed using a CASY Model TT cell counter (Roche), according to the manufacturer's instructions. A Neubauer cell counting chamber was used to determine the number of total viable single cells after enzymatic dissociation.

#### 2.6.2 Isometric contraction measurements

EHM force measurements were performed in temperature and pH regulated organ baths (FMI GMbH) as previously described (Soong et al. 2012, Zimmermann et al. 2000, Zimmermann et al. 2002). Briefly, following 25 days of dynamic loading on sili-

con stretchers, EHMs were suspended between a retaining hook and force transducer in organ baths containing 0.2 mmol/L [Ca<sup>2+</sup>] Tyrode's solution (Figure 10: Table 10). Temperature within the organ bath was maintained at 37° C with an adjusted pH of 7.4 maintained by a constant flow of carbogen (95% O<sub>2</sub>, 5% CO<sub>2</sub> gas mixture). Calcium concentration of Tyrode's solution was adjusted to 1.8 mmol/L using a CaCl<sub>2</sub> solution (0.2 mol/L) diluted from stock solution (2.25 mol/L). EHMs were equilibrated before an electrical field stimulus of 1.5 Hz was applied with monophasic pulse duration of 5 ms and an electrical current of 200 mA. A stepwise preload strain was applied to the EHMs under an extracellular calcium concentration of 1.8 mmol/L until a length of maximal force development (Lmax) was reached in accordance with the Frank-Starling mechanism. Following the preload stretching, 0.2 mmol/L [Ca<sup>2+</sup>] Tyrode's solution was exchanged and EHMs were equilibrated before force measurements were obtained. Contractile force measurements were acquired using PCbased acquisition software (BMON, Engineering firm G. Jaeckel, Hanau) under cumulatively increasing [Ca<sup>2+</sup>] (0.2-4.0 mmol/L). Force of contraction measurements. defined as the difference between maximum (peak systolic force) and minimum (diastolic) forces, were calculated from measurements taken at the steady state, approximately five minutes following addition of calcium. Analysis was performed using PC-based analysis software (AMON, Engineering firm G. Jaeckel, Hanau and Graph Pad Prism). Additional isoprenaline response and force frequency data were collected under half maximum effective concentration (EC<sub>50</sub>) for extracellular calcium (Figures 17 and 18).

# Stock I

Tyrode [Ca <sup>2+</sup> ]	0.2 mmol/L
NaCl (MW=58.44)	175 g
KCI (MW=74.56)	10 g
CaCl <sub>2</sub> stock (2.25 M)	2.22 ml
MgCl <sub>2</sub> stock (1.05 M)	25 ml
adjusted with ddH <sub>2</sub> O	1000 ml

# Stock II

NaHCO <sub>3</sub> (MW=84.01)	50 g
adjusted with ddH₂O	1000 ml

# Stock III

NaH <sub>2</sub> PO <sub>4</sub> (MW=137.99)	5.8 g
adjusted with ddH <sub>2</sub> O	1000 ml

\*All stock solutions can be stored at 4 °C until required.

# Tyrode's Working Solution

Stock I	40 ml	80 ml	120 ml	200 ml
Stock II	38 ml	76 ml	114 ml	140 ml
Stock III	10 ml	20 ml	30 ml	50 ml
Glucose	1 g	2 g	3 g	5 g
Ascorbic Acid	100 mg	200 mg	300 mg	500 mg
Adjusted with ddH <sub>2</sub> O	1000 ml	2000 ml	3000 ml	5000 ml

# Table 10: Tyrode's solutions for isometric force measurements.

\*All constituents were combined in a volumetric flask and thoroughly mixed before use. Working solution should be made fresh no more than 24 hours prior to isometric force measurement.



Figure 10: EHM in organ bath

Isometric force measurements were performed on EHMs suspended between two hooks attached to a force transducer in an insulated organ bath. Organ bath temperature was maintained at 37° C with an adjusted pH of 7.4 maintained by a constant flow of carbogen (95%  $O_2$ , 5%  $CO_2$  gas mixture). EHMs were subjected to cumulatively increasing extracellular calcium concentrations (0.2-4.0), isoprenaline (at  $EC_{50}$  [Ca<sup>2+</sup>], pacing frequencies of 1-4 Hertz in 0.5 Hz increments at 0.8 mmol/L [Ca<sup>2+</sup>], were applied with monophasic pulse duration of 5 ms and electric current of 200 mA.

# 2.6.3 Flow cytometry

Assessment of cardiomyocyte purity prior to EHM construction and following isometric force measurements were performed using flow cytometry with a LSRII cytometer (BD). For data analysis FACSDiva Software 6.0 (BD) was applied. Cell suspensions fixed in 70% EtOH and filtered through a 70 µm cell strainer (BD Biosciences) were

pelleted via centrifugation and permeabilized in blocking buffer (Appendix A5) for 10 minutes at 4  $^{\circ}$ C. Samples were incubated for 45 minutes at 4  $^{\circ}$ C with a primary antibody against  $\alpha$ -sarcomeric actinin (1:1,000; Sigma-Aldrich; Appendix A6) followed by incubation with a goat anti-mouse Alexa 488 secondary antibody in blocking buffer (Invitrogen, A-11001; 1:1,000; Appendix A6) and Hoechst 33342 for 30 minutes at 4  $^{\circ}$ C. A mouse immunoglobulin G (IgG) isotype control (R&D systems) was used as negative control for non-specific staining. Cells were washed (2x) with blocking buffer and resuspended in PBS (without Calcium and Magnesium) for analysis with a BD LSRII (BS Biosystems). Samples were first gated (G1) based on Hoechst 33342 staining of viable cells followed by a secondary gating (G2) to isolate single cell population. Cardiomyocyte purity was based on  $\alpha$ -actinin positive staining cells with an IgG isotype control used to determine non-specific staining (Figure 11). Analysis of samples was performed using Flowing Software 2.5.1.

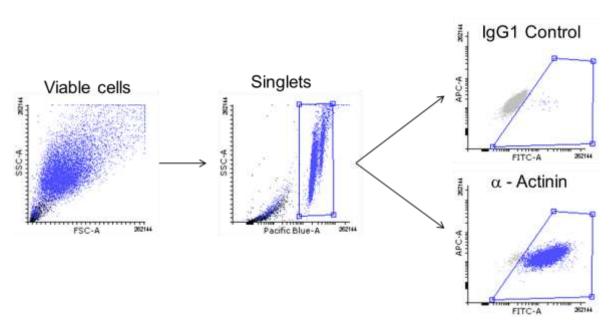


Figure 11: Gating strategy for  $\alpha$ -actinin positive cardiomyocytes.

Viable cells were selected based on Hoechst 33342 intensity. Doublets were excluded based on side scatter area (SSC-A). Cardiomyocytes were distinguished based on  $\alpha$ -actinin positivity. An IgG1 isotype control was used for exclusion of non-specific staining.

# 2.6.4 Immunofluorescence imaging

Sample aliquots of differentiated cardiomyocytes were seeded onto Matrigel<sup>TM</sup> coated coverslips (1:120) in hES medium and incubated at 37 °C for 4-7 days to ensure cell attachment and recovery. When cells began spontaneous contractions via visual inspection coverslips were washed with PBS and fixed with 4% Histofix for 10 minutes at room temperature. After another washing step with PBS, cells were permeabilized by incubation with blocking buffer (Appendix A5) for 30 minutes at RT. Primary antibody (Appendix A6) incubation directed against α-sarcomeric actinin (Sigma-Aldrich), connexin 43 (BD Biosciences), T-box transcription factor 5 (Tbx-5; Sigma-Aldrich), myosin light chain 2a (MLC2a; atrial isoform; Synaptic Systems), and myosin light chain 2v (MLC2v; ventricular isoform; Synaptic Systems) was performed at RT for 60 minutes. Following primary antibody incubation coverslips were gently washed with PBS and incubated with secondary antibodies conjugated with fluorescent dyes (anti-mouse Alexa Fluor® 546, anti-rabbit Alexa Fluor® 488, antimouse Alexa Fluor® 633; Molecular Probes® Appendix A6) and nuclei counterstained with Hoechst 33342 (1:1,000) at RT for 60 minutes. Samples were washed thoroughly with blocking buffer 3 times for 5 minutes with a final wash with only PBS for 5 minutes. Coverslips were mounted to superfrost microscopy slides with Fluoromount-G (Southern Biotech). The slides were air dried and stored in the dark until imaging was performed. Fluorescent images were obtained using a laser scanning confocal microscope (LSM710, Zeiss, Germany).

# 2.6.5 Whole mount imaging

Whole mount immunofluorescence imaging was performed to visualize cardiac muscle formation within EHMs. After cultivation, EHMs were washed once on 1X PBS and fixed in Roti® Histofix 4% overnight at 4 °C. Thereafter, EHMs were washed once in 1X PBS and incubated in a permeabilizing blocking buffer solution for 24 hours at 4 °C. EHMs were subsequently incubated with primary antibodies (Appendix A6) targeting α-actinin (1:1,000) and connexin 43 (1:1,000;) for 48 hours on a rotary microfuge holder at 4 °C. EHMs were then washed in blocking buffer (Appendix A5) for 24 hours on a rotary microfuge holder at 4 °C, followed by the incubation with the goat anti-mouse Alexa 488 secondary antibody (1:400; Appendix A6), a goat anti-rabbit Alexa 633 secondary antibody (1:400; Appendix A6) and a F-actin labelling dye, phalloidin (conjugated with Alexa Fluor® 546, 1:60; Appendix A6) and the

nuclei labelling dye, Hoechst 33342 for a further 48 hours on a rotary microfuge holder at 4 °C in the dark. EHMs were again thoroughly washed in blocking buffer (Appendix A5) for 24 hours and were immediately mounted onto superfrost microscopy slides with Fluoromount-G mounting medium (Southern Biotech) and stored in the dark until microscopic analysis was performed. Whole mount images were obtained using a Zeiss LSM710 confocal microscope.

# 2.7 Statistical analysis

Data were analyzed using Graph Pad Prism 7 software (Graph Pad Software Inc, San Diego) and displayed as mean  $\pm$  standard error of the mean (SEM). n indicates the number of samples. Statistical analyses used were unpaired, two-sided Student's t-test and one-way ANOVA followed by an appropriate post hoc test as indicated in the results section. A P < 0.05 was set to determine significant differences.

# 3. Results

The macaque model has been suggested as clinically predictive for investigating stem cell based therapeutics for the treatment of post myocardial infarction heart failure (Bel et al. 2010, Blin et al. 2010, Shiba et al. 2016). The primary aim of this research was to validate the ability of Rhesus macaque iPS cells to be directly differentiated into cardiomyocytes for use in generating functional EHMs. Two variants, a lentiviral and a Sendai viral transduced line of RhiPS cells, were utilized for cardiomyocyte differentiation and EHM construction.

# 3.1 Encapsulation of Lenti-iPS cells in an alginate:gelatin hydrogel

Our current monolayer differentiation protocol established for human ES and iPS cells has a 4-day period in which stem cells are cultured on Matrigel™ coated plates in E8 medium with 10 µmol/L RI prior to cardiac differentiation (Tiburcy et al. 2017). First attempts to direct the differentiation of the Lenti-iPS cell into cardiomyocytes using a previous version of this protocol that used a 50:50 hES:hFF conditioned medium rather than E8 medium did not yield reliable results. Thus, an alternative culturing strategy with alginate encapsulation to better control the cell growth milieu was investigated. It has been previously shown that encapsulation of stem cells in conjunction with culturing in fibroblast conditioned medium plus RI significantly enhances the viability of the cells (Sidhu et al. 2012). To investigate the effect of culturing and differentiating Lenti-iPS cells in alginate hydrogel beads, variable seeding densities and RI concentrations were tested. Four days prior to the start of directed differentiation, Lenti-iPS cells were dissociated from fibroblast feeder cultures and encapsulated in a 1% alginate:-gelatin hyrdogel at seeding densities of 2 x 106 cells/ml, 1.5 x 10<sup>6</sup> cells/ml, and 1.0 x 10<sup>6</sup> cells/ml as described in section 2.4.2 with either 5 μmol/L RI or 10μmol/L (Figure 12A). Encapsulation of Lenti-iPS cells in 25 μL alginate resulted in beads with a  $3 \pm 0.07$  mm (n = 7) diameter (Figure 12B) and a reproducible growth volume of  $15 \pm 1.2 \mu l$  (n = 7) within the beads (Figure 12C-E).

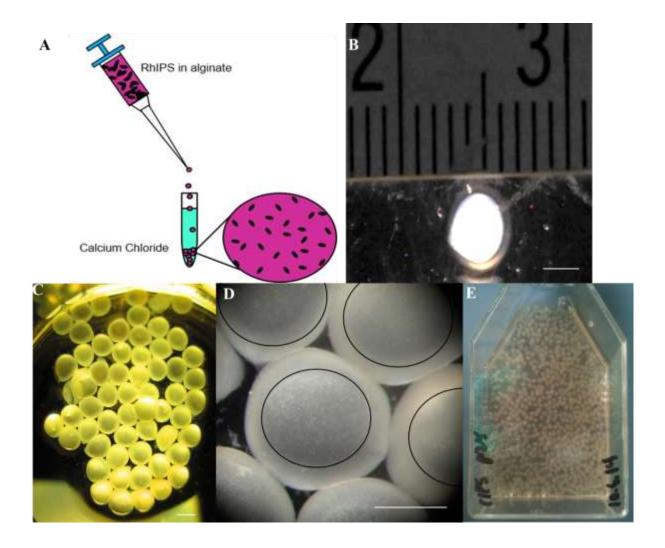


Figure 12: Encapsulation of Lenti-iPS cells:

A) Lenti-iPS cells were mixed with 1% alginate:gelatin and added dropwise (25  $\mu$ I) to a 50 mmol/L CaCl<sub>2</sub> bath in a 50 ml PP tube. Alginate was immediately crosslinked upon contact with calcium cations in the CaCl<sub>2</sub>. B) An alginate bead immediately after 25  $\mu$ I the alginate:cell suspension was dropped into 50 mmol/L of CaCl<sub>2</sub>. C) Consistent size and shape of beads formed by dropwise generation. D) 40X magnification of encapsulated Lenti-iPS cells, black circles signify interface between lique-fied center and crosslinked hydrogel shell, white dots within circles are encapsulated cells/aggregates. E) Over 600 alginate beads in a T175 flask, representing more than 30 x 10<sup>6</sup> RhIPS capable of differentiating into cardiomyocytes. Scale bars: 2 mm

At time points of 24, 48, 72, and 96 hours after encapsulation beads were removed

from culture (n = 5 beads/ time point; in triplicate), dissociated with an alginate dissociation solution (Appendix A9) and counted with a CASY counter to quantify the increase in cell number at each time point. Under treatment with 5  $\mu$ mol/L RI there was a significant increase in cell number after 72 hours in culture (P<0.05 vs. 0 hours post encapsulation) when cells were encapsulated at a concentration of 2 x 10<sup>6</sup> cells/ml (Figure 13A). However, this increase in cell number decreased to below the starting number of cells encapsulated at the 96-hour time point in which differentiation begins under our standard protocol. Culturing of encapsulated cells with 10  $\mu$ mol/L RI resulted in an overall increase in cell number within all three concentration groups at all time points tested. However, the increases in cell number were not significant when compared to the corresponding initial encapsulation concentrations (Figure 13B). Following successful encapsulation in alginate beads, we investigated the ability of Lenti-iPS cells to differentiate into cardiomyocytes.

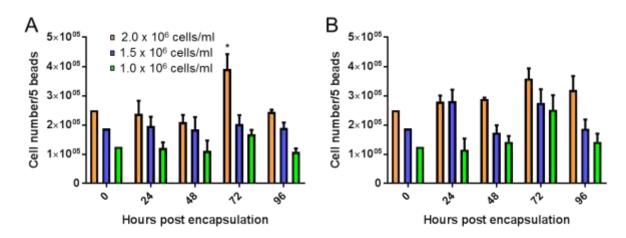


Figure 13: Increase in cell number following encapsulation.

Lenti-iPS cells were encapsulated at the indicated concentrations in alginate. Encapsulated cells were cultured for 96 hours in E8 medium with either 5  $\mu$ mol/L (A) or 10  $\mu$ mol/L (B) ROCK inhibitor. Every 24 hours cell content per bead was analyzed. n = 5 beads/time point, in triplicate; \*P<0.05 vs. 0 hours post encapsulation; 1-way ANOVA for correlated samples with Tukey's HSD test.

# 3.2 Directed differentiation of alginate encapsulated Rhesus Lenti-iPS cells in suspension culture

Following alginate encapsulation, Lenti-iPS cells were cultured for 4-days in 50:50

hES:hEF conditioned medium plus 10 µmol/L RI and 5ng/ml FGF-2, similar to our standard E8 differentiation protocol (Tiburcy et al. 2017). Directed progression into the mesodermal lineage was induced during the first 3 days through incubation with a small molecule Wnt signaling activator CHIR99021 (1 µmol/L), and the growth factors FGF-2 (5 ng/ml), Activin-A (9 ng/ml) and BMP4 (5 ng/ml). Cardiac specification occurred over the next 10 days through the inhibition of the Wnt signaling pathway by incubation with IWP4 (5 µmol/L). Cardiomyocyte selection and enrichment was preformed over the next 4-7 days through replacement of glucose with a lactate selection culture medium (Appendix A3). A post selection recovery period of 7-10 days was provided in which cells were cultured in standard SFBM + FGF-2 (5 ng/mL). On differentation day 28 cardiomyocytes were dissociated and counted. Successful differentiation of encapsulated Lenti-iPS cells resulted in an average cellular output of  $253,776 \pm 36,055$  cells/cm2 (n = 5). This equals a difference of  $203,776 \pm 36,055$ cells/ cm<sup>2</sup> or a  $5.1 \pm 0.7$ -fold increase over original seeding input of 50,000 cells/cm<sup>2</sup>. Next, cells were either with fixed and analyzed by flow cytometry (section 2.6.3) or combined with hFF cells for the generation of EHMs (section 2.5.2). Differentiated Lenti-iPS cell derived cardiomyocytes averaged a purity of 58 ± 7% (n=5) based on flow cytometry analysis of  $\alpha$ -actinin positive staining (Figure 14).

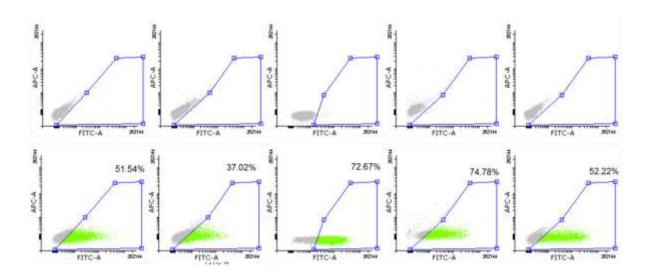


Figure 14: Cardiomyocyte purity following differentiation in alginate beads.

Prior to generation of EHMs, cardiomyocyte purity was assessed using FACS analysis. Cardiomyocytes were identified by a positive labelling for  $\alpha$ -actinin (bottom pan-

els). Isotype controls were included (top panels) to enable gating of the cardiomyocyte fraction. Original recordings from 5 independent experiments.

# 3.3 Directed differentiation of Rhesus Sendai iPS cells in monolayer culture

At differentiation day -4, undifferentiated Sendai iPS cells were plated in monolayers on Matrigel® at an average seeding density of 7.696 ± 1.328 cells/cm<sup>2</sup> (n = 18). A combination of variable culture flask areas (25 - 175 cm<sup>2</sup>) and variable seeding densities (4,000-25,000 cells/cm<sup>2</sup>) were investigated to identify the optimal seeding density. The cardiomyocyte differentiation followed the same protocol as described above for the Lenti-iPS cells (Tiburcy et al. 2017). Mesodermal lineage was induced during the first 3 days through incubation with CHIR99021, FGF-2, Activin-A and BMP4. Cardiac specification occurred over the next 10 days through the inhibition of the Wnt-pathway with IWP4. Cardiomyocyte selection and enrichment was preformed over the next 4-7 days through replacement of glucose with a lactate selection medium. A post selection recovery period of 7-10 days was provided with culturing in standard SFBM + FGF-2 (5 ng/ml). On differentiation day 28 cardiomyocytes were dissociated and counted. Successful differentiation of monolayer Sendai iPS cells resulted in an average cellular output of 64,117  $\pm$  16,789 cells/cm<sup>2</sup> (n = 8). This equals a difference of 54.945  $\pm$  14.635 cells/ cm<sup>2</sup> or a 7.5  $\pm$  1.7-fold increase over an average original seeding input of  $9.173 \pm 2.651$  cells/cm<sup>2</sup> (n = 8). Cells were either with fixed and analyzed by flow cytometry or combined with Rhesus macaque gingival fibroblasts for the generation of EHMs. Differentiated Sendai cardiomyocytes averaged a purity of 91  $\pm$  3% (n=6) based on flow cytometry analysis of  $\alpha$ -actinin positive staining (Figure 15).

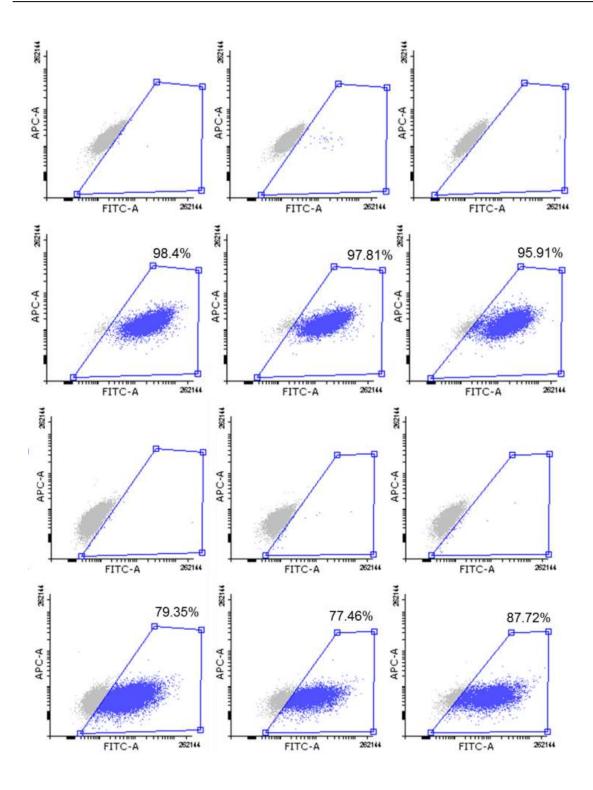


Figure 15: Cardiomyocyte purity following differentiation of Sendai iPS cells in monolayer culture.

Prior to generation of EHMs, cardiomyocyte purity was assessed using FACS analysis in differentiation culture. Cardiomyocytes were identified by a positive labelling for  $\alpha$ -actinin (blue). Isotype controls were included (grey) to enable gating of the cardiomyocyte fraction. Original recordings from the 6 independent experiments.

Fluorescence imaging analysis of Sendai-iPS-cell-derived cardiomyocytes after staining for  $\alpha$ -actinin staining and Tbx-5 demonstrated the presence of canonical cardiomyocyte markers. Additional staining for connexin 43 revealed the formation of gap junctions between cardiomyocytes (Figure 16).

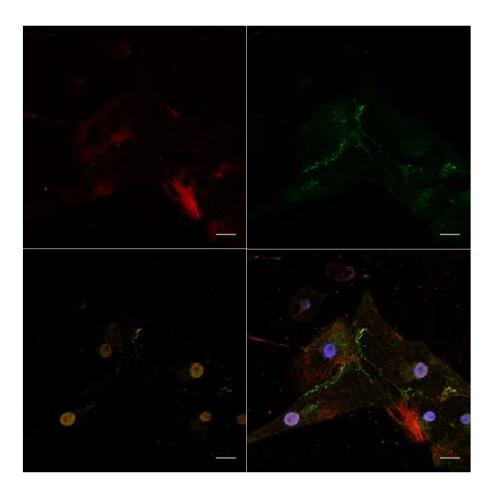


Figure 16: Fluorescence image of RhiPS derived cardiomyocytes

Positive staining of  $\alpha$ -actinin (red) and Tbx-5 (yellow) indicate successful differentiation of RhiPS through mesoderm induction into cardiomyocytes. Positive connexin 43 (green) staining identifies gap junctions between neighboring cardiomyocytes. DNA labelled in blue. Scale bars: 20  $\mu$ m.

# 3.4 Mamu screening

Targeted RNA-sequencing of transcripts encoding for Mamu antigens confirmed the identity of the C1 source fibroblasts and C1 iPS cell culture (Table 11). Sendai iPS cells were, as anticipated, clearly distinct from the C1 line as to Mamu haplotypes.

Mamu expression was similar in Sendai iPS cell-derived cardiomyocytes with and without LPS and INFy stimulation.

	Cell	C1 Fibro- blasts	C1 Lenti iPS cells	Sendai iPSC-CM	Sendai iPSC-CM (LPS)	Sendai iPSC-CM (INF <sub>Y</sub> )
	Origin	DPZ	DPZ	Stanford	Stanford	Stanford
	Mamu-A1	004	004	049	049	049
	Mamu-A2	nd	nd	05	05	05
	Mamu-A4	014	014	nd	nd	nd
	Mamu-AG	nd	nd	03	03	03
Mamu Class I (MHC Class I)	Mamu-B	012 022 030 057 069 082	012 022 031 045 049 057	035 051 063	051 060 063	035 051 060

Table 11: Rhesus macaque MHC Class I (Mamu) antigen screening

RNA sequencing was performed using Rhesus Lenti-iPS origin fibroblasts, Lenti-iPS, Sendai iPS (undifferentiated), and Sendai iPS cardiomyocytes with and without stim-

ulation with "pro-inflammatory cytokines". Sequencing was performed at the TAL-Göttingen. Sequencing analysis performed by Angela Noll (DPZ). Nd: not detected.

# 3.5 Teratoma assay

Teratoma formed in three out of four NOD-SCID mice injected with Sendai RhiPS cells. Two teratoma were removed 35 days post injection and had sizes of 314 mm<sup>3</sup> and 297 mm<sup>3</sup> and weights of 0.53 g and 0.36 g, respectively. The third teratoma was resected after 42 days with a diameter of 236 mm<sup>3</sup> and weight of 0.38 g. The fourth mouse developed no teratoma after 91 days. Teratomas were sectioned and stained with hematoxylin and eosin for embryonic germ layer identification (Figure 17).

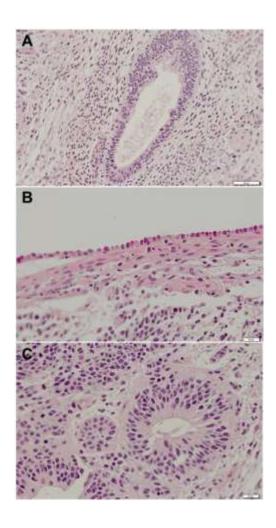


Figure 17: Teratoma assays confirmed pluripotency in Sendai RhiPS cells

H&E staining of teratoma in NOD SCID mice after subcutaneous injection of undifferentiated Sendai RhiPS cells. Structures suggesting endoderm (A), mesoderm (B),

and ectoderm (C) development. Study performed by Prof. R. Dressel. Scale bars: 20 µm

# 3.6 EHM Isometric force measurement

Spontaneous beating of Lenti-iPS-EHMs was detected at the earliest 10 days after EHMs were cast. This corresponds to approximately 7 days following the suspension of the EHMs on the dynamic mechanical stretch devices. EHM contractions were observed until EHM culture day 28. Spontaneous beating of Sendai RhEHMs was detected earliest at 13 days after EHMs were cast. This corresponds to approximately 10 days following the suspension of the EHMs on dynamic mechanical stretch devices. Isometric force measurements revealed positive inotropic responses to increasing concentrations of extracellular calcium with similar forces in Lenti-EHM (n = 23) and Sendai-EHM (n = 27; Figure 18). An apparent calcium sensitivity (EC<sub>50</sub>) was calculated for both iPS cell line at  $0.8 \pm 0.03$  mmol/L [Ca<sup>2+</sup>] (n = 50; Figure 19).

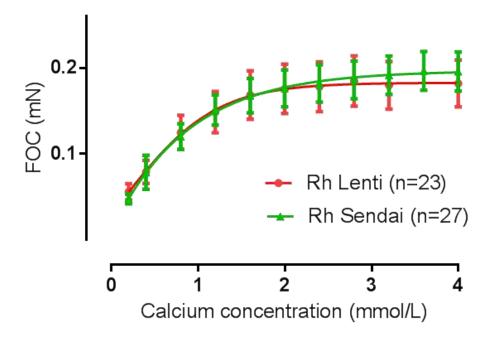


Figure 18: Calcium concentration response in the two tested iPS cell-derived EHM.

Isometric force measurements performed at 37 °C and under electrical field stimulation at 1.5 Hz in EHM generated either from Lenti- (Lentivirus) or Sendai-iPS-

cardiomyocytes. Extracellular calcium was cumulatively increased as indicated on the abscissa. FOC: force of contraction in mN.

Testing response to the classical positive inotrope isoprenaline (1  $\mu$ mol/L) revealed that  $\beta$ -adrenergic responsiveness was similarly developed in either EHM format (Figure 19).

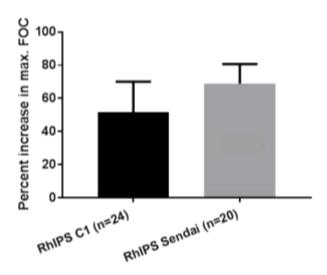


Figure 19: Positive inotropic responses to isoprenaline in Rhesus EHM

Isometric force measurements performed at 37 °C and under electrical field stimulation at 1.5 Hz in EHM generated either from Lenti- or Sendai-iPS cell-derived cardiomyocytes. Extracellular calcium was individually adapted to EC<sub>50</sub> prior to bolus stimulation with isoprenaline (1  $\mu$ mol/L). Positive ionotropic responses were generated in both Lenti EHM and Sendai EHM resulting in an increased FOC of 51 ± 18% (n= 24) and 69 ± 12% (n= 20) respectively. \*P<0.05; one-tailed student T test. FOC: Force of Contraction

A positive force frequency behavior is considered a sign of advanced myocardial maturation. In contrast to human EHM developed under a similar protocol, force frequency behavior appeared negative in Rhesus EHM (Figure 20).

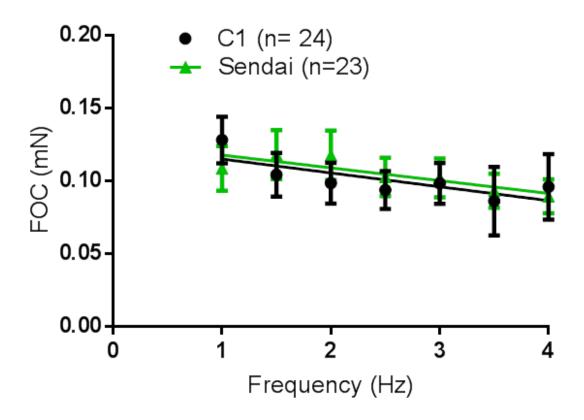


Figure 20: Negative force-frequency response in Rhesus EHM.

Isometric force measurements performed at 37 °C and under electrical field stimulation at the indicated stimulation frequencies in EHM generated either from Lenti- or Sendai-iPS cell-derived cardiomyocytes. Extracellular calcium concentration was maintained at 0.8 mmol/L for the duration of the experiment.

# 3.7 EHM cardiomyocyte content

Following isometric force measurements, EHMs were dissociated and analyzed via flow cytometry for  $\alpha$ -actinin positive staining to identify the EHM cardiomyocyte content. Lenti-EHMs contained 66  $\pm$  1% (n= 3) cardiomyocytes (Figure 21A). Average cardiomyocyte content in Sendai-EHM was 84  $\pm$  2% (n= 3; Figure 21B). Total cell number per EHM was 189,500  $\pm$  12,251 and 995,733  $\pm$  206,107 in Lenti- and Sendai-EHM, respectively.

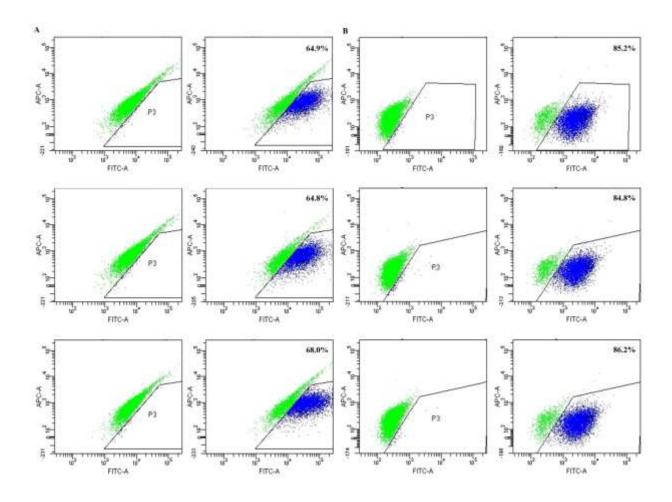


Figure 21: Cardiomyocyte content of Rhesus EHM.

Flow cytometry analysis of  $\alpha$ -actinin cardiomyocytes in EHM. Analyses were performed in EHM subjected to contraction experiments. A) Cardiomyocyte content in Lenti-iPS cell EHMs (n=3). B) Cardiomyocyte content in Sendai-iPS cell EHMs (n=3). Original flow cytometry plots are displayed from all experiments. The cardiomyocyte gate (P3) was set based on the signal obtained from isotype control experiments (green cell population).

Whole mount staining was performed to visualize the structural arrangement of RhiPS-derived cardiomyocytes in EHM (Figure 22). Staining for  $\alpha$ -sarcomeric actinin and f-actin revealed heart muscle network formation.

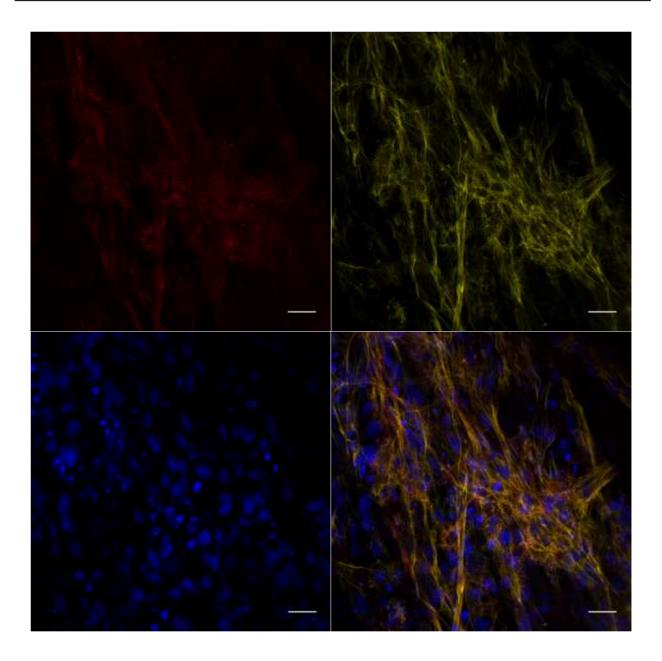


Figure 22: Whole mount staining of RhEHM.

Whole mount EHM with immune labeling of  $\alpha$ -actinin (red) and f-actin (green). DNA (blue). Scale bars: 20  $\mu$ m.

## 3.8 EHM post transplantation gene expression

EHMs generated from Lenti-iPS cell-derived cardiomyocytes were transplanted into non-infarcted nude rats for four weeks (performed by the Wu lab at Stanford, US). Following euthanasia and excision of the hearts no teratoma were observed. Subsequently, RNA was extracted for qPCR analysis for the expression of pluripotency and cardiac marker transcripts. Expression of pluripotency genes POU5F1 and MYC

were found to be 61% and 27% higher than in monolayer differentiated cardiomyocytes from Sendai iPS cells, respectively (Table 12). Cardiomyocyte markers were much less abundant in the Lenti-EHM compared to the Sendai iPS cell-derived cardiomyocyte cultures.

RNA expression ratio					
Lenti-EHM: 2D Sendai cardiomyocyte					
Lenti-EHM Send					
	(explant)	Cardiomyocyte			
Rhesus GAPDH	1	1			
POU5F1	60.97876003	0.999908			
ABCG2	0.990132138	0.999659			
MYC	26.94916477	0.989154			
MYL2	0.004635175	0.884931			
MYH7	0.000123657	0.255618			
MYH6	0.007126534	0.224855			
TNNT2	0.015418664	0.241777			
NKX25	0.008650089	0.962193			

Table 12: Comparative RNA expression of pluripotency and cardiomyocyte marker genes in Lenti-EHM explants and Sendai-cardiomyocytes.

Following four-week implantation in nude rats, a Lenti-EHM (n=1) was excised and a comparison of pluripotency (blue) and cardiomyocytes (red) marker gene expression levels with Sendai iPS cell-derived cardiomyocyte monolayer culture (n=1) was performed via qPCR. Data provided from the Wu laboratory, Stanford University. Data was normalized to GAPDH.

The high level of Rhesus POU5F1 and MYC was taken as a sign of an activation of endogenous stemness markers and thus thought to be suboptimal for further development of the Lenti-EHM approach.

#### 3.9 Superovulation and oocyte retrieval of Rhesus macaque oocytes

As an alternative to iPS cells, parthenogenetic stem cells (pSC) may evolve as a more immune privileged cell source (Didie et al. 2013). To translate earlier work into large animal models, oocytes from five female macaques were collected following three separate hormone-induced superovulation cycles with a one month break inbetween cycles for hormone clearance. Super ovulation was monitored via ultrasound examination to track progression of follicle growth and to identify if a longer cycle of hormone injections was required (Figure 23). Follicle aspiration was employed for ovulation cycle 1 and ovulation cycle 2, whereas complete ovariectomy was performed following ovulation cycle 3 (Figure 24). Ovaries from a sixth female were provided by the DPZ after necropsy. Oocytes retrieved from ovaries of the sixth monkey were subjected to an IVM protocol (Section 2.2.6) in attempt to develop MII oocytes *in vitro*.



Figure 23: Ultrasound examination of hormone stimulated Rhesus ovaries.

Ultrasound examination of Rhesus macaques 2616 (Left), 2551 (middle), and 2500 (right) during super ovulation cycle 3 show stimulated ovaries with multiple follicles having diameter lengths ranging from 0.46 cm to 0.71 cm (optimal follicle diameter is >0.6 cm). Monkeys with ovaries that had smaller and/or fewer follicles received additional hormone injections. L: length in cm; re: right; le: left.

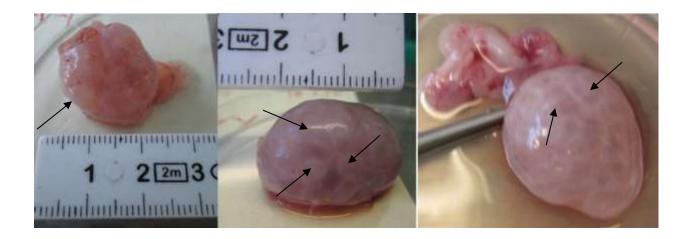


Figure 24: Excised ovaries from super-ovulated Rhesus macaques

Ovaries were excised by bilateral ovariectomy from Rhesus macaques 2551 (Left), 2616 (middle), and 2506 (right) after hormone induced superovulation. Ovary diameters ranged between 2 cm and 3 cm and contained multiple enlarged follicles (arrows).

A total of 224 oocytes were collected from five female Rhesus macaques over the course of three hormone-induced superovulation cycles. An additionally 23 oocytes were collected from a sixth monkey (#2379), not originally included in the study, that was euthanized for medical reasons by the veterinarians at the DPZ. 38 oocytes were harvested from five monkeys after superovulation cycle 1 (8  $\pm$  3/macaque) and a total of 34 for superovulation cycle 2 (7  $\pm$  2/macaque). Harvesting of oocytes was greatly increased in superovulation cycle 3 to a total of 152 (30  $\pm$  8/macaque) due to having the full ovary available for dissection (Figure 25). In the case of the sixth monkey 23 oocytes were collected and subjected to *in vitro* maturation.

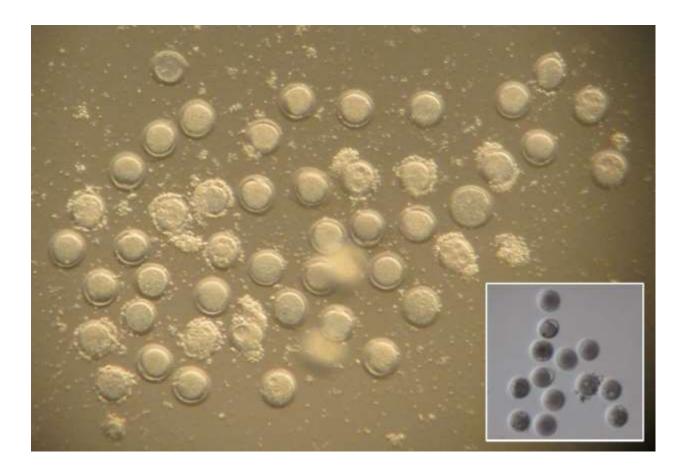


Figure 25: Oocytes collected from excised ovaries

A larger number of oocytes from Rhesus #2506 were collected from whole ovaries following bilateral ovariectomy when compared to number of eggs collected via follicle aspiration (inset).

## 3.10 Activation of retrieved oocytes

Successfully retrieved oocytes from hormone-induced superovulated ovaries were exposed to 0.5 mg/ml hyaluronidase (Type IV; Sigma Aldrich) for 5-10 minutes to remove the cumulus cells. Oocytes were then washed three times with TALP-Hepes medium. Activation of oocytes was performed with exposure to a calcium ionophore (Ionomycin) followed by either 6-DMAP or puromycin. We obtained an overall average activation rate  $27 \pm 0.1\%$  (n = 16) from all oocyte collections (superovulation + IVM). In all ovulation cycles, activation was classified as cells that either divided or fragmented following activation with Ionomycin. Oocytes that remained intact single cells were not considered to be activated.

During normal fertilization, preimplantation embryo development progression consists of an initial division from 1 to 2 cells 24 hours after fertilization. Further cell divisions occur approximately every 12-18 hours thereafter until the morula stage at which point the cells become autonomous in their division cycles (Braude et al. 1988, Chason et al. 2011). We were able to obtain precise timepoints of cellular divisions using the video-optic capabilities of the Embryoscope™. Image analysis indicated that successfully activated Rhesus oocytes followed approximately this same cell division timeline as occurs in normal fertilization (Figure 26).

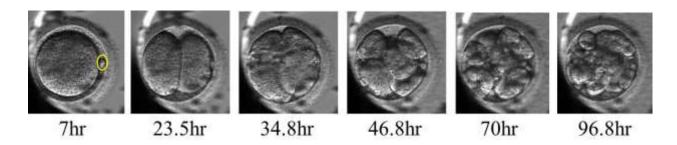


Figure 26: Cellular division of activated Rhesus oocytes

Following successful chemical activation, oocytes begin division after 24 hours and continue dividing approximately every 12-18 hours to the morula stage at which point cells division becomes autonomous. Yellow circle depicts first polar body extrusion signifying an MII oocyte.

Oocytes from ovulation cycle 1 were exposed to 6-DMAP for 4 hours in attempt to block 1<sup>st</sup> polar body (PB) extrusion, which would result in a partially heterozygous pSC (Figure 3). A total of 14 oocytes were activated from the 38 oocytes collected (37%; Table 13). A total of 9 activated oocytes (24% of total) progressed to the morula stage and were transferred onto feeder cultures (Figure 27A).

Rhesus	#2500	#2616	#2483	#2506	#2551	Total
Collected	3	3	7	20	5	38
Activated	1	1	5	7	0	14
Activation rate	33%	33%	71%	35%	0%	37%
2 cell	0	1	4	4	0	9

4 cell	0	1	4	4	0	9
8 cell	0	1	4	4	0	9
Morula	0	1	4	4	0	9
Transferred	0	1	4	4	0	9

Table 13: Oocyte activations of Rhesus macaques from ovulation cycle 1

Oocytes from 5 female macaques were collected via follicle aspiration and activated with ionomycin and 6DMAP treatment. Long term cultures were performed as clusters of oocytes in medium drops in a humidified cell culture incubator. 9 activated oocytes were transferred into feeder layer cultures.

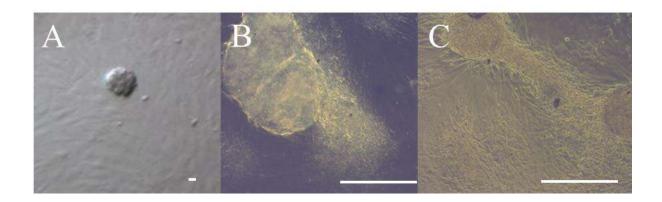


Figure 27: Rhesus ICM plating

Inner cell mass of activated Rhesus oocyte plated onto inactivated human fibroblasts (A). Cellular outgrowths from Rhesus parthenote blastomeres (B and C). Scale:  $50 \, \mu m$ 

Oocytes from ovulation cycle 2, ovulation cycle 3 and IVM cycles were exposed to puromycin for 5 hours in attempt to block 2<sup>nd</sup> PB extrusion which would result in a fully homozygous pSC (Figure 3). A total of 13 oocytes were activated from a total of 38 oocytes collected from ovulation cycle 2 (38%; Table 14). All 13 activated oocytes progressed to the 8-cell stage however, only 11 progressed into the morula stage and were transferred onto feeder cultured (32% of total).

Rhesus	#2500	#2616	#2483	#2506	#2551	Total
Collected	4	14	1	5	10	34
Activated	0	8	0	5	0	13
Activation rate	0%	57%	0%	100%	0%	38%
2 cell	0	8	0	5	0	13
4 cell	0	8	0	5	0	13
8 cell	0	8	0	5	0	13
Morula	0	8	0	3	0	11
Transferred	0	8	0	3	0	11

Table 14: Oocyte activations of Rhesus macaques from ovulation cycle 2

Oocytes from 5 female macaques were collected via follicle aspiration and activated with ionomycin and puromycin treatment. Long term cultures were performed as clusters of oocytes in medium drops in a humidified cell culture incubator. 11 activated oocytes were transferred into feeder layer cultures.

A total of 22 oocytes were activated from a total of 152 oocytes collected from ovulation cycle 3 (14%; Table 15). Only 3 oocytes from 22 activated (2% of total) progressed to morula stage, however, due to the better monitoring capabilities by video-optic recordings of the developmental process via an automated Embryoscope™ 12 activated oocytes at different stages of division were transferred onto feeder layers. Two of the transferred parthenotes from macaques #2551 and #2616 began cellular outgrowths onto the feeder cultures (Figure 27B and C).

Rhesus	#2500	#2616	#2483	#2506	#2551	Total
Collected	27	16	19	58	32	152
Activated	2	8	2	6	4	22
Activation rate	7%	50%	11%	10%	13%	14%
2 cell	2	5	0	5	2	14
4 cell	2	2	0	2	0	6
8 cell	1	2	0	1	0	4

Morula	0	2	0	1	0	3
Transferred	2	7	0	1	2	12

Table 15: Oocyte activations of Rhesus macaques from ovulation cycle 3

Oocytes from 5 female macaques were collected from ovaries following bilateral ovariectomy, activated with ionomycin and puromycin treatment. Long term culture was performed in an Embryoscope<sup>TM</sup> as single cells in a culture well under constant video-optic monitoring. 12 activated oocytes were transferred into feeder layer cultures and 2 blastomeres developed outgrowths.

A total of 2 oocytes were activated from a total of 23 oocytes collected from unstimulated ovaries of a sixth Rhesus (#2379) that was euthanized for medical reasons by the veterinarians at the DPZ. The oocytes underwent 24 hours of *in vitro* maturation prior to parthenogenetic activation. Only 1 activated oocyte was transferred onto feeder layer cultures in attempt to propagate 4-cell outgrowths (Table 16).

Rhesus	#2379
Collected	23
Activated	2
Activation rate	9%
2 cell	2
4 cell	1
8 cell	0
Morula	0
Transferred	1

Table 16: Oocyte activation of Rhesus #2379

Oocytes from Rhesus #2379 were collected from unstimulated ovaries following euthanasia by veterinarians at the DPZ for medical reasons. Oocytes were subjected to *in vitro* maturation prior to parthenogenetic activation. 1 Oocyte was successfully activated and transferred into feeder layer cultures after progressing to the 4-cell stage of development.

The bar graph in Figure 28 summarizes the data from all Rhesus macaque oocyte retrievals and activations.

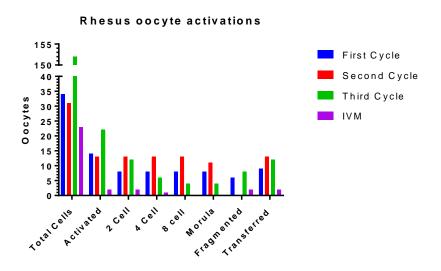


Figure 28: Summary of all Rhesus macaque oocyte activations.

Oocytes were retrieved either by follicle aspiration (cycle 1 and 2) or ovariectomy (cycle 3) after superovulation or necropsy followed by *in vitro* maturation (IVM). Parthenogenetic activation was either by Ionomycin/6-DMAP (1<sup>st</sup> cycle) or Ionomycin/Puromycin (2<sup>nd</sup>, 3<sup>rd</sup> cycles, and IVM).

## 4. Discussion

Heart failure is an increasing social and economic burden across the globe. In recent years therapeutic strategies for the repair of damaged myocardium through tissue replacement have intensified (Feric and Radisic 2016). These investigations have shown promising results in the treatment of heart failure not only in providing paracrine support (Wei et al. 2015, reviewed in Ye et al. 2013) to slow the progression, but also in providing a contractile function to the failing heart (Riegler et al. 2015, Shiba et al. 2012, Weinberger et al. 2016). These studies, it must be noted, were performed in small animal models.

As reviewed in Plews et al (2012) current rodent models are not optimal predictors of efficiency and safety for clinical application, because of their short life span, rapid heart rate and most importantly, the fundamental difference between rodent and human cardiomyocyte morphology and physiology. In order for stem cell based therapeutics to become a viable therapeutic option for the treatment of heart failure, basic scientific findings derived from rodent models must be replicated in a clinically relevant large animal model (Harding et al. 2013, Ogle et al. 2016). Multiple large mammalian species have been used to test the safety and efficacy of stem cell treatments in heart failure caused by myocardial infarction. However, most of these studies utilized xenograft transplantation of human cells into immune suppressed animals such as pig and dog (van der Spoel et al. 2011), sheep (Dayan et al. 2016) and nonhuman primate (Blin et al. 2010, Chong et al. 2014). The lack in pluripotent stem cell models in pigs, sheep, and dogs makes the non-human-primate, with well documented derivation of stem cells (Chen et al. 2008, Debowski et al. 2015, Fang et al. 2014, Liu et al. 2008, Wunderlich et al. 2014, Zhu et al. 2011), and here in particular the macaque model, the preferred large animal for auto- and allograft testing (Shiba et al. 2016). Smaller monkeys such as the common marmoset are, because of bone marrow chimerism in monoplacental species, less suited for transplant studies (Averdam et al. 2007, Genain et al. 1994). Thus, this dissertation focused on Rhesus macaque pluripotent stem cell models, i.e., embryonic stem cells (ESCs), induced pluripotent stem cells (iPSCs), and parthenogenetic stem cells (pSCs). A Rhesus ESC line described in (Thomson et al. 1995) was obtained from WiCell, but could not be maintained in culture. In contrast, iPS cells derived in Göttingen and Stanford

could be cultured and differentiated. First Rhesus pSCs were derived, but not yet further characterized.

The main findings of this dissertation can be summarized as follows:

- Rhesus macaque iPS cells can be directionally differentiated into spontaneously beating cardiomyocytes,
- EHM can be constructed from Rhesus macaque iPS cell derived cardiomyocytes,
- Rhesus macaque EHM have similar properties as human EHM, but appear functionally less mature (lack of a positive force frequency relationship),
- Sendai viral non-integrating reprogramming methods appear to be superior to lentivirally reprogramming of iPS cells in relation to their differentiation capacity and reduced likelihood of oncogene expression,
- Lenti-iPS cells required a preparatory alginate encapsulation approach for successful differentiation into cardiomyocytes,
- Rhesus macaque parthenogenetic stem cells can be derived from unfertilized oocytes after superovulation.

#### 4.1 RhiPS cells can differentiate into spontaneously beating cardiomyocytes

To test the ability of stem cell based therapies in a large animal model to repair damaged myocardium, in which approximately 1 billion cells are lost, a reliable source of replacement cells must be available. Ideally, these cells should be allogeneic to the large animal model in which they will be tested. Previous investigations using allogeneic cells from NHPs include, stem cell derived cell sheets (Bel et al. 2010), cardiovascular progenitor cells (Blin et al. 2010) and iPS derived cardiomyocytes (Kawamura et al. 2016, Shiba et al. 2016). In the study by Shiba et al (2016), iPS

derived cardiomyocytes were shown to have a regenerative function in regards to repairing damaged myocardium similar to other studies using stem cells (reviewed in Der Sarkissian et al. 2017, Hao et al. 2017). Additionally, Kawamura et al (2016) were cable to show that there was a reduced immunological response to homozygous matched transplants. It must be noted that both studies used the Cynomolgus macaque (Macaca fascicularis) ESC and iPS cells respectively.

In this work, we focused on the Rhesus macaque (Macaca mulatta) as our NHP model and were able to successfully differentiate two independent RhiPS cell lines, Lenti- and Sendai-iPS cells, into spontaneously beating cardiomyocytes. Cardiomyocyte differentiation was achieved via small molecule and growth factor directed differentiation that has been proven to be highly effective in human cardiomyocyte differentation(Tiburcy et al. 2017). Cardiomyocyte purity of the RhiPS-derived cardiomyocytes ranged between  $58 \pm 7\%$  for the Lenti- to  $91 \pm 3\%$  for the Sendai-iPS cell lines. Previous protocols for the differentiation of Rhesus ESC into cardiomyocytes utilized the EB format. In this format, between 18% (Schwanke et al. 2006) and 50% (Leschik et al. 2008) of spontaneously beating EBs have been reported. More recently, monolayer differentiation cultures of RhiPS cells have also been shown to produce a cardiomyocyte purity of 92% based on cardiac troponin T positive staining (cTNT) (Zhang et al. 2017). In the studies by Kawamura at al. (2016) and Shiba et al. (2016) cardiomyocyte purities of 80% in EB culture and 84% in monolayer culture were achieved respectively.

Interestingly, human stem cell differentiation protocols in monolayer and EB cultures have been able to achieve purity levels ranging between 30-90% cTNT positive, spontaneously beating cardiomyocytes. (Kadari et al. 2015, Laflamme et al. 2007, Sharma et al. 2015, Tan et al. 2016, Weng et al. 2014, Yang et al. 2008). Additionally, the above studies demonstrate that there is high line-to-line variability when differentiating stem cells into cardiomyocytes. Our Lenti- and Sendai-iPS differentiations provide additional supporting evidence in confirmation of this phenomenon in NHPs. Taken together, our data from Rhesus macaque, provide additional complementary evidence that cardiomyocytes can be efficiently differentiated in high purity from multiple human and non-human primate ES and iPS cells with the caveat that individual

cell lines must be optimized to current differentiation protocols. To further test the applicability for cardiac repair, Rhesus iPS cell derived cardiomyocytes were used in the generation of engineered heart muscle.

#### 4.2 EHM can be constructed from RhiPS cell-derived cardiomyocytes

Current methods of cellular transplantation into infarcted regions of the heart are either by direct injection (intravenous, intracoronary, or intramyocardial) or application of patch like constructs. Incidentally, direct injection methods have been shown have low rates of cellular retention (75-90% lose within 24 hours) due immediate leakage at the injection site or circulatory clearance of non-adherent cells (reviewed in Tian et al. 2015). Conversely, patch like constructs have been shown to increase cell retention and have additional therapeutic benefits (Eschenhagen et al. 2012, Feric and Radisic 2016). These engineered constructs have been shown to stably engraft to recipient hearts, provide paracrine support, and produce an improvement in ventricular function (Didie et al. 2013, Riegler et al. 2015, Weinberger et al. 2016, Wendel et al. 2015, Zhu et al. 2017, Zimmermann et al. 2006). These studies have been conducted with neonatal rat cardiomyocyte allografts, murine parthenogenetic stem cell derived cardiomyocyte allografts or human xenograft cardiomyocytes. Following successful differentiation of RhiPS cells into cardiomyocytes the next step was construction of functional Rhesus EHM for subsequent *in vivo* testing.

Lenti-iPS cell-derived cardiomyocytes were combined with hFF and Sendai-iPS cell-derived cardiomyocytes were combined with Rhesus gingival fibroblasts in a collagen matrix to produce EHMs. After 28 days in culture (25 days under dynamic stretching) the beating syncytia in Lenti- and Sendai-EHM were similar (Lenti: 0.19 ± 0.03 mN, n = 23; Sendai: 0.2 ± 0.02 mN, n = 27). The cross-sectional area of Rhesus EHM was similar to the cross-sectional area previously recorded in human EHM – 0.5 mm². Force of contraction in Rhesus EHM were similar to the forces developed by neonatal human myocardium heart, i.e., 0.3-0.4 mN/mm² (Wiegerinck et al. 2009) and ~5% of the forces recorded in adult human myocardium, i.e., 20 mN/mm² (Mulieri et al. 1992). This will have to be considered and forces further optimized to meet the specific in vivo demands. Here it is also important to note that the macaque heart (40 g) is roughly 10% of the human heart (400 g). Human EHM developed un-

der a similar protocol as the Rhesus EHM in this study developed up to 6 mN/mm<sup>2</sup> (Tiburcy et al. 2017). Additional refinements of the Rhesus protocol as to electromechanical stimulation may assist in achieving these values. For investigations on the smaller Rhesus macaque heart, this may not be necessary if allometric scaling would be acceptable.

### 4.3 Rhesus macaque EHM appear less mature than human EHM

Positive inotropy is the ability to stimulate an increase in the contractility of the myocardium. Pharmaceutical drug treatments for heart failure focus on induction of positive inotropic response in the adult heart to counteract the decrease in myocardial function in a failing heart (Slinker et al. 2000, Tamargo et al. 2009). Identifying the state of functional maturity in stem cell derived cardiomyocytes and their derivatives is investigated through responses to increasing concentrations of extracellular calcium, bolus stimulation with the β-adrenergic agonist isprenaline, and increasing electrical stimulation frequencies (force-frequency behavior). Maximal inotropy of Rhesus EHMs (Lenti- and Sendai) was achieved at 2-4 mmol/L calcium with an EC<sub>50</sub> at 0.8 mmol/L, indicating a maturing of the calcium handling machinery. Positive inotropic effects to isoprenaline suggested the presence of physiologically active β-adrenergic signaling (Tiburcy et al. 2017). Investigations of calcium handling and inotropic responses in early human ESC derived cardiomyocytes (9-19 days after the onset of spontaneous contraction) have shown that immature stem cell derived cardiomyocytes possess proper calcium handling machinery however lack a β-adrenergic response found in mature adult myocardium responsible for contractility stimulation (Pillekamp et al. 2012). The greater increased response in Sendai EHMs may be due to the higher purity of cardiomyocytes used in their production, 91 ± 3% for Sendai EHMs as compared to the 58 ± 7% for Lenti-EHMs. This difference in cardiomyocyte input would result in Sendai EHMs being comprised of more cardiomyocytes that would mature into a more postnatal phenotype and respond to β-adrenergic stimulation. Interestingly, investigation of the force frequency relationship of the EHMs resulted in a lack of positive response from both Lenti- and Sendai-EHMs. This is in contrast to findings in an apparently more mature human EHM model at the same time point of 4 weeks in culture (Tiburcy et al. 2017). To further mature Rhesus EHM it may be imperative to pre-condition Rhesus EHM with electrical pacing during

culture to reach an optimal degree of functional maturation. Adjustment to nearphysiological stimulation frequencies was shown previously to enhance cardiomyocyte maturation in a rat EHM model (Godier-Furnemont et al. 2015).

# 4.4 iPS reprogramming by integrating vectors increases the likelihood of oncogene expression

One of biggest challenges in transplanting pluripotent stem cell based therapeutics is ensuring the safety of not delivering undifferentiated or incompletely differentiated pluripotent cells that retain the potential to form tumors (Forsberg and Hovatta 2012, Hong et al. 2013). Blin et al (2010) have demonstrated that a purified population of SSEA<sup>+</sup> Rhesus cardiac progenitor cells directly injected into an infarcted Rhesus heart can differentiate into ventricular myocytes and reconstituting 20% of an infarct scar. Conversely, they also demonstrated that an unpurified population results in teratoma formation. El Khatib et al (2016) have shown that iPS tumorigenicity is caused by residual pluripotent stem cells and excessive c-MYC expression caused by lentiviral genome integration. The observation of high expression of c-MYC in Lenti-EHM compared to Sendai-iPS monolayer derived cardiomyocytes agrees with this finding.

It is believed that c-MYC regulates expression levels of 15% of all genes (Gearhart et al. 2007). Okita et al (2007) demonstrated that reactivation of the c-MYC transgene was responsible for approximately 20% of tumors developing in chimeras generated from iPS cells. c-MYC has also been shown to be induced in cardiac hypertrophy, where cell size increases not cell number. This effect of c-MYC is in contrast to its regulation of cell number rather than cell size in embryos (Zhong et al. 2006). It is unknown if the high expression of the MYC gene was causing increased iPS proliferation or increased cardiomyocyte size within the Lenti-EHM. Findings from (Martinez-Fernandez et al. 2009) demonstrate that exclusion of c-MYC from iPS reprogramming factors resulted in an increase in cardiogenic lineage fate of differentiated iPS when compared to differentiated iPS cells when c-MYC was included in the reprogramming factors. This data provides support to the speculation that the high expression of c-MYC in Lenti-EHM was causing a cardiogenic hindrance during differentiation as demonstrated by the low purity of α-actinin positive cardiomyocytes.

This would also support the observation of lower cardiomyocyte purities following differentiation when compared to Sendai iPS differentiated cardiomyocytes.

The pluripotency gene POU5F1 was also shown to have increase expression in Lenti-EHMs. The POU5F1 gene encodes for the OCT4 protein involved in the self-renewal, proliferative and pluripotent capabilities of undifferentiated stem cells (Yin et al. 2015). OCT4 works in cooperation with SOX2 and Nanog to maintain the self-renewal and pluripotency capacity in ES and iPS cells (reviewed in Zeineddine et al. 2014). During normal differentiation OCT4 is upregulated and directly involved in mesoderm linage fate and cardiac specification, followed by a diminishing expression in terminally differentiated cardiac cells (Zeineddine et al. 2006). Prior to generation of Lenti-EHM an average of 19.6 ± 5.3% OCT4 positive cells (n = 5; data from A. Krahn, Institute of Pharmacology and Toxicology) were produced by Lenti-iPS differentiations. Since both c-MYC and OCT4 were contained in the STEMCCA integrating reprogramming vector it is highly likely that SOX-2 and NANOG may also show high expression levels in Lenti-EHMs although this was not tested.

Due to the high expression of c-MYC and OCT4 following transplantation in NOD-SCID mice combined with data indicating  $19.6 \pm 5.3\%$  OCT4 positive cells (n = 5; data from A. Krahn, Institute of Pharmacology and Toxicology) remaining following Lenti-iPS differentations it was concluded that Lenti-iPS may be tumorigenic in prolonged implantation studies. Despite not having observed tumors in a 4 week pilot study, it appears plausible to conclude that reprogramming via integrating vectors should not be a preferred method for generating stem cell sources for use in therapeutic applications.

#### 4.5 Encapsulation enhanced directed differentiation in Lenti-iPS cells

Maintaining stem cells in an undifferentiated state requires the *in vitro* creation of the *in vivo* microenvironment that provides growth factors, cell-to-cell interactions and cell matrix adhesions (reveiwed in Moore and Lemischka 2006). This can be supplied through feeder layer cultures or synthetic matrices with growth factor supplementation (Jozefczuk et al. 2012, Lin and Talbot 2011). Conversely, differentiation of stem cells requires the removal of stem cells from this environment of self-renewal

and pluripotency plus the time specific treatment with growth factors and small molecules (Tiburcy et al. 2017).

In our first attempt to differentiate Lenti-iPS cells, removal from the feeder layer culture and cultured in monolayers on Matrigel® resulted in spontaneous differentiation and were not receptive to the recently established and highly effective human stem cell directed differentiation protocols (Riegler et al. 2015, Tiburcy et al. 2017). It is unknown if this was due to the removal of stemness supporting factors excreted by the fibroblast feeder layer, or the reported high biological activity of the Matrigel® (Kleinman and Martin 2005). An alternative strategy was devised that employed the encapsulation of the Lenti-iPS cell in an alginate hydrogel. It has been shown that encapsulation within alginate hydrogels creates a niche environment for stem cells to not only remain in a pluripotent state and propagate, but also differentiate (Dixon et al. 2014, Kryukov et al. 2014, Serra et al. 2011, Sidhu et al. 2012, Siti-Ismail et al. 2008). Our data confirms these findings, as cell number did increase prior to differentiation. Additionally, we have shown that alginate encapsulation enables the successful differentiation of Lenti-iPS cells into spontaneously beating cardiomyocytes, albeit with lower efficiency and purity than their Sendai counterparts.

Coincidently, in a T25 culture flask with a surface area of 25 cm², our current human monolayer differentiation cultures are seeded at a density of 25,000 cell/cm². Encapsulating cells at a concentration of 2.0 x  $10^6$  cells/ml and culturing them in a free-floating suspension resulted in an average of 50,000 cells/bead or 1,250,000 cells in the same culture dish (25 beads/flask). Culture after alginate encapsulation during 4 days of culture before the start of directed differentiation in the presence of  $10 \mu mol/L$  RI 4 resulted in an average cell increase of  $28 \pm 19\%$  (n = 5 beads/time point) in predifferentiation cell number. Furthermore, an additional adjustment to the differentiation protocol to three days of feeder free culture with 5  $\mu$ M RI would result in an increase in cell number by  $57 \pm 20\%$  (n = 5 beads/time point). Despite having a significant increase in cell number at the 72-hour time point after encapsulation using 2 x10^6 cells and 5  $\mu$ mol/IL RI it was determined that fitting the encapsulation procedure to the standard differentiation protocol with  $10\mu$ mol/L RI as well as having a more stable cellular content would be more beneficial for the generation of cardio-

myocytes. Interestingly, directed differentiation was possible after this 4- day encapsulation pre-culture period, suggesting that the microenvironment in the beads rendered the Lenti-iPS cells inducible for directed differentiation. Notably, encapsulation was not required with the Sendai-iPS cells suggesting these cells possess a more stable pluripotent state than in Lenti-iPS cells. The instability of the Lenti-iPS cells may be potentially due to the continuous expression of the genomically integrated pluripotency factors from lentiviral reprogramming which has been previously reported.

#### 4.6 RhpSC can be derived from unfertilized oocytes after superovulation

Cardiomyocytes generated from MHC homozygous iPS cells have previously been shown to have decreased immunogenicity when transplanted into an allogeneic MHC matched Rhesus macaque (Kawamura et al. 2016, Shiba et al. 2016). Prior work from our group has shown that murine pSC derived EHMs transplanted into MHC matched and mismatched mice exhibit immunological acceptance (Didie et al. 2013). We hypothesized that a similar immunological acceptance of MHC haploidentical Rhesus pSC would further reduce the immunogenicity of differentiated cardiomyocytes when transplanted as an EHM. Therefore, generation of pSC lines was attempted with Rhesus monkeys scheduled for future EHM safety studies.

To test this hypothesis pSC lines must first be generated. The Rhesus Mamu homologs to the human leukocyte antigen are present on chromosome 6. Coincidently, these alleles are located on the long arm of the chromosome in Rhesus macaques where there is a higher likelihood of crossing over events occurring during meiosis I as opposed to being located on the short arm of the chromosome in humans. Thus, generating a heterozygous cell line through retention of the second polar body may not be as beneficial in future Rhesus transplantation models as would be in human application. Therefore, attention should be given to the generation of homozygous parthenote cell line.

Generation of pSCs requires the activation of MII oocytes through chemical agents that mimic the fusion of PLC- $\zeta$  with the IP3R. Fusion of PLC- $\zeta$  with the IP3R initiates calcium oscillations in MII oocytes resulting in resumption of MII arrest (Saunders et

al. 2002, Swann and Yu 2008). MII arrest is caused by a cytostatic factor (CSF) regulated by cyclin B1 (Winston 1997). Successful generation of pSC requires the manipulation of the cyclin B1 equilibrium to produce either heterozygous or homozygous pSC.

In our attempts to produce pSC either 6DMAP or puromycin were used following activation by Ionomycin to manipulate cyclin B1 for either extrusion or retention of the 2<sup>nd</sup> PB. Extrusion of the second polar body requires the proper formation of the meiotic spindle. Proper spindle formation requires the degradation of Cyclin B1 causing a decrease in CSF and the resumption of MII. The mode of action of 6DMAP is through the stabilization of cyclin B1 following Ca<sup>2+</sup> activation of the MII oocyte (Bodart et al. 1999) so that it cannot be degraded thus keeping the spindle fibers from properly forming and extruding the second polar body. Therefore, use of 6DMAP will produce a heterozygous pSC due to the retention of the second polar body followed by chromatid replication returning the haploid cell to a diploid state.

Conversely, the use of puromycin after oocyte activation will result in the formation of a homozygous pSC. Puromycin's mode of action is through inhibition of protein synthesis thus increasing the degradation of Cyclin B1 resulting in the decrease of CSF that leads to the cessation of MII arrest. Resumption of MII enables meiotic spindle formation (Homer 2013, Polański et al. 2012) and extrusion of the 2<sup>nd</sup> PB to proceed as would occur during fertilization. Following 2<sup>nd</sup> PB extrusion, spontaneous replication of the single chromatid occurs returning the haploid cell to a stable diploid state (reviewed in Daughtry and Mitalipov 2014).

33 parthenogenetic blastomeres, 9 from 6DMAP treatment and 24 from puromycin treatment, were isolated from activated oocytes and placed into feeder layer cultures. Two of these transferred blastomeres, from the third ovulation cycle, demonstrated cellular outgrowths. Previous derivation of pSC from the Cynomolgus fascicularis monkey by Vrana et al (2003) resulted in one line (Cyno-1) being successfully generated. The Cyno-1 line was the result of 4 ICMs being plated from a total of 28 activated MII occytes obtained from resected ovaries. We could obtain blastomeric outgrowths from 2 separate monkeys obtained from a total of 9 plated blastomeres

out of 22 activated MII oocytes (from 5 monkeys) following ovariectomy. This is in comparison to the 20 plated blastomers from 27 activated MII oocytes following follicle aspiration that gave no outgrowths. These combined data indicate that the most optimal path of pSC line derivation is through isolation of oocytes from whole ovaries.

Incidentally, derivation of a Rhesus macaque haploidentical stem cell line was not completed due to the blastomeres not stabilizing in feeder culture for propagation. This is likely due to one of two factors (or both): (1) the inactivated hFF feeder layer did not properly support the outgrowth of cells despite successful human pSC culture being derived on hFF and/or (2) the culture medium used (Revazova et al. 2007) was not optimal for the propagation of Rhesus pSC as it is for human pSC. We believe that use of inactive mouse embryonic fibroblast feeder cultures will properly support the derivation of the pSC Rhesus macaque and will be utilized in future experiments.

#### 5.0 Summary and outlook

In conclusion, this dissertation has collectively provided evidence that RhiPS cells from integrating and non-integrating reprogramming methods can be successfully differentiated into functional cardiomyocytes. The efficiency and purity of these differentiations is comparable to other non-human and human primate stem cell based cardiomyocyte differentiations. Furthermore, these cardiomyocytes can be used to generate force producing EHM constructs similar to human EHMs generated in the same fashion, albeit with a less mature functional phenotype. Moreover, we generated two independent Rhesus macaque parthenogenetic stem cell lines from unfertilized oocytes. Experience and knowledge gained thus far in the derivation of pSC cells will enable us to establish stable cell lines soon.

We acknowledge that scaling of the differentiation protocol to increase the yield of cardiomyocyte production to reach the billions of cardiomyocytes required for in vivo EHM transplantation studies, is still a work in progress. However, with the data provided by the encapsulation experiments, we envision large scale culture and differentiation are possible in bioreactors. Further critical experiments need to be performed to develop methodologies for increasing the level of physiological maturity of Rhesus EHMs. These findings serve as a stable foundation for the translation of EHM constructs from small animal proof of concept and principle into safety and efficacy investigations utilizing the Rhesus macaque as a surrogate for pre-clinical human studies.

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### **Appendix**

### A1: Reagents

#### Accutase digestion solution

Accutase solution (SCR005; Millipore)
0.125 % Trypsin (1:20 dilution from Trypsin 2.5 % stock [Gibco, 15090-046])
20 μg/ml DNase I

### Activin A stock solution

Dissolve Activin A (338-AC, R&D systems) according to manufactures protocol to obtain a stock concentration of 10  $\mu$ g/ml. Aliquot into sterile 1 ml tubes and store at -20 °C.

#### FGF-2 stock solution

Dissolve FGF-2 (130-093-841; Stemgent) according to manufacturer's protocol to obtain a stock concentration of 10  $\mu$ g/ml. Aliquot into sterile microfuge tubes and store at -20 °C. Aliquot into sterile 1 ml tubes and store at -20 °C.

#### BMP4 stock solution

Dissolve BMP4 (314-BP, R&D systems) according to manufactures protocol to obtain a stock concentration of 10  $\mu$ g/ml. Aliquot into sterile 1 ml tubes and store at -20 °C.

#### CHIR stock solution

Dissolve CHIR (04-0004, Stemgent) in DMSO to obtain a stock concentration of 10 mmol/L. Aliquot into sterile 1 ml tubes and store at -20 °C.

#### IWP4 stock solution

Dissolve IWP4 (04-0036, Stemgent) in DMSO to obtain a stock concentration of 5 mmol/L. Aliquot into sterile 1 ml tubes and store at -20 °C.

#### Collagenase I solution

100 mg of collagenase I dissolved in 40 ml of 1X DPBS (containing  $Ca^{2+}/Mg^{2+}$ ) and 10 ml FBS (10270; Gibco) . Sterile filter (0.22um), aliquot 10 ml into 15 ml tubes and store at -20 °C.

### EDTA Solution (0.5 mmol/L)

Add 500 µl of the 0.5 mol/L EDTA stock solution (pH 8.0), to 500 ml of 1X PBS 14190; Gibco), containing 0.45 g NaCl. Sterile filter (0.22 um), aliquot 10 ml into 15 ml tubes and store at 4 °C up to 6 months.

### Freezing medium

90 % FBS

10 % DMSO (276855; Sigma Aldrich)

### hFF medium

500 ml DMEM (42430; Gibco)

88.8 ml FBS

3 ml P/S

### hES medium

500 ml KO-DMEM(10829-018, Gibco)

128.2 ml KOSR (10828028, Gibco)

6.4 ml L-glutamine

6.4 ml MEM NEAA

Add 10 µl of -2FGF stock solution into 10 ml of hES medium (final FGF-2 concentration: 10 ng/ml).

### TeSR™-E8™ medium

Make up TeSR™-E8 medium (05940; StemCell Technologies) according to manufacturer protocol, including 1% P/S.

### 10X DMEM

669 mg DMEM powder (52100-039; Gibco) dissolved in 5 ml sterile H<sub>2</sub>0.

Filter the solution through a 0.22  $\mu m$  syringe filter. Store at 4 °C for 1-2 months.

## 2X DMEM

1 ml 10X DMEM

120 ml FBS

100 μl L-glutamine

100 μl P/S

Volume adjusted to 5 ml with  $ddH_2O$ .

### A2: Serum free basal media

Serum free basal medium (SFBM)	Volume	Vendor	Order No.
RPMI 1640 plus Glutamax		Gibco	61870-044
B27 plus insulin	2%	Gibco	17504-044
Non-essential amino acids	1%	Gibco	11140-035
Ascorbic acid	200 μmol/L	Sigma	A8960
Penicillin	100 U/ml	Gibco	15140-122
Streptomycin	100 μg/ml	Gibco	15140-122
Supplements	Volume	Vendor	Cat. No.
FGF - 2	5 ng/ml	Stemgent	130-093-841
Activin A	9 ng/ml	R&D Systems	338-AC

BMP4	5 ng/ml	R&D Systems	314-BP
CHIR99021	1 μmol/L	Stemgent	04-0004
IWP4	5 μmol/L	Stemgent	04-0036C

## A3: Lactate selection medium

Lactate selection medium	Concentration	Vendor	Order No.
RPMI w/o glucose w/o glutamine		Gibco	11879020
Sodium lactate	2.2 mmol/L	Sigma-Aldrich	71723
50 mM 2-mercaptoethanol	100 μmol/L	ThermoFisher	31350010
Penicillin	100 U/ml	Gibco	15140-122
Streptomycin	100 μg/ml	Gibco	15140-122

## A4: Serum free maturation medium

	Concentration	Vendor	Order No.
Iscove-DMEM		Gibco	31980-022
B27 minus insulin	4%	Gibco	A418950-01
Non-essential amino acids	1%	Gibco	11140-035
Ascorbic acid	200 µmol/ml	Sigma-Aldrich	A8960
IGF-1	100 ng/ml	Peprotech	AF-100-11

FGF-2	10 ng/ml	Stemgent	130-093-841
VEGF <sub>165</sub>	5 ng/ml	Peprotech	AF-100-20
Penicillin	100 U/ml	Gibco	15140-122
Streptomycin	100 μg/ml	Gibco	15140-122
Supplement	Concentration	Vendor	Order No.
TGF-β	5 ng/ml	Peprotech	AF-100-21C

# A5: Blocking buffer

1X PBS (0.05 M, pH 7.4)	500ml	Gibco	14190-094
Fetal Bovine Serum	25 ml	Gibco	10370-106
Bovine Serum Albumin (BSA)	5 grams	Sigma-Aldrich	A3311
Triton – X 100	2.5 ml	Sigma-Aldrich	T8787

# A6: Immunofluorescence antibodies

Primary Antibody	Dilution	Vendor	Order No./Clone
Anti - α-actinin	1:1000	Sigma-Aldrich	A7811
Anti - MLC2v	1:500	Synaptic Systems	310003
Anti - MLC2a	1:500	Synaptic Systems	56F5
Anti – Tbx-5	1:25	Sigma-Aldrich	HPA008786

Anti - CX43	1:250	BD Biosystems	610062
Secondary Antibody	Dilution	Vendor	Order No./Clone
Goat anti mouse Alexa Fluor 488	1:1000	ThermoFisher	A-11001
Goat anti mouse Alexa Fluor 633	1:1000	ThermoFisher	A-21052
Goat anti rabbit Alexa Fluor 633	1:1000	ThermoFisher	A-21070
Goat anti rabbit Alexa Fluor 546	1:1000	ThermoFisher	A-11010

# A7: Operation anesthesia

### Rhesus 2551

Operation	Weight (kg)	Ketamine (7 mg/kg) (total ml)	Medetomidine (0.04 mg/kg) (total ml)	Metamizole (20-50 mg/kg) (total ml)	Butorphanol (0.01-0.02 mg/kg) (total ml)
Ultrasound	5.7	0.5	0.6		
Follicle aspiration	5.1	0.35	0.2	0.2	0.16
Ovariectomy	5.8	1.32	0.5	0.1	0.5

### Rhesus 2506

Operation	Weight (kg)	Ketamine (7 mg/kg) (total ml)	Medetomidine (0.04 mg/kg) (total ml)	Metamizole (20-50 mg/kg) (total ml)	Butorphanol (0.01-0.02 mg/kg) (total ml)
Ultrasound	7.4	0.6	0.35		
Follicle aspiration	7.3	0.55	0.03	0.2	0.2
Ovariectomy	7.4	0.6	0.35	0.7	0.14

## Rhesus 2500

Operation	Weight (kg)	Ketamine (7 mg/kg) (total ml)	Medetomidine (0.04 mg/kg) (total ml)	Metamizole (20-50 mg/kg) (total ml)	Butorphanol (0.01-0.02 mg/kg) (total ml)
Ultrasound	7.6	0.5	0.3		
Follicle aspi- ration	6.5	0.5	0.3	0.2	
Ovariectomy	7.5	0.7	0.3	0.2	0.1

### Rhesus 2483

Operation	Weight (kg)	Ketamine (7 mg/kg) (total ml)	Medetomidine (0.04 mg/kg) (total ml)	Metamizole (20-50 mg/kg) (total ml)	Butorphanol (0.01-0.02 mg/kg) (total ml)
Follicle aspiration	5.1	0.35	0.2	0.2	0.16
Ovariectomy	4.9	0.65	0.4	0.35	0.1

### Rhesus 2616

Operation	Weight (kg)	Ketamine (7 mg/kg) (total ml)	Medetomidine (0.04 mg/kg) (total ml)	Metamizole (20-50 mg/kg) (total ml)	Butorphanol (0.01-0.02 mg/kg) (total ml)
Ultrasound	6.2	0.65	0.2		
Follicle aspiration	5.2	0.65	0.2	0.2	0.16
Ovariectomy	6.2	0.92	0.25	0.4	0.22

### A8: In vitro maturation medium

	Concentration	Vendor	Order no.
hES medium (Appendix A1)			
Ascorbic Acid	50 μg/ml	Sigma-Aldrich	A8960
Sodium Lactate	10 mmol/L	Sigma-Aldrich	71723
hCG	10 IU/ml	Merck Serono	Ovitrelle; 250mg
hFSH	10 IU/ml	Merck Serono	Gonal F; 75 IU
Sodium pyruvate	0.5 mmol/l	Gibco	11360070
Penicillin	50 U/ml	Gibco	15140-122
Streptomycin	50 μg/ml	Gibco	15140-122

#### **Curriculum vitae**

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- 08. 10. October 2015: **DGK Herbsttagung**, Berlin, Germany, poster presentation: Engineered Heart Muscle from Rhesus Macaque Induced Pluripotent Stem Cell-Derived Cardiomyocytes. 3. Prize Hans-Jürgen-Bretschneider-Posterpreis des Basic Science Meetings.
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