



**Diversity and distribution patterns of foliar fungal
endophytes in *Theobroma cacao* in Central Sulawesi and
interactions between endophytes and host plant**

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Summary

Tropical plants are colonized by a highly diverse community of fungal endophytes. Factors contributing to fungal endophyte diversity pattern and underlying mechanisms are, however, largely unknown. Interactions between host plants and their endophytes are regarded to be highly complex and dynamic. The life cycles of endophytic fungi, their distribution, and diversity are driven by host plant conditions and environmental factors. Some fungal endophytes may be latent pathogens, mutualists or anti-pathogens, depending on their host plants or not yet identified signals inducing switches in their life cycles.

The introduction of cultivated plants to new geographic regions provides an opportunity to compare the fungal endophyte diversity between continents. Cacao (*Theobroma cacao* L.) has been introduced to Sulawesi less than 25 years ago and is therefore reflecting local endophyte communities on a quite recently co-adapted level.

This study aimed at evaluating the changes in fungal endophyte communities in cacao leaves, based on interactions with the host plant and environmental parameters. It provides information about endophyte diversity of the area and therefore about potential outbreaks of latent pathogens in cacao or intercrops, due to environmental conditions.

We investigated the diversity and environmental parameters of the fungal leaf endophyte community of 23 cacao plantations and compared the diversity with existing data from Latin America, the area of origin of cacao.

We found that:

The endophyte diversity of fungi of cacao leaves in Sulawesi is reduced as compared to Latin America.

The composition of the endophyte community within the region is determined mainly by the diversity of shade trees planted in between cacao plants and therefore by the attenuation of radiation provided by the additional canopy.

There is clear evidence for spatial structure of the endophyte diversity, with the similarity of fungal endophyte composition continuously declining between plots with distance.

We analysed the change of fungal endophyte diversity and composition with regard to changing host plant conditions by exposing an artificial drought period in a cacao plantation in Central Sulawesi, Indonesia.

We found that:

After 13 month of throughfall displacement cacao trees exhibited a significant decrease in fungal endophyte diversity.

Given the impact of drought, the endophyte composition within cacao trees became more similar.

The dominance of the species abundance changed, with *Fusarium spp.* becoming the most abundant taxa.

The fungal endophyte community in cacao leaves was sensitive to seasonality.

Furthermore, we analysed the diversity and phylogeny of the various pathogenic *Fusarium* taxa obtained from cacao leaves using molecular methods.

We found

four species known to science (*F. decemcellulare*, *G. fujikuroi*, *F. lateritium*, *F.mangiferae*), having pathogenic potential for cacao or nearby crops (rice, sweet potato, mango)

two unknown *Fusarium spp.*

that all species obtained (except 2) have been also reported from Vanilla, commonly planted in cocoa plantations in Sulawesi.

Fusarium diversity was following a temperature gradient

General Introduction

Fungal life on earth is highly varied. The estimation of fungal diversity was revised when Petrini reported in 1986, that every observed plant exhibited fungal organisms living within plant tissues. Only 74-120 thousand fungal species are known in contrast to at least 1.5 million expected species (Hawksworth 1991, Hawksworth 2001). The undiscovered fungi were assumed to be associated with plants, lichens or insects (Hawksworth & Rosman 1997). In 1995 Willson defined plant associated fungi that occur with complete or partial developmental stages in living tissues, without harming their host plant, such as necrosis or pathogenicity as fungal endophytes. The fungal leaf endophytes were proven to be hyperdivers in tropical trees (Arnold et al. 2000). In leaves of several species a dense patchwork community of abutting fungi was found (Lodge et al. 1996, Gamboa & Bayman 2001, Herre et al. 2007). Additional, host preferences (Arnold et al. 2001) led to the suggestion that a thrust worthy estimation of fungal global species number is relying mainly on the ratio fungal to plant species and plant diversity (Hyde 2001). Conversely, fungal endophytes are able to affect the diversity, structure and dynamics of plant communities (Saikkonen et al. 1998, Clay & Holah 1999). Several studies investigated the various host plant-fungal endophyte interactions and found the endophytes to be mutualists, anti-pathogens, and latent pathogens affecting host plants (Photita et al. 2004, Müller & Kraus 2005, Schulz & Boyle 2005, Kogel et al 2006, Arnold & Engelbrecht 2007, Arnold 2007). An increased knowledge about dynamics in fungal communities and about the interactions with the host plant, could reveal potential biological agents that could stem important pests and diseases in agriculture (e.g. Tong-Kwee et al. 1989, Krauss & Soberanis 2001, Posada & Vega 2005, Rubini et al. 2005, Tondje et al. 2007, Mejia et al 2008, Bailey et al 2008, Zabalgoitia 2008, Vega et al. 2008, Vega et al. 2009a, Ownley et al. 2010). Yet there is only little known about the coherence of endophyte community within a plant, which might complicate the application of fungal bio control agents.

There would be nearly

For example the world's cash crop number one cacao (*Theobroma cacao* L.), an ancient neotropical crop, is grown today in more than 50 countries around the tropics (Lass 2004, Schroth & Harvey 2007). Nevertheless cacao production is threatened by numerous diseases: 1) The "black pod" disease is caused by *Phytophthora palmivora* and *Phytophthora megakarya*. These Oomycetes attacks all parts of the cacao tree, is causing most crucial losses world wide (Appiah et al. 2004a, Appiah et al. 2004b, Guest 2007, Clough et al. unpublished). 2) The vascular streak dieback caused by the highly specialized basidiomycete *Oncobasidium theobromae* led to severe losses in the cacao production in South East Asia,

by wiping entire cacao plantations (Guest & Keane 2007, Ploetz 2007). 3) One of the main cacao diseases in the neotropics is the “witches broom” disease caused by the basidiomycete *Moniliophthora perniciosa*. The infection with the fungus leads to necrosis of plant tissues, after hypertrophy and hyperplasia (Griffith 2004, Aime & Phillips-Mora 2005). 4) Another threat to cacao production in South America is the “frosty pod” disease by the agent *Moniliophthora roreri*, which infects and destroys cacao pods (Griffith 2004). 5) Insect pests such as mirids (e.g. *Sahlbergella* spp., *Helopeltis* spp., *Monalonain* spp., *Distantiella theobroma*) or the cocoa pod borer (*Conopomorpha cramerella*) are causing further serious yield losses by damaging ripe cacao pods (Ploetz 2007, IOOC Database). Beside the direct damage, the insects act as vectors for fungal diseases and enhance fungal infestation rates by additional infections via lacerations of plants epidermis (Williams 1953, Bisseleua 2007). 6) If pathogens with minor potential are introduced in the same cacao plants, pathogens might interact and enhance their pathogenicity. If for example *Fusarium decemcellulare*, the agent responsible for cushion galls and *Lasiodiplodia theobromae* co-occur in cacao plants, they will additionally inflict stem cancer or lead to diebacks (William 1953, Bisseleua 2007, Adu-Acheampong 2009).

The knowledge about the endophyte communities of cacao plantations in Africa and Asia is very limited (but see Crozier et al. 2006). While professional cacao production was established in Africa during the late 19th century (Edwin & Masters 2005, Laird et al. 2007, Sonwa et al. 2007), the main cacao production in Asia started with the cacao boom 25 years ago (Clough et al. 2009). The introduction of cultivated plants to new geographic regions provides an opportunity to compare the fungal endophyte diversity between continents. The endophyte community of a host plant is known to change, when introduced in a new region with a different climate (Hoffman & Arnold 2008). Additionally it was shown, that endophyte composition changes along precipitation gradients (Suryanarayanan et al. 2002) and that even in desert areas, adapted plant species with rich endophyte communities can be found (e.g. El-Zayat 2008, Porras-Alfaro 2008). Though the impact of gradually- as well as suddenly occurring changes in microclimate are yet unknown.

The global climate change influences the intensity and frequency of the El Niño-Southern Oscillation (ENSO) phenomenon in the tropics (IPCC 2001, Walter et al. 2002, Thomas et al. 2004, Parmesan 2006). South East Asia suffered in recent ENSO phenomena severe droughts periods (Quinn et al. 1978, Kerr 1998, Timmermann et al. 1999). If latest climate-prediction-models prove to be true, Indonesia will be threatened in future more frequent by severe drought events (Sheffield & Wood 2008). The impact of long term drought on the physiology of evergreen tropical tree species used to stable climatic conditions is widely unclear. The cacao production of Central Sulawesi decreased by 38% associated with the

recent ENSO related drought events (Keil et al. 2008), suggesting that cacao trees are susceptible to drought stress. The changes in plant physiology of an affected tree may also affect the fungal endophytes within the climatic sensitive leaves. All changes in endophyte community, might affect the interactions between endophyte community and host plants. On one hand interaction between endophytes and host plant might get out of balance and latent pathogens might break out causing diseases (Schulz et al. 1999). On the other hand there are some endophytes are known to induce some drought resistance by stimulating drought stress related genes (Sherameti et al 2008), or regulating photosynthesis rates (Bacon 1993, Arnold & Engelbrecht 2007). The reactions of foliar fungal endophytes community to drought stressed cacao as host plants and consequential changes in host plants physiology are largely unknown. Investigations of the endophytic fungal community and their physiology in cacao plant are necessary for diseases predictions, bio control agent monitoring, and as basic knowledge for a successfull and sustainable establishment of bio control agents.

The aim of this investigation was to increase the knowledge about the diversity and distribution patterns of fungal endophytes in cacao agricultural management. Additionally the impact of drastic climatic change on the foliar fungal endophyte composition of one cacao plantation was investigated, with respect to fungi-host plant interactions.

Objectives

The aim of this work was to investigate the foliar fungal endophyte composition of cacao (*Theobroma cacao*) in Central Sulawesi, Indonesia. The focus laid on diversity, distribution patterns, interactions with the host plant in agricultural management and impact of Climate Change.

The diversity of leaf-inhabiting fungal endophytes of 23 cacao plantations was investigated and compared to existing data from South America, the area of origin of cacao. The fungal endophyte composition was investigated for environmental parameters.

Changes of fungal endophyte diversity and composition with regard to changing host plant conditions by exposing an artificial drought period in a cacao plantation were analysed to understand host plant – endophyte interactions under extreme conditions.

Diversity and phylogeny of the various pathogenic genus *Fusarium* obtained from cacao plantations in Central Sulawesi has been investigated, using the LSU of the ribosomal DNA and EF-1 α .

Chapter 1:

The Lost Endophytes: Endophytic Fungal Diversity of a Neotropical Tree (*Theobroma cacao*) in Central Sulawesi, Indonesia

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Abstract

Patterns of fungal endophyte diversity in trees and underlying mechanisms are largely unknown. Introductions of cultivated plants to new regions provide an opportunity to compare endophyte diversity between continents. This study investigated diversity and environmental drivers of leaf endophyte community of *Theobroma cacao* L. in 23 plantations in Central Sulawesi, Indonesia. With 149 morphospecies and 32 distinguished taxa the diversity is reduced compared to South America, where cacao originates. The differences in endophyte community suggest that host preferences of local endophytes cause this pattern. The composition of the endophyte community within the region was determined mainly by the diversity of shade trees planted in between cacao plants and, to some extent, by the attenuation of radiation by additional canopy. In addition, there was clear evidence for spatial structure, with the similarity of fungal endophyte composition between plots continuously declining with distance.

Keywords: cocoa/ diversity/ endophytes/ similarity/ species richness/

Subject Category: Microbial ecology and functional diversity of natural habitats

Introduction

Tropical plants are colonised by a highly diverse community of fungal endophytes, as has been shown for several tree species (Brown et al. 1998, Arnold et al. 2000, Arnold et al. 2001, Hyde 2001, Suryanarayanan et al. 2002, U'ren et al. 2007). The ecological roles of fungal endophytes are diverse and variable (Saikkonen et al. 1998): Fungal endophytes may occur as pathogens or mutualists, inducing systemic resistances or providing their hosts with antipathogenic compounds depending on the host plant or releasing unknown signals to switch life cycles (Müller & Kraus 2005, Schulz & Boyle 2005, Kogel et al. 2006, Arnold 2007, Zabalgoitia 2008). Given the ecological importance of endophytes, disentangling the drivers of endophyte community diversity and composition is of major importance (Saikkonen et al. 1998, Arnold 2007).

The agricultural history of cacao reaches back to the Mayas, which contributed to the widespread use of cacao in South America (Hurst et al. 2002, Motamayor et al. 2002). Recently, molecular investigations using microsatellites identified the upper Amazonian region in Brazil to be the geographic origin of cacao (Serenio et al. 2006, Zhang et al. 2009). In the early 19th century cacao (namely the Amelonado cultivar) was introduced for the first time to Africa (Edwin & Masters 2005). In 1886 the Trinitario variety was introduced to this region and became the dominant cultivar because of its productivity as well as its resistance against plant diseases (Laird et al. 2007, Sonwa et al. 2007). Today cacao is grown in more than 50 countries around the tropics, including several African and Asian countries (Lass 2004, Schroth & Harvey 2007). Compared to the long cacao growing history in South America and Africa, Asian cacao production is very young.

Several surveys confirmed a high diversity of fungal endophytes associated with cacao (*Theobroma cacao* L.) and related plant species in South America (Arnold et al. 2003, Evans et al. 2003, Rubini et al. 2005, Thomas et al. 2008). However, studies on the endophytes of cacao outside the neotropics are rare (Crozier et al. 2006).

So far there have been no comparative studies on the characteristics of fungal endophyte diversity in cacao grown in Asia compared to South America. In this study we therefore investigated the diversity of fungal endophytes in leaves of cacao trees from Central Sulawesi in Indonesia, where cacao was introduced less than 25 years ago. We deliberately used the same methodological approach for the assessment of endophyte diversity in Central Sulawesi as in published studies from Panama (Arnold et al. 2001, Arnold et al. 2003), aiming at comparing the intercontinental endophyte diversity with the same methodological approach. We are aware of drawbacks inherent to this approach; a more timely approach would have been assessing the diversity of endophytes by molecular

methods. However, with regard to our hypotheses proposed below comparisons would have been impossible when using these more recent methods as compared to the isolation methods used in these previous studies (Arnold et al. 2000, Arnold et al. 2003).

Plants introduced outside of their native ranges typically host a reduced plant pathogen diversity (Parker & Gilbert 2004), which needs decades to re-establish (Mitchell et al. 2010). Based on this well established evidence for plant pathogens we hypothesized that several fungal endophyte species should have been lost by the introduction of cacao in Sulawesi, and, due to the short history of cacao growth in this region, should be either permanently missing from the species pool or should not yet have re-established in this plant species. Therefore we expected the diversity of endophytic fungi in Sulawesi to be less species rich compared to Panama. Furthermore we hypothesized that the cacao plants in Sulawesi are mostly colonized by unspecific local endophytes, recruited on the base of environmental factors.

Material and Methods

Area of investigation

The plantations were selected along the Kulawi valley, located at the western border of the Lore Lindu National Park in Central Sulawesi (Fig. 1). The humidity in the investigation area is between 77% and 85% while minimum temperature ranges between 12 and 17°C and maximum temperatures between 26 and 35°C. Rainfall is very variable and may range between 84 and 2110mm per month (The Nature Conservancy, 2004). During data collection in March 2007, the mean temperature in the Kulawi valley ranged from 22-25°C depending on the plantation elevation (Tab. 1). The precipitation in March reached 242-250mm and the last dry month before was in October 2006 when the rainfall was below 46mm. The wind was blowing mainly northwards (down the valley) with mean wind speed of 1.13m/s. (Kreilein, pers. communication). The 23 investigated plantations were chosen within 40km along environmental gradients of shading, shade tree composition and distance to natural rainforest. In each plantation one 40 x 40m investigation plot was established. The distances between plots were greater than 850m, while the distances between plots and natural rainforest ranged between 10m and 2500m. Cacao trees in the investigation plots originated from nurseries and are grown under overstory canopy of natural forest, or planted shade trees. These shade trees originated from cuttings. Most common planted shade tree species are *Gliricidia sepium* (Jacq.) Walp. and *Erythrina subumbrans* Merr.. Furthermore fruit trees were planted to increase the shade, like candlenut (*Aleurites moluccana* (L.) Willd.), rambutan (*Nephelium lappaceum* L.), avocado (*Persea americana* Mill.), langsung, (*Lansium domesticum* Correa) and durian (*Durio zibethinus* Merr.). Species that are both planted and naturally occurring are sugar palm (*Arenga pinnata* (Wurmb) Merr.) and sago palm

(*Metroxylon sagu* Rottb.). In plantations formerly covered with rainforest *Ficus* sp., *Pterospermum celebicum* Miq. and *Bischofia javanica* Blume commonly remain in the cacao plantations as shade trees. In addition to the shade trees, other crops, such as coffee (*Coffea* sp.), or chili (*Capsicum annuum* L.) are grown within the cacao plots. All together 150 tree species were recorded in the cacao plantations, with a maximum diversity of 20 species in one plot (Clough et al. 2009). The density of tall shade trees ($h > 15\text{m}$) ranged from two to 42 individuals per plot. (Tab. 1). Because of the different number of shade trees, the investigated plantations differed in solar exposition. The openness as an index for radiation was measured using a digital camera system with a calibrated fish-eye lens converter (WINSCANOPY Basic Mini) in a self-leveling mount with remote control at noon on days with clear sky (Propastin & Erasmi 2010). Data were recorded in the centre as well as in all four corners of each plot and subsequently averaged across the plot using the software CanEye. The canopy openness was measured over the cacao canopy in a height of 5.80m as well as under the cacao canopy in 1.30m height. Values ranged from 21.79 to 76.65% in 5.80m and 8.05 to 20.20% in 1.30m height. Under the cacao in the herbaceous layer a grid (5x5m) was installed and weed diversity was recorded twice a year. The diversity of weeds ranged from two to 19 species (Cicuzza et al. 2010). According to the farmers, no fungicides had ever been applied to the cacao trees in the experimental plots.

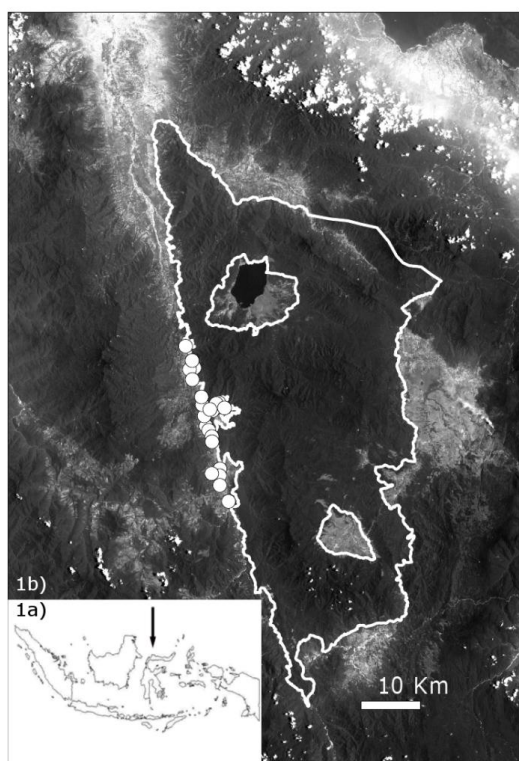


Figure 1 a) Study area in Sulawesi (Indonesia) marked with an arrow; b) Lore Lindu National Park (border marked with white line) and sampled cocoa plantations (white dots).

Collection and isolation of endophytes

The sampling of foliar endophytes on cacao trees was carried out from March 10th to 15th, 2007. 544 mature and healthy cacao leaves were collected. From each tree (*Theobroma*

cacao L.) four leaves from separate branches of two different canopy layers were sampled. Because foliar endophyte diversity is known to increase with exposition of the leaves to the environment (Arnold & Herre 2003), only mature leaves were harvested. Immediately after sampling, the leaves were individually enclosed in sterile polyethene bags and transported to the laboratory of the University of Tadulako (Central Sulawesi), where they were stored at 8°C and processed within the following two days. For surface sterilization the whole leaves were bleached in 3% NaOCl for 3 minutes, afterwards washed in 70% ethanol for 3 minutes and then rinsed in distilled water for 3 minutes. To obtain a representative diversity of endophytes per leaf, five leaf discs were cut out from each leaf, using a flame sterilized circle-cutter (area= 78.5mm²). While one leaf disc was cut of midvein on the leaf tip, four further discs were cut aside the midvein of the leaf. Distance between leaf discs was not less than 3cm. To control for a successful surface sterilization process the leaf surfaces were pressed on Malt Extract Agar (MEA. (Roth, Karlsruhe, Germany) and, after incubation for seven days, Petri dishes were examined for fungal infections. The five leaf discs of one leaf were placed on antibiotic (streptomycin 600ppm) 2.5% MEA in a 9cm Petri Dish sealed with Parafilm and incubated at 25°C. All fungi that grew out from each leaf disc were isolated, purified and grouped to morphospecies (hereafter called morphospecies) based on cultural characteristics, using the following parameters: colony surface textures, hyphal pigments, colours of exudates, and growth rates (Brown et al. 1998, Arnold et al. 2000, Suryanarayanan & Vijaykrishna 2001). We excluded isolates of *Aspergillus*, *Penicillium* and *Mucor* species following the suggestion of Hyde & Soyong (2008), as we expected them to be contaminants, invading through lacerations of parafilm during isolate shipment to Germany. For descriptive identifications purified fungi were placed on a thin layer of nutrient less MEA and stored in a UV-Chamber at 20°C with a light regiment of 12h light: 12h darkness. MEA was used because it is regarded to encourage higher sporulation in many genera of fungi (Brown et al. 1998). Following one month of UV radiation, fungal isolates were characterized based on their spores (hereafter called fungal taxa). Because several of the fungal endophytes remained sterile even under UV- and nutrient stress conditions, we classified those cultures who failed to sporulate using characteristics of the mycelium. Only morphospecies data were used for the statistical analyses. *Nodulisporium* species are considered anamorphs of some *Xylaria* species and form different morphospecies. As this study has been designed to closely mimic the sampling and data analyses methods of a former study in Panama (Arnold et al. 2000), pleomorphs were not considered.

Statistical Analysis

We analyzed leaf-level presence/absence data of the fungal endophyte morphospecies. Because four leaves were collected from each tree, each tree could reach a maximum frequency of four, while each site could reach a maximum frequency of 16. Based on the

results of Herre et al. (2007), who report an endophyte species density of one fungal endophyte each 2mm² leaf fragment on *Theobroma cacao* L., we regard it valid to treat foliar fungal endophyte samples separated by more than 2mm as abundances. Dominances (D) were calculated and categorized in dominance classes (Engelmann 1978). Furthermore alpha diversity indices (Fishers- α and Shannon index) as well as gamma diversity were calculated to assess the species richness of the area. This allows comparative studies of endophyte communities regardless of hosts and country. In proportion to the mean leaf area (34.3m²± 14.2) (Köhler et al. 2009) of one cacao tree in the investigation area, the 20 per tree sampled leaf discs (1.5 *10³ m²) were randomly selective to the inhabiting endophyte community. Therefore we also calculated the Simpson Index (SI) which takes into account that species are randomly sampled from a population (Simpson 1949).

The similarity in endophyte composition between plots was calculated to test for spatial structure and environmental drivers. Jaccard- (JI) and Soerensen (Sol) indices, which are based on presence/absence data were calculated, as they have previously been used in fungal endophyte ecology (Arnold et al. 2000). Abundance-based Morisita-Horn (MH. and Bray-Curtis dissimilarity (BC) indices were also computed (Arnold et al. 2001, Arnold et al. 2003, Vega et al. 2009). A matrix of between-plantation distance classes (0 to 28.000m, 1000m steps) was generated. A linear regression was used to relate distance with similarity in endophyte composition. For the regression and significance tests the program Statistica (version 2.0) was used. Diversity and shared species analysis was calculated using Estimate S (version 8.2) with 1000 randomizations (Colwell 2008).

Results

Though all 544 leaves (2720 leaf discs) contained fungi (data not shown), fungi considered to be endophytes could not be isolated from all leaves (Tab. 1). Altogether 2885 fungal isolates were cultured, from which 32 fungal taxa were distinguished based on spore and morphological characters (Tab. 2). Based on the morphospecies concept, 149 morphospecies were obtained representing a Fishers- α of 41.34± 2.78 as well as an exponential Shannon Index of 78.35± 2.38 over all plantations in the Kulawi valley. Species richness ranged from seven to 64 distinguished morphospecies per plot, with a mean value of 44.47± 13.31 (Tab. 1). Environmental factors showed no significant influence on species richness. The likelihood of two randomly sampled endophytes in this valley belonging to the same morphospecies was 52.66± 3.17% (SI. (Simpson 1949).

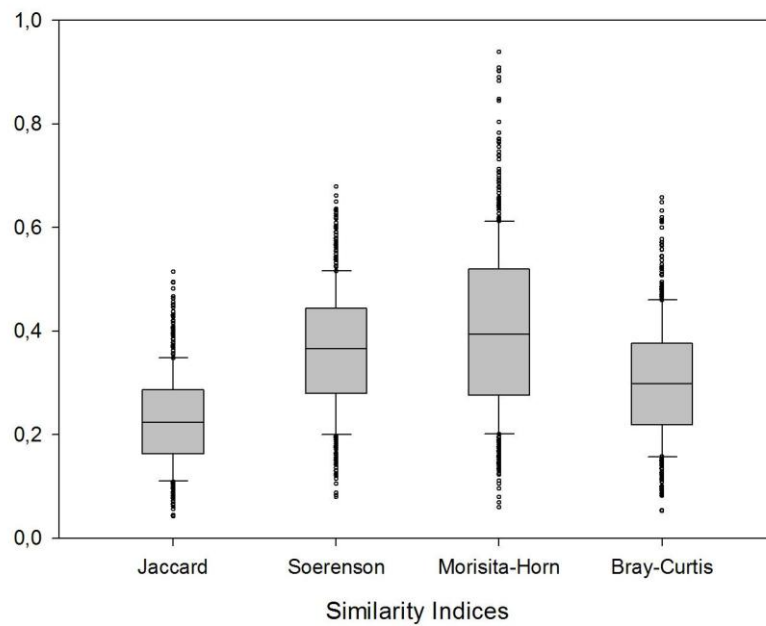


Figure 2 Indices of beta diversity of fungal foliar endophyte composition based on morphospecies for all cocoa plantations in Kulawi valley, Central Sulawesi.

Accumulation Curve

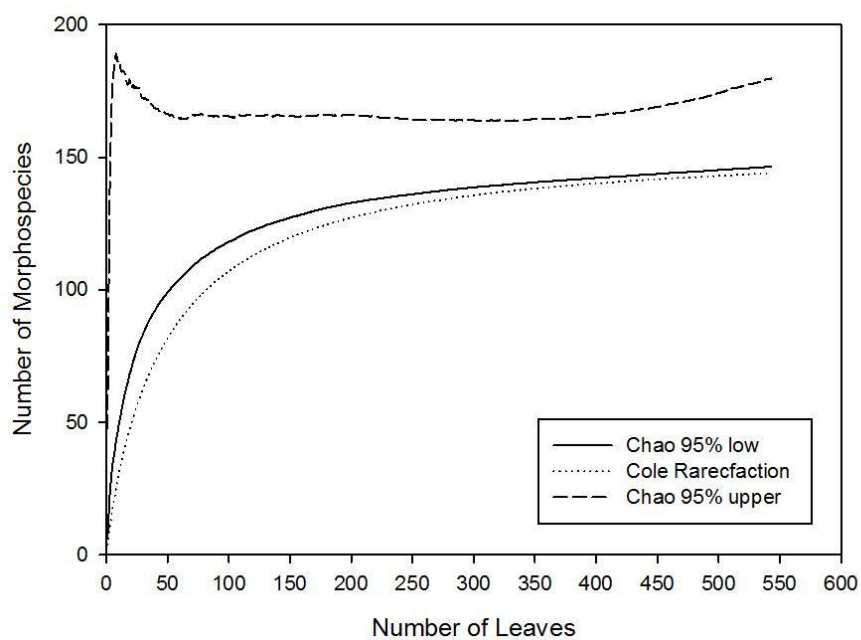


Figure 3 Rarefaction curve for expected and estimated number of foliar fungal endophytes per cocoa leaf in central Sulawesi.

The rarefaction curve showed an initially strong rate of increase of accumulated morphospecies, which declined slightly after 45 sampled leaves. After 200 collected leaves the majority of the fungal endophytes diversity was obtained (Fig. 3). With regard to spore characterisations, the endophyte composition was dominated by the genera *Nodulisporium* with 303 samples (D =10.5%) besides subdominant occurring genera of *Fusarium* with 280 samples (D= 9.7%) and *Xylariaceae* with 214 samples (D=7.4%) (Tab. 2). Medium similarity of the endophyte morpho-species composition in all investigated plantations of the Kulawi valley was 0.23 ± 0.09 (JI + standard deviation SD. 0.37 ± 0.12 (SI+ SD) 0.40 ± 0.16 (MH+SD) 0.30 ± 0.12 (BC+SD) (Fig. 2).

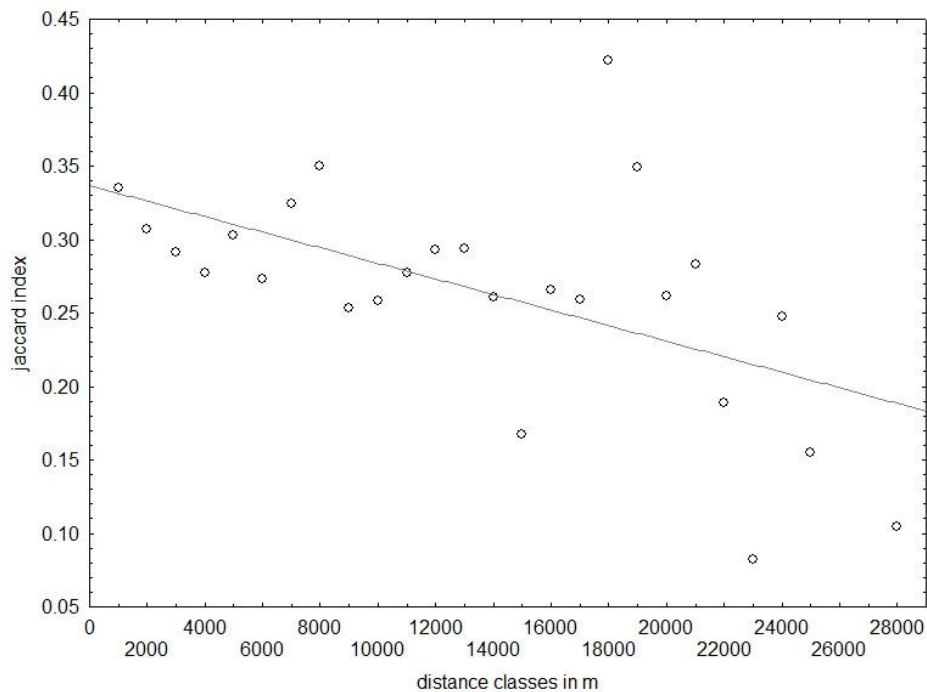


Figure 4 Jaccard index for fungal endophyte composition in leaves of cocoa plantations for all possible site pairs, grouped by distance: $y = 0.34 - 5.30E-6 \cdot x$; $r = -0.55$; $p = 0.004$; $r^2 = 0.30$ (for explanation of distance grouping calculations see text).

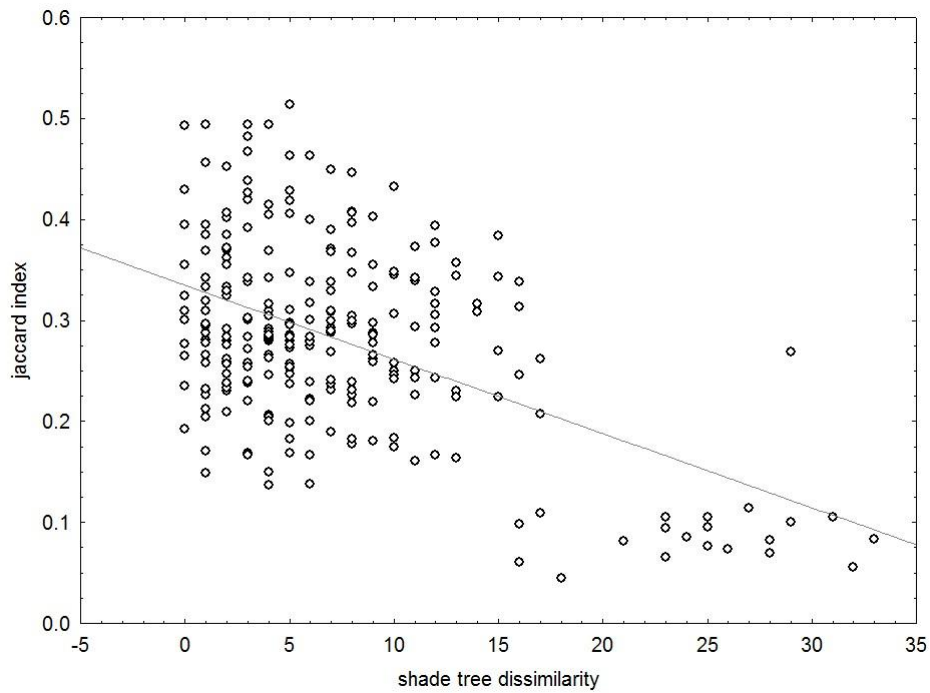


Figure 5 Correlation between similarity (Jaccard index) of foliar fungal endophyte composition and differences in species richness of shade trees for all possible site pairs. $y = 0.34 - 0.01 \cdot x$; $r = 0.52$, $p < 0.000$; $r^2 = 0.27$.

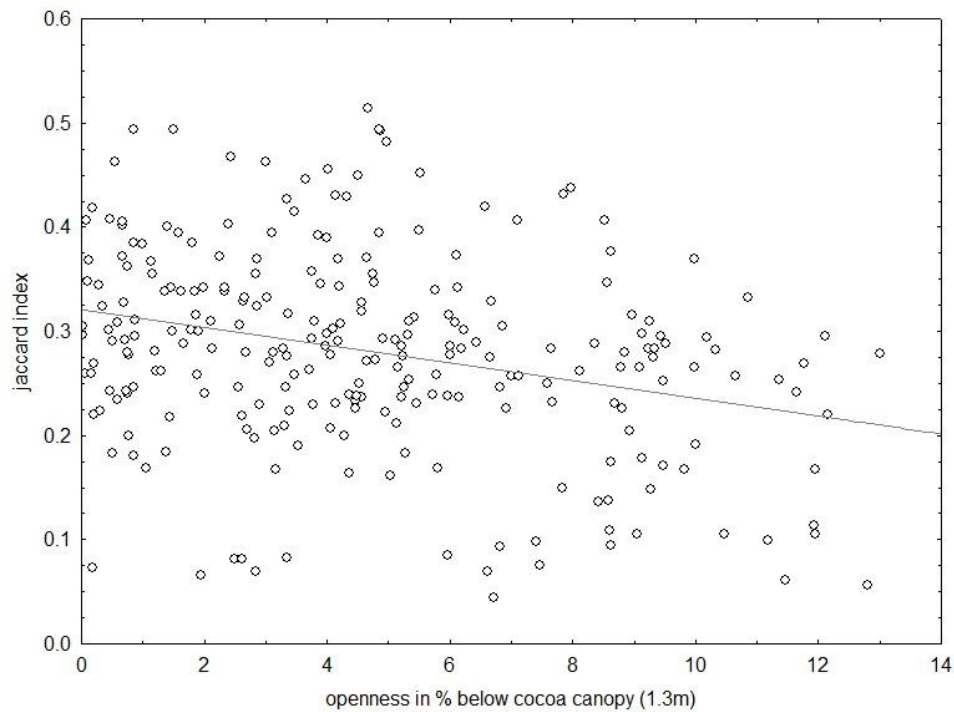


Figure 6 Correlation between similarity (Jaccard index) of foliar fungal endophyte composition and radiation (openness in % in 1.3m height) differences for all possible site pairs. $y = 0.32 - 0.01 \cdot x$; $r = -0.29$; $p < 0.000$; $r^2 = 0.09$.

The similarity between closely located plantations was higher compared to plantations separated by larger distance classes (Fig. 4). The regression between distance class and similarity (JI) was highly significant and explained 30% of the variance ($y = 0.34 - 5.23E-6 \cdot x$; $r = -0.55$, $p < 0.001$; $r^2 = 0.30$). Species richness of shade trees also influenced fungal endophyte composition (Figure 5). Plantations with a similar amount of shade tree species shared significantly more fungal endophytes with 27% explained variance (JI, $y = 0.33 - 0.01 \cdot x$; $r = 0.52$, $p < 0.001$; $r^2 = 0.27$). The diversity of weeds, as well as the distance to natural rain forests, did not show any significant effects on the endophyte composition (Tab. 3). For abiotic factors correlated with altitude, as annual mean and minimum temperature, a significant impact could not be proved. Canopy openness measured under the canopy of cacao correlated negatively with similarity (JI in endophyte composition (Fig. 6; $y = 0.32 - 0.01 \cdot x$; $r = -0.29$; $p < 0.001$; $r^2 = 0.08$).

Discussion

Total diversity

Host plant population characteristics, environmental conditions and whether a plant species is native to a given region will affect the diversity of interacting species, such as endophytes. Non-native plants are known to harbour a lower diversity of fungal endophytes than closely related native species in the same environment (Hoffman & Arnold 2007). In this study 149 fungal morphospecies were found in 544 cacao leaves. The estimated morphospecies richness (Fig. 3) suggests the regional endophyte community was well-sampled. This number is relatively low, when compared to the endophyte communities reported by Arnold et al. (2003), where 344 morphospecies were isolated from 126 cacao leaves in Panama, Central America. Arnold et al. (2000 and 2001) calculated species accumulation curves for endophytes of the hosts *Heisteria concinna* Standl. and *Ouratea lucens* Engl. in Panama, that suggest increase of sample size would lead to extraordinary high number of endophytes. Between-country differences in fungal endophyte diversity are not restricted to T. Cacao L.. Vega et al. (2009) investigated several tissues of non-resident coffee (*Coffea arabica* L.), from Colombia, Hawai'i, Mexico and Puerto Rico. The number of obtained genotypes for all tissue types ranged from 113 genotypes representing a Fishers alpha of 75.3 in Colombia to 32 genotypes representing a Fishers alpha of 14.9 in Mexico. The same variation occurs in *Musa accuminata* Colla, where 32 morphospecies could be found in Hong Kong, while in three different places in Queensland (Australia) could be found 15, 18 and 25 morphospecies (Brown et al. 1998). In the case of cacao, we expected a lower diversity in South East Asia than in South America, given that 1) it is likely that only a fraction of the endophyte community was introduced to Sulawesi together with the crop and 2) given the short time for adaptation, few fungi have at present adapted to become endophytic fungi within cacao,

being an introduced crop species. Unspecialised soil-borne fungal endophytes are challenged by different abiotic conditions within the new location and need to prevail against soil microorganism to re-infect leaves (Parker & Gilbert 2004, Herre et al 2007). Furthermore, an important prerequisite for any differences in fungal species composition between Sulawesi and Central America is that specialised endophytic fungi make up a significant part of the hosts species richness in the area of origin (Hoffman & Arnold 2008). The identity and specialisation of cacao endophytic fungi may help in understanding the mechanisms behind the low total diversity found in this study.

The dominant and subdominant genera found in this study have been frequently reported in previous studies. *Nodulisporium* species have been isolated as endophytes before (Petrini & Fisher 1990, Rodrigues & Samuels 1990, Fisher et al. 1992, Polishook et al. 1996, Polishook et al. 1999). Some *Nodulisporium* species act as pathogens (Stao et al. 1995), but some other isolates also showed anti-pathogenic potential in combinations with the fungal endophyte *Cordana* sp. against Anthracnose Disease in Banana (Nuangmek et al. 2008). *Fusarium* occurs nearly all over the world as a pathogen, endophyte or even antipathogen (Lodge et al. 1996, Evans et al. 2003, Kim et al. 2007, Vega et al. 2009). Unlike the genera named above, the widespread family of *Xylariaceae* is known to be species-rich and widely distributed in the American and Asian tropics. The genus *Xylaria*, especially, is a highly diverse taxon (Lodge et al. 1996, Bayman et al. 1998; Rogers 2000, Rubini et al. 2005, U'ren et al. 2009, Vega et al. 2009). These groups seems to contain fast growing competitive generalists, which have the capacity to colonize plants more successfully and thereby dominate the endophyte communities. A caveat may be that a bias through isolation technique favouring particular taxa cannot be entirely excluded (Hyde & Soyong 2008).

Dissimilarities in plant defences or plant metabolites between closely relative species may divide fungal endophyte communities in groups the tree is susceptible or unsusceptible for (Schulz & Boyle 2005). Arnold et al. (2003) found a significant amount of endophyte nonsingleton morphospecies occurring in only one of the tested host species (*T. cacao* L., *H. concinna* Standl. and *O. lucens* Engl.). The authors therefore concluded that host specificity might be prevalent among tropical endophytes. Further support comes from a study by Vega et al. (2009), who obtained only four genotypes of foliar endophytes occurring in more than one country in his survey for coffee plants endophytes in four tropical countries in Latin America. Because coffee does not originate from these countries, he suggested these endophytes are either generalists, or have coevolved and were distributed together with the hosts. Based on the hypothesis of host affinity, the comparably low number of fungal endophytes of cacao plants in Sulawesi represents decreased endophyte diversity. Some foliar endophytes specialized in cacao might has been lost as the host plant was introduced.

At the same time, other fungal endophytes may have established, forming new dominant groups in the host plant. *Xylariaceae* and *Fusarium* species for instance are frequently occurring in *Theobroma spp.* (Evans et al. 2003, Rubini et al. 2005, Crozier et al. 2006, Thomas et al. 2008, Mejia et al. 2008), while there have been no previous reports about *Nodulisporium* showing dominance in cacao endophyte communities. In our study, only a small amount of non-singletons were found in more than one site (Tab. 2). This may indicate host preference, by a small group of specialists, or the presence of a small pool of generalists with a broad environmental tolerance.

Because there are no studies available about endophyte diversity of African cacao trees, comparisons between Indonesia, Africa and South America are not yet possible. We expect the endophyte diversity in African cacao to be also dominated by generalists and reduced compared to South America, but possibly not as much as Asian cacao, given that a longer history of cacao cultivation may have provided enough evolutionary time for naïve fungal species to adapt to the new host (see Mitchell et al. 2010). The distribution of endophytes seems to be strongly affected by environmental factors like temperature, radiation and precipitation (Suryanarayanan et al. 2002, Hoffman & Arnold 2007, Arnold 2008). Comparisons between regions are difficult due to a lack of replication at that scale. However, our data allowed us to correlate within-region patterns in composition to environmental variables and discern spatial autocorrelation.

Diversity Gradient

The community of fungal leaf endophytes was dominated by a small group of numerically dominant morphospecies associated with an assemblage of sporadically occurring species. As a result the mean similarity of leaf endophyte composition among sites was low (Fig. 2). Investigating leaf-litter fungi of two tree species (*Guarea guidonia* (L.) Sleumer and *Manilkara bidentata* (A.DC.) A. Chev.) in two sites separated by 200m in Puerto Rico, Polishook et al. (1996) obtained comparable degrees of similarity (JI= 0.34, 0.38). Arnold et al. (2000) instead obtained a higher similarity value (JI= 0.48) for *H. concinna* associated leaf endophytes on sites 500m apart. The leaf endophyte composition predictably changed along the valley of the study area in Sulawesi. The highest similarity (JI) was found between neighbouring plantations (Fig. 2), suggesting either similarity in management, environmental conditions or cacao type, or a dispersal-limitation of cacao leaf endophytes, resulting in spatial structure. At distances of more than 300m, similarity decreases with distance between two sites (Fig.4). A similar pattern was found in cacao in Panama by Arnold et al. (2003), where the similarity (JI) of endophyte composition in cacao was reduced from 0.458 to 0.023 within 325km, and for *Cirsium arvense* (L.) in England (Gange et al. 2007), where similarity between endophyte composition was declining significantly within 52km. On the other hand,

Vega et al. (2009), who investigated the endophyte composition of coffee plants in four different countries found low similarities between sites, with a Jaccard index ranging from 0.226 to 0.092, regardless of distance between plots. In the investigation area in Sulawesi similarity between individual distant plots pairs did occur, and was likely due to similarity in environmental conditions (Fig.4). Very high similarity occurred between two plantations at high altitude, which presented, due to a steep slope in south direction, the same annual minimal temperature as sites at altitudes 100m below. This suggests that similarity of endophyte composition is not only a matter of endophyte dispersal, but as well affected by temperature, even though a general effect of temperature could not be verified in this study (Tab. 3). However, canopy openness slightly influenced endophyte composition. The similarity (JI) in endophyte composition between plantations was positively, although only weakly, correlated with the similarity in openness (Fig. 6). Openness below the cacao canopy depends on the density of the shade tree canopy and the cacao canopy, which can both be managed by planting, felling or pruning shade trees, or pruning the cacao trees. The openness above the cacao trees, representing the shade tree layer, was not significant suggesting an important role of the cacao layer. Shade tree diversity was much more important for the fungal endophyte composition (Fig. 5). A similar number of shade tree species grown within cacao plantations resulted in similar endophyte composition in the cacao canopy. Closely neighboured trees even of different species are known to share more endophytes with each other, than with distant trees of the same species or their relatives (Arnold et al. 2000, Arnold et al. 2003, Hoffman & Arnold 2008). Therefore it is likely, that most of the collected endophytes in this study are cosmopolitan generalists, or generalists reflecting a part of the regional/local occurring diversity. The amount of endophytes distributed with the host may be smaller than expected. Anyhow, the sample size with 16 leaves per plot may not have been sufficient to assess the complete gamma diversity of each plot (Fig. 3) which could lead to artificially higher dissimilarity, and a lack of power to detect further, but less important environmental gradients. An exhaustive sampling on the complete pool of plots may reveal further patterns, but was beyond the scope of this study.

Conclusions

Compared to South America, where the species originates, cacao (*Theobroma cacao* L.) trees grown in Indonesia harbour reduced diversity of foliar fungal endophytes. Endophyte diversity is spatially structured, following a gradient along the valley along which the sites were located. Endophyte composition was more similar grown with similar diversity of shade tree species and in similarly shaded environments. The results contribute to our understanding of the spatial patterns of leaf endophytes and environmental determinants. Manipulative studies and long-term surveys of endophyte community changes will be

required to better understand spatio-temporal patterns in endophyte communities and their functional consequences.

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Appendix

Table 1 Mean number of fungal endophytes isolated per cocoa leave, species richness of endophytes, and explanatory variables used in general linear models to explain fungal endophyte diversity in cocoa plantations along the Kulawi valley in Central Sulawesi. The variables are: Distance of each plantation to natural forest, altitude above sea level, radiation (measured by openness in %) above (5.8m) and below (1.3m) cocoa canopy, mean temperature within plantations, the number of non-cocoa tree species, the number of weed species, and the number of tall shade trees (h>15m).

Plantation No.	No./ Leave	Species richness	Distance to forest in m	Altitude ASL in m	Openness in % at 1.30m	Openness in % at 5.80m	Mean temperature per plot	n tree species	n weed species	n tall trees
1	0.5	7	550	675	20.01	55.99	23.60	13	19	16
2	3.56	38	adjacent	650	12.61	21.79	22.70	20	10	21
3	6.13	51	650	600	13.20	64.21	24.15	13	13	12
4	6.63	56	50	675	10.95	30.66	23.21	11	19	24
5	4.19	44	500	725	11.41	35.96	22.63	19	19	20
6	3.44	39	adjacent	750	17.18	64.44	22.43	8	19	4
7	6.38	57	300	675	14.05	33.02	23.07	12	2	27
8	3.94	42	700	650	8.08	34.81	22.90	9	6	9
9	2.06	26	650	625	8.25	24.80	23.39	7	8	22
10	6.63	56	350	575	9.54	65.46	23.97	5	15	2
11	4.56	50	650	550	7.20	20.26	23.31	4	10	32
12	3.25	37	500	550	8.83	29.27	24.24	7	5	17
13	4.44	45	2500	475	8.55	34.03	24.30	20	15	25
14	7.00	58	700	500	18.06	36.42	23.85	13	22	25
15	8.31	59	2000	425	17.52	76.65	24.61	8	14	11
16	3.81	37	1600	400	20.20	60.40	24.72	10	12	14
17	3.44	35	adjacent	400	8.05	51.21	24.28	13	14	12

18	3.63	39	650	700	11.39	36.04	23.58	11	20	21
19	7.13	64	400	775	12.54	27.15	22.41	11	12	29
20	7.19	60	100	800	17.39	36.19	22.80	15	20	20
21	6.63	54	300	800	13.39	53.95	22.67	8	11	5
22	2.94	39	adjacent	925	13.29	33.34	22.33	18	13	15
23	2.75	30	800	560	16.67	52.00	24.41	3	np	np

Table 2 Taxa of fungal leaf endophytes extracted from *Theobroma cacao* L. identified to the genus level based on spore characteristics after one month of UV-exposition. Non-sporulating isolates were grouped by mycelium characteristics.

Taxon	No.
<i>Nodulisporium</i> sp.	303
<i>Fusarium</i> sp.	280
<i>Xylariaceae</i>	214
Isolate A	165
Isolate B	101
Isolate C	47
<i>Colletotrichum</i> sp.	46
<i>Phomopsis</i> sp.	41
Isolate E	39
Isolate D	29
non-typed	27
<i>Lasiodiplodia theobromae</i> (Pat.)	26
<i>Acremonium</i> sp.	24
sporulating form x	21
<i>Virgaria</i> sp.	20
<i>Phoma</i> sp.	19
Pycnidial form	13
Hyphomycetes x	8
<i>Cladosporium</i> sp.	6
<i>Scopulariopsis</i> sp.	2
<i>Trichoderma</i> sp.	2
<i>Paecilomyces</i> sp.	2
Isolate F	2
Isolate G	2
Basidiomycota	1

<i>Geotrichum sp.</i>	1
<i>Gliomastix sp.</i>	1
<i>Guignardia sp.</i>	1
<i>Pestalotia sp.</i>	1
<i>Verticillium sp.</i>	1
Phialidic form	1
<i>Phialophora sp.</i>	1
<i>Botryotrichum sp.</i>	1

Table 3 Impact of assumed environmental parameters of diversity on similarity of endophyte composition calculated using the Jaccard Index between each possible pairs of plantations.

tested parameter	r ²	p
distance to forest	0.003	0.353
distance between plots	0.302	0.004 ***
altitude	0.011	0.121
openness in 1.3m	0.084	< 0.001***
openness in 5.8m	0.003	0.389
temperature mean	0.003	0.382
temperature minimum	0.002	0.527
tree species	0.271	<0.001***
weed species	0.006	0.219
n tall trees	0.011	0.103

Chapter 2:

Endophytes stressed out: Effect of a simulated ENSO-drought on fungal endophyte communities in cacao trees

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Summary

The distribution and diversity of fungal endophytes are driven by environmental factors as well as by host plant conditions. Climate change may alter the interactions between fungal endophytes and their host plants. Here, we analysed the composition of the foliar fungal endophyte diversity in cacao trees before and after an experimentally-imposed drought in Central Sulawesi, Indonesia. Cacao trees exposed to this drought treatment (a 13-month throughfall displacement) harboured significantly less endophyte diversity in their leaves than control trees. Additionally, fungal endophyte composition became more similar under the influence of drought. Fungal species known to have pathogenic potential against other endophytes outcompeted other taxa and became more abundant. The changes in the fungal endophyte composition induced by drought were related to seasonal changes, indicating that seasonal dry spells may also have contributed to these findings. This study highlights the close interaction between fungal endophytes and their host plants and suggests that the endophytes are potential indicators for climate change-induced impacts on the physiology of their host plants.

Keywords: cocoa, diversity, *Theobroma cacao*, seasonality, drought stress, climate change

Introduction

According to climate models, predicted changes in the El Niño Southern Oscillation (ENSO) (IPCC 2001) will cause droughts and other extreme weather events to become more frequent and severe in subtropical and tropical regions including Southeast Asia (Sheffield & Wood 2008). Drought extremes will greatly affect rain-fed crop plants and food resources worldwide (Patz et al. 2005). Most of the common crop plants are highly susceptible to changes in precipitation rates (Le Houérou 1996, Malinowski & Belesky 2000, Lloret et al 2004, Sivakumar et al. 2005), while trees in agroforestry plantations have deeper roots and are able to tolerate drought stress to some extent (Le Houérou 1996).

In many tropical countries including Indonesia, cacao (*Theobroma cacao* L.) agroforestry plantations are a mainstay of the economy and provide a sustainable livelihood for smallholder farmers (Smith et al. 1996, Duguma et al. 2001, Belsky & Siebert 2003, ICCO 2009). Cacao plantations in Indonesia, however, have suffered from severe ENSO related-droughts in recent decades (Quinn et al 1978, D'Arrigo et al 2006). Cacao production in the province of Central Sulawesi, Indonesia, is especially prone to drought, and cacao yields decreased up to 38% during an ENSO-related drought (Keil et al. 2008).

Plant tolerance to drought and drought-associated heat can be increased by certain fungi that live in the plant tissues without detriment to their host. Some of these endophytic fungi are able to stimulate genes related to drought stress (Sherameti et al 2008) and to regulate photosynthesis rates (Bacon 1993, Arnold & Engelbrecht 2007, Arnold & Herre 2003). Other fungal endophytes contribute to the recovery of host plants from drought effects by enhancing regrowth (Latch et al. 1985, Arechavaleta et al. 1989, Rahman & Saiga 2005). Host plants of arid deserts (El-Zayat et al. 2008, Porras-Alfaro et al. 2008), temperate regions (Espinosa-Garcia & Langenheim 1990, Unterseher et al. 2007), and tropical regions (Arnold et al. 2000, Arnold et al. 2001, Suryanarayanan et al. 2002, U'ren et al. 2007, Thomas et al. 2008) contain a species-rich community of fungal endophytes. The diversity and distribution of fungal endophytes are especially high in tropical plants (Arnold et al. 2000). Although these fungal endophytes are usually not strictly host specific (Cannon & Simmons 2002), Hoffmann & Arnold (2008) demonstrated that fungal endophyte diversity was less in non-native plants than in native plants in arid areas.

Cacao trees were introduced into Sulawesi from Latin America via the Philippines less than 25 years ago, and fungal endophyte diversity in cacao is lower in Sulawesi than in Latin America (Schmidt et al. unpublished). Because the fungal endophytes of cacao trees in Sulawesi are of tropical origin, the endophytes could be sensitive to drought, and drought could cause the endophytes to damage their hosts. Some fungal endophytes are able to

switch from a mutualistic to a parasitic or saprophytic lifestyle, depending on the species and the physiological status of the host plant (Müller & Kraus 2005, Schulz & Boyle 2005, Kogel et al 2006, Arnold & Engelbrecht 2007, Arnold 2007). These so-called latent pathogens may become actively pathogenic during drought stress.

In the current study, we experimentally induced a permanent drought event in a 7-year-old cocoa plantation in the upland region of Central Sulawesi, by establishing a roof below the canopy of the trees. Soil water content and sap flux of the host plants were measured continuously during the experiment, and foliar fungal endophyte composition was monitored before the experiment started and after 13 months of permanent drought stress. We tested the hypothesis that i) the species richness of fungal endophytes in the cacao trees would decrease within 1 year in response to the drought stress, and that ii) the abundance of latent pathogens would increase as cacao trees were subjected to drought.

Results

From 192 mature leaves of cacao trees, representing two treatments (roof and control) and two sampling periods (period 1 and 2), a total of 86 fungal morphospecies were recovered. Morphospecies richness per plot ranged from 13 to 32 before the drought treatment commenced (period 1). After 13 months of drought treatment (period 2), morphospecies richness ranged from 17 to 27 for the control and 11 to 16 for the roof plots (Tab. 1). Thus, the endophyte morphospecies richness in cacao leaves in period 2 was significantly less ($F = 7.9$, $df = 5$, $P = 0.044$) in the roof plots than in the control plots (Fig. 1). In addition, the number of morphospecies in the control plots was slightly smaller in period 1 than in period 2 (Tab. 1). The number of fungal morphospecies did not significantly differ between control plots and roof plots for period 1 ($F = 15.7$, $df = 5$, $P = 0.308$) or between roof plots for period 1 and period 2 ($F = 11.9$, $df = 5$, $P = 0.402$).

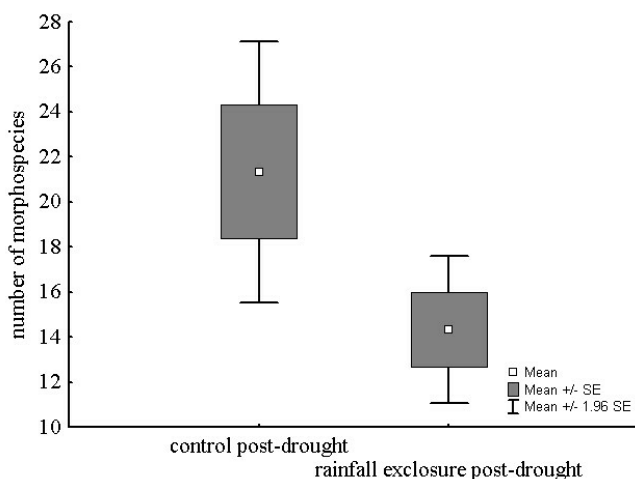


Figure 1 Difference between number of morphospecies in roof treatment and control in period 2. (generalized linear model based on Poisson distribution, intercept $<2e-16$, $p = 0.044$, dispersion parameter = 0.909).

Table 1 Species richness (N) for control (Cont) and rainfall exclusion plots (RFE), mean species richness per leaf in each plot (N/leaf) and alpha diversity index (Fishers alpha; +SD).

	Period 1					Period 2				
	Fishers					Fishers				
	N	N/leaf	SD	α	SD	N	N/leaf	SD	α	SD
I (Cont)	30	2.75	2.57	25.93	7.87	27	3.75	1.98	29.82	2.8
II (RFE)	15	1.88	1.78	27.13	5.04	16	2.25	2.18	29.93	2.64
III (RFE)	30	3.38	2.78	28.32	4.13	11	1.56	1.21	30.02	2.51
IV (Cont)	32	4.5	2.83	29.01	3.61	17	2.19	2.01	30.04	2.4
V (Cont)	20	2.88	2.36	29.39	3.25	20	2.56	1.86	30.04	2.31
VI (RFE)	13	0.81	1.76	29.63	2.99	16	2.56	1.15	30.01	2.23

We found a similar pattern when comparing fungal species composition pooled per treatment. A high similarity in species composition was found between control and roof treatments in period 2 (JI = 0.45, MHI = 0.88), while the similarities for roof treatments in period 1 vs. period 2 (JI = 0.30, MHI = 0.56) and for control treatments (JI = 0.35, MHI = 0.40) in period 1 vs. period 2 were very low (Tab. 2). Comparisons of all possible plot pairs (Morisita-Horn) revealed medium similarities for period 1, but relatively high similarities for period 2 (Fig. 2).

Table 2 Similarity indices comparing species richness for treatments and periods.

	Morisita-Horn index
pre- vs. post drought sampling	0.510
rainfall exclusion: pre- vs. post drought sampling	0.563
control: pre- vs. post drought sampling	0.397
pre-drought sampling: control vs. rainfall exclusion	0.799
post-drought sampling: control vs. rainfall exclusion	0.877

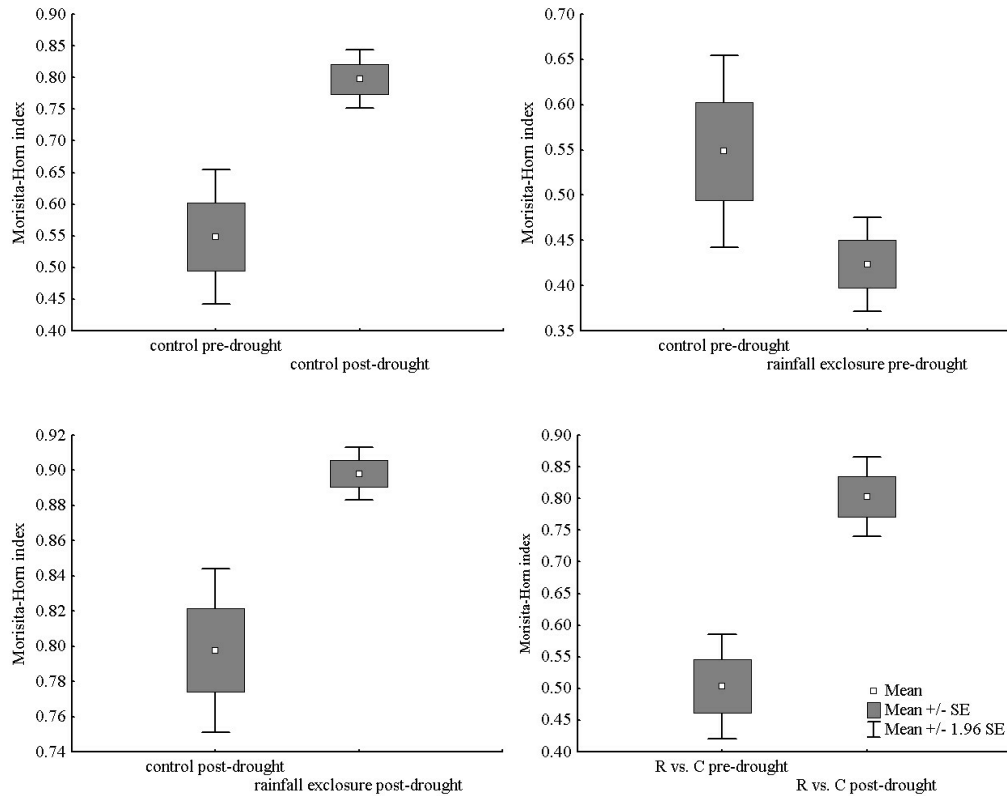


Figure 3 Comparing mean similarity (Morisita-Horn index) of a) controls of both periods ($p=0.014$), b) controls and roof plots of period 1 ($p=0.108$), c) controls and roof plots of period 2 ($p=0.016$) and d) all mixed pairs of roof and control comparing period 1 and 2 ($p>0.000$) with general linear model based on Gaussian distribution.

The mean similarity between all roof pairs in period 1 was not significantly different from the mean similarity of all control pairs in period 1 (Fig. 3b, $F = 4.2$, $df = 4$, $P = 0.108$). However, the mean similarity in period 2 was significantly higher in roof plots than in control plots (Fig. 3c, $F = 9.6$, $df = 4$, $P = 0.016$). Overall, the similarity of fungal endophyte morphospecies composition was significantly higher in period 2 than in period 1 within control plots (Fig. 3a, $F = 5.3$, $df = 4$, $P = 0.014$), within roof plots (not displayed, $P < 0.000$), and within all possible mixed pairs of roof and control plots (Fig. 1, Fig. 3d, $F = 12.0$, $df = 4$, $P < 0.000$). There were no significant correlations between species composition and the mean sap flux during the sampling periods for each sampled cacao tree (data not shown).

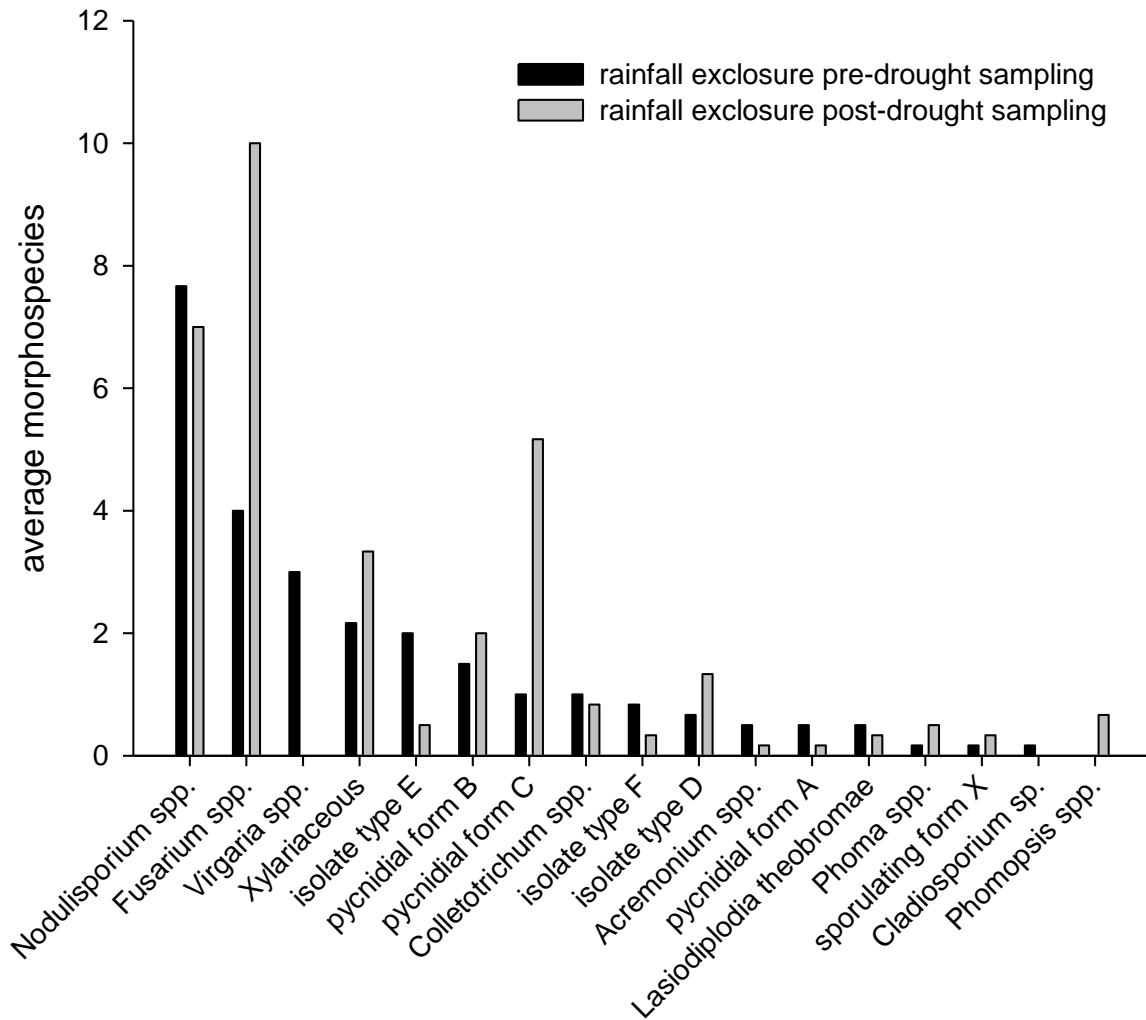


Figure 3 Abundances of all taxa identified based on spore or mycelium morphology of both periods. All subdominant taxa (categorized after Engelmann 1978) are labelled with value dominance.

Based on spore and mycelium characteristics, the 86 morphospecies were classified into 17 distinct taxa (Fig. 4). In period 1, *Nodulisporium* spp. (D = 30%) were the most abundant in both treatments, followed by *Fusarium* spp. (D = 16%), *Virgaria* spp. (D = 12%), and *Xylariaceae* spp. (D = 8%). In period 2, more species of the genus *Fusarium* (D = 31%) were found than species of the genus *Nodulisporium* (D = 21%) or of the genus *Xylariaceae* (D = 10%). *Virgaria* spp. were not found in period 2, while more pycnidium-forming species (type 3) were found (D = 16%) in period 2 than period 1.

Discussion

Species diversity

In this study, we isolated 86 morphospecies from 192 leaves of cacao trees growing in control and roof plots. Based on the results of previous studies, in which 149 morphospecies were isolated from 544 cacao leaves of 22 different cacao plantations in Central Sulawesi (Schmidt et al. 2010), we expected to find at least 100 fungal morphospecies. Previous surveys of the fungal endophyte diversity in cacao leaves in Panama detected 344 morphospecies from 126 cacao leaves (Arnold et al. 2003). As irradiance diversity has been shown to be a major driver for fungal endophyte diversity (Schmidt et al. unpublished), the low diversity of shade trees, represented by *Gliricidia* and coconut palms in the experimental plots, may explain the lower number of morphospecies found in the control and roof plots in the Marena area. Moreover, the high management intensity of the Marena plantation, including annual fertilizer applications and herbicide use previous to the experiment, might have affected fungal soil and leaf litter communities (Sarathchandra et al 2001). Because foliar fungal endophytes are horizontally dispersing by spores (e.g., Arnold 2007), foliar fungal endophyte diversity is related to the fungal leaf litter communities (Herre et al. 2007, Suryanarayanan et al. 2009, U'ren et al 2010). This relationship between endophytes in the litter and leaves and the previous treatment of litter with fertilizers and herbicides may explain the overall low diversity but high variability between the plots, resulting in low similarities (JI) between all treatments and periods (Table 2). The total species richness was additionally reduced in roof and control plots during the experiment (Table 1), and this may have partially masked the impact of the roof installation. We conclude that a drought stress that simulates an ENSO event significantly reduces the species richness of fungal endophytes.

Species composition

During the 13 months of experimental drought, the roof installation linearly decreased soil water content in roof plots to a depth of 2.5 m, diverting about 80% of the throughfall (Moser et al. 2010, Schwendenmann et al. 2010). Cacao trees responded to the reduced water availability with a monthly linearly decreasing sap flux density (Köhler et al. 2010), suggesting that the trees were indeed water stressed. Although water availability was continuously reduced in the roof plots during this period, cacao trees displayed no visible stress reactions, e.g., higher leaf fall as reported by Ling (1986) or Heuvel dop et al. (1988). Compared to control plots, the increase in defoliation on trees within roof plots was only minor (8%) (Schwendenmann et al. 2010). Apparently, cacao trees were able to mitigate the drought effects primarily by their shallow root system (Moser et al. 2010), which guaranteed

access to the throughfall that occasionally penetrated the roofs and rewetted the topsoil (Köhler et al. 2010).

Although the cacao trees appeared to tolerate the imposed drought, the endophytic fungi inside the leaves were sensitive to the altered environmental conditions, resulting in decreased species richness (Fig. 2). Because the fungal endophytes are located within the leaves and therefore were continuously exposed to humid conditions above the roof, it seems likely that the fungal endophytes were sensitive to changes in tree physiology rather than to the direct effects of reduced rain and humidity. The loss of species richness in period 2 (Fig. 1) coincided with the significantly higher similarities (MHI) in the roof plots than in the control plots (Fig. 3d), suggesting that selective forces specifically limited the re-colonisation of new leaves on cacao trees on stressed trees, thus contributing to the lower endophyte diversity.

Based on measurements of trees sap flux and tree water use, stress values were calculated by Köhler et al. (2009) for the same tree species used in our survey. Sap flux density in cacao trees was quite variable and changed during daytime (Köhler et al. 2009), and thus correlations between sap flow and the fungal endophyte communities in the leaves, which change only slowly in time (Arnold & Herre 2003), could not be expected. Thus, the reduced use of water by trees is not regarded as a driving force for decreasing fungal endophyte diversity. However, the endophytes may have responded to other changes in plant physiology. Saunders & Kohn (2009) have shown that plant defence compounds influence fungal endophyte communities. Because aboveground biomass of the trees was not directly affected by reduced access to water (Moser et al. 2010), we hypothesize that plant cells might have accumulated metabolites that prevented or reduced drought-induced damage by functioning as osmolytes (e.g., sugars), antioxidants (e.g., peroxiredoxins), or scavengers (e.g., reactive oxygen species) as described by Bartels & Sunkar (2005) and Seki et al (2007). It seems possible that an increase in concentrations of specific plant metabolites might have affected at least some fungal endophyte species, causing the observed change in endophyte composition (Schulz et al. 1999).

Seasonal changes and fungal species shift

The significant differences in the similarity of the fungal endophyte composition between plot pairs, observed within control plots in both sampling periods (Fig. 2, Fig. 3a) and observed when comparing plot pairs of both treatments from both periods (Fig. 3d) may be explained by i) a response to the naturally occurring dry spell from January to March 2008, which enhanced drought effects in roof and control plots (Schwendenmann et al. 2010) or by ii) a

seasonal variation in endophyte occurrence as reported by Suryanarayanan & Thennarasan (2004), Unterseher et al. (2007), and Porras-Alfaro et al. (2008). However, the endophyte communities of both periods were characterized by a group of commonly occurring and abundant endophytes (Fig. 4) known to be the most common endophytes in cocoa leaves in the Kulawi Valley (Schmidt et al. unpublished). Because most of the fungal endophytes recovered from the cacao leaves were found only infrequently as singletons or doubletons in the plots, the main changes in the similarity of the endophyte diversity during the two periods were due to a shift of dominances among the frequently occurring and often abundant endophyte species. For example, *Nodulisporium* spp. were the most abundant endophytes in period 1 but were second most abundant in period 2, whereas *Fusarium* spp. were the second most abundant in period 1 but the most abundant in period 2. *Fusarium* spp. produce a wide variety of mycotoxins (Guo et al. 2008) that could reduce the occurrence of other pathogenic fungi. The drought spell may have enhanced the mycotoxin production of the endophytic *Fusarium* spp., increasing its competitiveness with other fungal endophytes. By this mechanism, *Fusarium* spp. may have dispersed within the leaves, occupied many niches, and reduced the colonisation by other species, thereby increasing the similarity between the fungal endophyte communities. *Fusarium* spp. are causal agents of several plant diseases occurring near the plantation in the current study, and these include Fusarium wilt in *Musa* spp., *Coffea* spp., *Gossypium* spp., and *Elais guineensis*, and malformation of *Magnifera indica* fruits. These alternative hosts might have contributed to the higher colonisation rate by *Fusarium* spp. during period 2. *Fusarium* spp. are also the causal agent of cushion galls in cacao (Ploetz 2007), but no significant increase in plant pathogen incidence was observed on the trees (Schwendenmann et al. 2010). Contrary to other published reports (Suryanarayanan & Thennarasan 2004, Porras-Alfaro et al. 2008), *Xylariaceae* spp. were isolated more frequently following a period of low precipitation. This may indicate that the effects of roof-induced drought differ from those of naturally occurring drought as reported by Suryanarayanan & Thennarasan (2004) and Porras-Alfaro et al. (2008). In the current experiment, the leaves of the cacao trees in the roof plots were exposed to precipitation, whereas precipitation was drastically reduced and leaves were not exposed to precipitation in the other studies mentioned above. Temperature, radiation intensity, and humidity also differ between a roof-induced drought and a natural drought.

Conclusion

In terms of short-term biomass accumulation, the cacao trees showed no obvious response to the drought stress (Moser et al. 2010) and can therefore be regarded as resistant to ENSO-drought events, which typically last 12 months (Trenberth 1997). Cacao yields, however, decreased by 45% following the experimental drought (Schwendenmann et al.

2010), indicating that cacao production may be substantially reduced by ENSO drought events. Following a 13-month throughfall displacement in the cacao 'roof' plots, the fungal endophyte composition in the leaves of cacao trees was significantly reduced relative to control plots. Because the roof in this experiment was built below the canopy of the cacao trees, adaptations of the fungal endophyte composition must have been driven indirectly via the colonised host plants. The reduced diversity of fungal endophytes and the high similarity in species composition in drought-exposed plots indicates that host-plant chemistry seems to control endophyte diversity. Fungal endophytes are evidently highly sensitive to host-plant conditions and may be better indicators of stress induced by climate change than host-plant growth or sap flux. Other effects associated with natural droughts (e.g., increased radiation, heat-damaged leaves, and reduced humidity) that have been shown to affect fungal endophyte composition (Suryanarayanan et al. 2002, Schmidt et al. unpublished) were not manipulated in this experiment and did not differ between roof and control plots (Moser et al. 2010). Understanding the complex interactions between endophytes and their host plants under climate change will require additional research on how plant metabolism changes in response to drought.

Experimental Procedures

Study site

The drought simulation experiment was conducted in a 7-year-old cocoa agroforestry plantation in the western margin of the Lore Lindu National Park in Central Sulawesi (1.552°S, 120.020°E); this is in the centre of the cocoa production region in Indonesia. The plantation was located 560 m above sea level on a gentle slope (8–12°) where the ground water table was deeper than the tree rooting zone (Moser et al. 2010). The soil was a sandy-loam Cambisol (Schwendenmann et al. 2010). The mean annual temperature at the nearby meteorological station was 25.5°, with an average annual precipitation of 2092 mm year⁻¹ (Schwendenmann et al. 2010). The cacao trees were growing under an overstorey of *Gliricidia spium* trees interspersed with a few coconut palms (*Cocos nucifera*) in a plantation comprising 8400 m². During the experiment, the cocoa trees were pruned in July and December 2007. No other agricultural measures than biweekly harvesting of the cocoa pods and manual removal of weeds were performed. Before the experiment started, the plantation had been fertilized annually.

Artificial drought

The plantation was divided into six plots of 40 x 35 m each. Three plots were randomly designated as undisturbed control plots, and the other three plots were designated as treatment or "roof" plots. In the latter plots, a transparent roof was installed below the cocoa

canopy. The roof diverted about 80% of the throughfall, while temperature, humidity, and incident radiation under the cover were unaffected by the roof (for further details, see Schwendenmann et al. 2010). Because the roof was located below the cacao canopy, cacao leaves were still exposed to precipitation, which is regarded as important for colonisation of foliage by fungal endophytes (Arnold & Herre 2003). The experiment started in late January 2007, when the roofs were closed and ended with a rewetting phase in April 2008, when the roofs were reopened. During the experiment, soil water content was measured in soil-pits (Köhler et al. 2010). Sap flux of specific cacao trees, monitored for their fungal endophytes, was measured and tree water consumption was calculated in order to confirm water stress (Köhler et al. 2009, Köhler et al. 2010).

Sample collection

The endophytes were collected from four randomly selected *Theobroma cacao* trees in each plot. Edge effects were avoided by excluding trees on the plot margins. Leaves were first sampled on 12 March 2007, in advance of the drought effects created by the installed roofs. This sampling of unaffected roof plots and control plots is referred to as period 1. Shortly before roofs were opened on 8 April 2008, when the drought was at its maximum (Köhler et al. 2010), leaves were sampled again from the same trees; this sampling is referred to as period 2. Leaves were sampled according to the methods of Schmidt et al. (unpublished). One leaf was collected from each of four branches from each cacao tree, giving a total of four leaves per tree. Two of the branches on each tree were located high in the canopy and two were located low in the canopy. Exclusive harvesting of matured leaves was assuring the assessment of a fully established endophyte community (Arnold et al. 2003). After sampling, the leaves were sealed in sterile polyethylene bags and transported to the laboratory at the Tadulako University (Palu, Central Sulawesi, Indonesia), where they were stored at 8°C. The leaves were shipped to Germany by airmail for isolation of endophytic fungi.

Fungal endophyte isolation

Endophytic fungi were isolated following the protocol of Arnold et al. (2000). The collected leaves were surface sterilized in 3% NaOCl for 3 minutes, placed in 70% ethanol for 3 minutes, and then rinsed in sterile-distilled water for 3 minutes. From each sterilized leaf, five discs were cut with a flame-sterilized circle-cutter size 6 (78.54 mm²). For each leaf, one disc was cut from the endophyte-rich midvein on the leaf tip (Cannon & Simmons 2002), and four discs were cut along the midvein, with 3 cm between each disc. The five leaf discs were placed on 2% MEA (malt extract agar, Roth GmbH, Germany) containing streptomycin (600 mg/L) in one 9-cm Petri dish, which was sealed with Parafilm. After 7 days at 25°C, fungi growing out of the leaf discs were isolated and identified to morphospecies based on the

following characteristics: colony surface texture, hyphal pigmentation, colour of exudate and growth rate (Brown et al. 1998, Arnold et al. 2000, Suryanarayanan & Vijaykrishna 2001). We excluded isolates of *Aspergillus*, *Penicillium*, and *Mucor* species following the suggestions of Hyde & Soyong (2008), because we assumed that they were contaminants. For sporulation, the isolated fungi were transferred to a thin layer of nutrient less MEA (Brown et al 1998) and stored in a UV-chamber at 20°C with a light regime of 12 h light:12 h darkness. Following 1 month of UV radiation, fungal morphospecies were identified based on their spores. Several of the fungal endophytes remained sterile under UV- and nutrient-stress conditions, and these were classified based on the characteristics of their mycelia.

Statistical analysis

We used leaf-level presence/absence data of the fungal endophyte morphospecies for the analyses. Because four leaves were collected from each tree, each tree had a maximal fungal endophyte abundance of four, while each site had a maximal fungal endophyte abundance of 16. Besides species richness, Fishers α and dominances (D, categorized in dominance classes, following Engelman 1978) were calculated to assess the gamma diversity of the cacao plots. For testing the differences between treatments, a generalized linear model was calculated with adjustments in case of over-dispersion using the program R (version 2.11.1) and after the suitability of the Poisson distribution for the sampled endophyte species richness data was tested (Onofri et al. 2010). Because of the time interval of 13 months between the sampling periods, leaves collected during period 2 were not present during period 1, and we therefore regarded the samples as independent of each other. A linear regression was calculated using Statistica (Version 2.0) to relate endophyte species richness to the mean monthly sap flux; sap flux data were obtained from Köhler et al. (2010). Similarity indices of endophyte composition were used to calculate differences between roof plots and control plots before and after the roof closure. Both the presence/absence-based Jaccard index (JI) and the abundance-based Morisita-Horn index (MHI) were computed because these indices use different weightings of community characteristics and therefore differ in sensitivity. Both species diversity and shared species analysis were calculated using Estimate S version 8.2 with 1000 randomizations (Colwell 2008). Because species with high abundance in all plots are weighted higher, the MHI was regarded as more appropriate for measuring similarities and changes in fungal endophyte composition between the treatments. Values for MHI between all combinations of plot pairs were grouped for all combinations of treatments and periods. Differences between treatments and periods were tested with a generalized linear model based on a normal distribution and using the program R.

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Chapter 3:

Distribution and Diversity of *Fusarium spp.* in Cacao Plantations of Central Sulawesi - Indonesia

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Abstract

The ecology and distribution of endophytic occurring *Fusarium spp.* in the tropics is largely unknown. In this investigation the phylogenetic relation among 172 *Fusarium* isolates, sampled in 23 cacao (*Theobromae cacao*) plantations along environmental gradients, was studied using partial sequences of the ribosomal large subunit (LSU) and transcription elongation factor 1-alpha (TEF 1 α). The observed diversity of endophytic *Fusarium spp.* was comparable to the *Fusarium* diversity that was found previously in cacao of South America. The composition of phylogenetically distinct *Fusarium* groups in the cacao plantations was dependent on local plant composition, and was reflecting Sulawesi's characteristic *Fusarium* diversity. The abundance of *Fusarium* phylogenetic groups was changing along local temperature gradients and seasonal patterns. Two shade tree species (*Piper anduncum* and *Toona ciliata*) within the cacao plantations could be discriminated based on their *Fusarium ssp.* composition. We hypothesize that the latter trees influence the *Fusarium* composition of a habitat by antifungal compounds. The neighbourhood of Mango trees (*Mangiferae spp.*) showed an enhanced *Fusarium* diversity compares to cacao trees. Neither yield losses nor symptoms of *Fusarium* induced diseases have been observed in the plantations investigated in this study, despite of the high diversity or abundance of *Fusarium spp.*

Introduction

Due to its high species diversity and its wide host range, *Fusarium* is one of the most widely distributed microbe associated to plants (Summerell et al. 2003, Leslie & Summerell 2006). Many *Fusarium spp.* occurs as pathogens of various crops from temperate areas (e.g. Yli-Mattila et al. 2004, Logrieco et al. 2002, Punja et al. 2008, Nitschke et al. 2009) to the tropics (e.g. Ploetz 2006, Marasas et al. 2006, Dita et al. 2010). *Fusarium* pathogens cause worldwide considerable economic yield losses in field crops and agroforestry (see Leslie & Summerell 2006). Furthermore, non-pathogen *Fusarium spp.* can also occur as endophytes without detriment to their host plants (Rubini et al. 2005, Evans et al. 2003, Vega et al. 2009, Pinaria et al. 2010). Some endophytes gained high popularity as biocontrol agents. Specifically non-pathogenic species and strains of the genus of *Fusarium* attracted high interest (e.g. Boyette & Walker 1985, Freeman et al. 2002, Larkin & Fravel 2002, Fravel et al. 2002, Menjivar Barahona 2010). This suggests that investigations of *Fusarium* diversity in unexplored habitats might be a promising approach to discover new potential biocontrol agents. Because endophytic fungal diversity is generally correlated to plant species diversity (Hyde 2001, Cannon & Simmons 2002, Schmidt et al. unpublished), tropical rainforests can be regarded as a valuable source of yet unknown *Fusarium spp.*, strains or varieties (Arnold et al. 2000, Arnold et al. 2001, Arnold & Lutzoni 2007). In an investigation of leaf endophytes in 23 cacao (*Theobroma cacao*) plantations in Central Sulawesi (Indonesia) *Fusarium* was found to be the second most abundant taxa (Schmidt et al. unpublished). Because the fungal endophyte morphospecies richness correlated to the diversity of planted shade trees along the investigated plantations, it seems most likely that the cacao trees hosted more than one *Fusarium spp.* This assumption was supported by the results of a study, in which the changes in fungal endophyte composition of a cacao plantation were investigated under drought stress (Schmidt & Vidal unpublished). After 13 month of artificial drought *Fusarium spp.* reacted to the induced changes in host plant conditions and became the dominant genus, pointing towards a possible increase in *Fusarium* diversity in cacao. Further support to the above mentioned hypothesis comes from Pinaria et al. (2010), who investigated *Fusarium* species associated with *Vanillae planifolia*, which is in Sulawesi commonly grown around stems of cacao plants, in seven locations in Indonesia. Half (six of twelve) of the *Fusarium* species which had been identified in Pinaria et al. (2010), were previously known from North Sulawesi (Pinaria et al. 2010), suggesting a high diversity of *Fusarium* species in Central Sulawesi.

Some *Fusarium spp.* are known to have distinct climatic preferences for their distribution (Summerell et al. 2003). But besides regional climates, the drivers of *Fusarium* diversity are largely unknown. The aim of the present research was to screen *Fusarium* diversity based on the molecular identification of fungal isolates collected from 23 cacao plantations in two

previous studies (Schmidt et al. unpublished, Schmidt & Vidal unpublished). Another focus of the investigation was to survey for characteristics in *Fusarium* diversity along the Kulawi valley. In a previous investigation about endophyte diversity in cacao after drought simulation, several *Fusarium spp.* were isolated from sample leaves (Schmidt et al. unpublished). These results and isolates were used in the present study to determine whether *Fusarium spp.* populations varied following drought stress and whether this yielded to an increase in *Fusarium spp.* diversity.

Material and Methods

Investigation area and sampling

Fungi were isolated from cacao leaves of 23 cocoa plantations along the western margin of Lore Lindu National Park in Central Sulawesi, Indonesia. A total of 22 of the investigated cacao plantations were managed by small holders along the Kulawi valley (hereafter addressed). In an additional plantation close to the village Marena (hereinafter called Marena plantation), an artificial drought was established for a period of 13 month. The conditions and performance of the drought experiment was already described in Schmidt & Vidal (unpublished). All samplings of fungal endophytes were conducted in March 2007. The Marena plantation was sampled again in April 2008, when the introduced drought reached its peak level (see. Schwendenmann et al. 2010). All investigated plantations succeeded the following environmental gradients: distance to natural rainforest, temperature, weed diversity, shadetree composition and degree of shading and solar exposition (see Schmidt et al. unpublished).

Isolation of *Fusarium* species

The sampled cacao leaves were immediately enclosed in sterile polyethene bags and transported to the laboratory of the University of Tadulako (Central Sulawesi), where they were stored at 8°C and processed within the following two days. The surface sterilization of the collected leaves was performed following the protocol suggested by Arnold et al. (2000). In 3min steps, whole leaves were bleached in 3% NaOCl, washed in 70% ethanol and rinsed in distilled water. Five leave discs were cut out of each sterile leaf using a flame sterilized circle-cutter size 6 ($A=78.54\text{mm}^2$). One of the five leaf discs was cut from the leaf tip, which is known to contain most endophyte diversity (Cannon & Simmons 2002). Four further leave discs were cut in the leave periphery aside the midvein to assure a leaf-representative endophyte extraction. The five leave discs of one leaf were transferred on antibiotic (streptomycin 600ppm) malt extract agar (Roth, Germany) in a 9cm petri dish sealed with parafilm and incubated at 25°C for seven days. Fungi that grew out from the leaf discs were transferred into a new plate and purified through several isolations. Pure fungal culture were

stimulated for sporulation using nutrient less malt extract agar and UV (light regiment UV darkness 12:12h). After one month of UV radiation, *Fusarium spp.* were characterized based on their spores morphology.

Fungal DNA extraction

For fungal DNA isolation, 4ml malt extract broth media (Roth) were inoculated with the fungus and shaken for growth at 25°C and 300rpm for five days. Hereinafter the hyphae were collected by vacuum-filtration and the mycelia were then freeze-dried for two days. Dried fungal material was used for DNA extraction according to the CTAB-protocol of Doyle & Doyle (1990). The sedimented DNA was suspended in 30µl TE-buffer.

Diversity study of selected *Fusarium* isolates

Following comparable fungal diversity investigations, analysis by PCR amplification of the LSU followed by sequencing of the obtained PCR products was performed (Aime & Phillips-Mora 2005, Crozier et al. 2006). For this purpose the primers LR0R (5'ACCCGCTGAACTTAAGC 3', Moncalvo et al. 1995, Maier et al. 2003) and LR6 (5'CGCCAGTTCTGCTTACC3', Vilgalys & Hester 1990, Aime & Phillips-Mora 2005) were used. The PCR reactions contained 25ng fungal DNA, 1x PCR-buffer (10x, Bioline), 1.5mM MgCl₂ (Bioline), 0.4µM each of forward/reverse primer, 200µM of a dNTP-mix and 1 Unit of Taq-DNA-Polymerase (Bioline). After a first denaturation step of 5min at 94°C a total of 35 cycles each of 94°C for 30s, 50°C for 45s and 72°C for 1 min were run followed by a final extension step at 72°C of 7 min.

To achieve differentiation between species of the genus *Fusarium*, the translation elongation factor (TEF) 1α gene was selected (O'Donnell et al. 1998, Nitschke et al. 2009). For PCR amplification the primers EF1 and EF2 (O'Donnell et al. 1998), have been modified by adding a restriction site on the 5'-end of each primer, resulting in 5'TCAGTAGCGGCCGCATGGGTAAGGARGACAAGAC3' and 5'AGACCCTGCAGGGGARGTACCAGTSATCATGTT3'. The PCR was conducted in 25µl reactions containing 0.3µM primers, 1x PCR-buffer (10x, Bioline), 2mM MgCl₂ (Bioline), 200µM dNTP-Mix, 1 Unit of Taq-DNA-Polymerase (Bioline) and 10ng fungal DNA and used in the following PCR-program: 94°C for Xmin, 30 cycles of denaturing at 94°C for 20s, annealing at 54°C for 20s, and extension at 72° for 35s followed by an additional extension time for 2 min at 72°C.

Sequence analysis

LSU and TEF-1α PCR products were purified using SureClean (Bioline, Luckenwalde, Germany) following instructions of manufacturer. In order to proof quality and quantity of the PCR products these were separated on a 1.6% agarose-gel and then send for sequencing.

All LSU PCR products were sequenced by Macrogen (Korea) and all TEF-1 α PCR products were sent for sequencing to MWG Biotech (Germany). The obtained sequences were aligned with known sequences in the databases NCBI and EMBL. Alignments and phylogenetic analysis were performed with MEGA4. Before conduction of phylogenetic analysis, the sequences were manually aligned on their 5' site and hereafter the 3'-ends were trimmed in order to obtain sequences of equal length. All LSU- and EF-1 α sequences were aligned by ClustalW (Thompson et al., 1994). Maximum Likelihood of alignments was performed to analyze the phylogenetic relationship of the given samples. For the clustering gaps were treated as missing data. The estimation of nodal support was performed by 1000 bootstrap replications.

Additional statistics

Different statistical tests were used to correlate *Fusarium* diversity with environmental conditions based either on morphological or molecular data. The integration of *Fusarium* in the fungal endophyte community was calculated by a non-metric multidimensional model (NMDS) (Minchin 1987) based on the morphological data (Schmidt et al. unpublished). Thereafter to identify ecological drivers of diversity, the environmental data of the plantations were tested against the NMDS using the program R (version 2.11.1). The impacts of: environmental factors, time discrimination and shade tree species on the distribution and diversity of *Fusarium spp.*, represented by LSU sequences were tested. Therefore a generalized linear model analysis of the phylogenetic clusters from the LSU alignment based on poisson distribution was calculated (Venables & Ripley 2002, Onofri et al. 2009). The model was performed and tested in R. Afterwards phylogenetic clusters of the LSU were correlated with environmental factors – the significance of the correlation was tested using a linear regression with the program Statistica (Version 2.0).

Results

Cluster analysis of the LSU of 172 fungal isolates

The phylogenetic analysis of 172 *Fusarium* isolates and seven species from the NCBI database reveals two main clusters with bootstrap values higher than 85% (Fishbein et al. 2001). The biggest cluster, with a bootstrap value of 92% comprises the sub-clusters B, C and D (Fig. 1, A, B, C and D). Cluster A comprises three sub-clusters, in which most of the previously described *Fusarium spp.* arrange. In this cluster there is only one of the *Fusarium* isolates (160), clustering close to *F. ambrosium*. The next cluster (B) consists in a sub-cluster with three isolates (247, 129 and 221), which are grouping with 243. In cluster C there are 14 isolates and no known species defining the origin of the isolates, while cluster D houses the rest of the isolates (153) and *Fusarium oxysporum*. This last cluster (D) can be

divided into two groups (1 and 2), distinguished by a low bootstrap value of 61%. Interestingly, both groups included two different sequences of *Fusarium oxysporum*, corroborating the solidity of cluster D in terms of the distinction of one specific taxa, in this case, *F. oxysporum*. The *Fusarium spp.* clustered in geographical ($p=0.026$) and seasonal groups ($p<0.001$, Fig. 2). For example the whole cluster C and group 1 of cluster D (LSU D1) mainly contained *Fusarium spp.* Isolated from the plantations along the Kulawi valley sampled in 2007. In the contrary isolates of cluster A and group 2 of cluster D (LSU D2) were mainly sampled in 2008 in the Marena plantation when the drought achieved its maximum level. There was no difference in *Fusarium* diversity or abundance between the 22 plantations in the Kulawi valley and in the Marena plantation before the drought experiment was conducted.

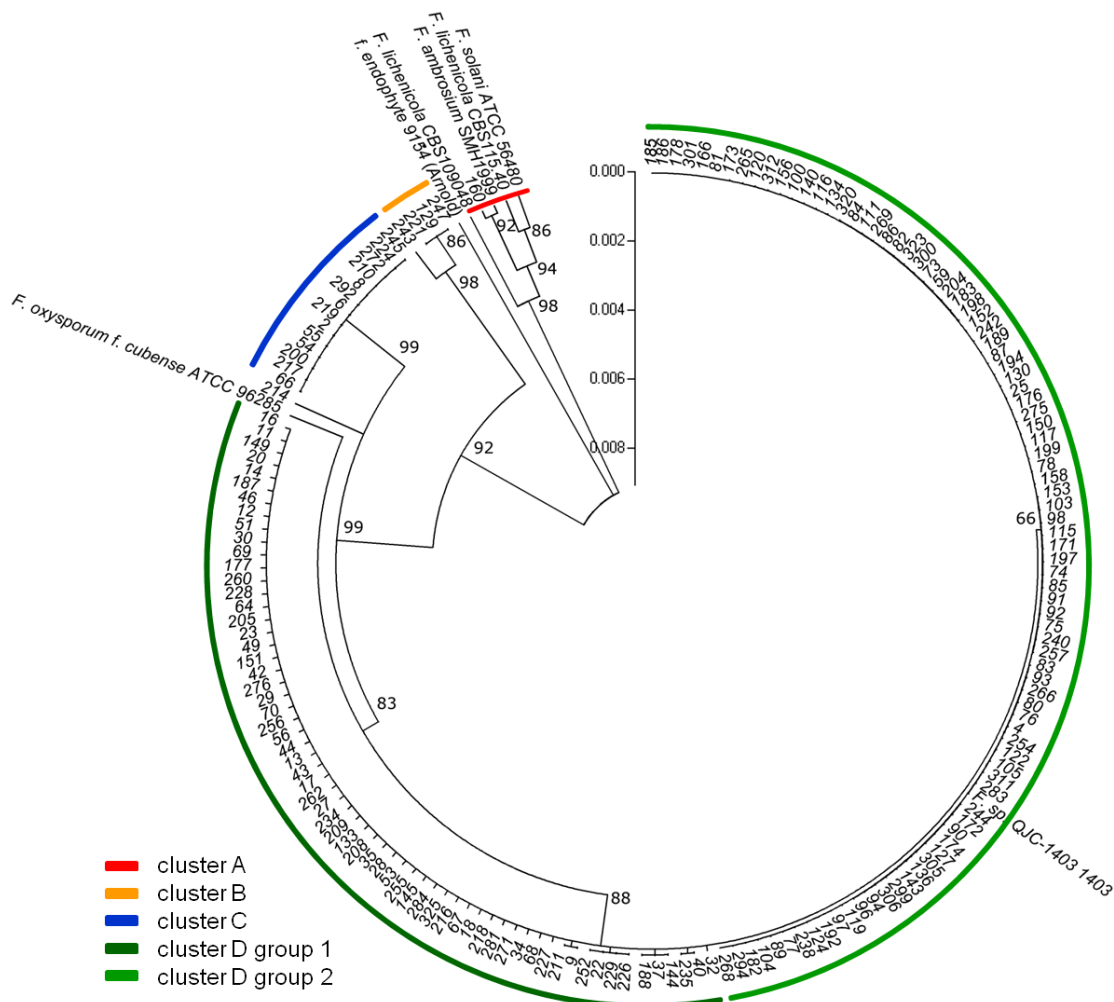


Figure 1 Phylogenetic analysis of 179 *Fusarium spp.* The evolutionary history was inferred using the Neighbor-Joining method. The optimal tree with the sum of branch length = 0.05362127 is shown. The tree is drawn to scale, with branch lengths in the same units as those of the evolutionary distances used to infer the phylogenetic tree. The evolutionary distances were computed using the Maximum Composite Likelihood method and are in the units of the number of base substitutions per site. All positions containing gaps and missing data were eliminated from the dataset. This analysis was conducted in MEGA4.

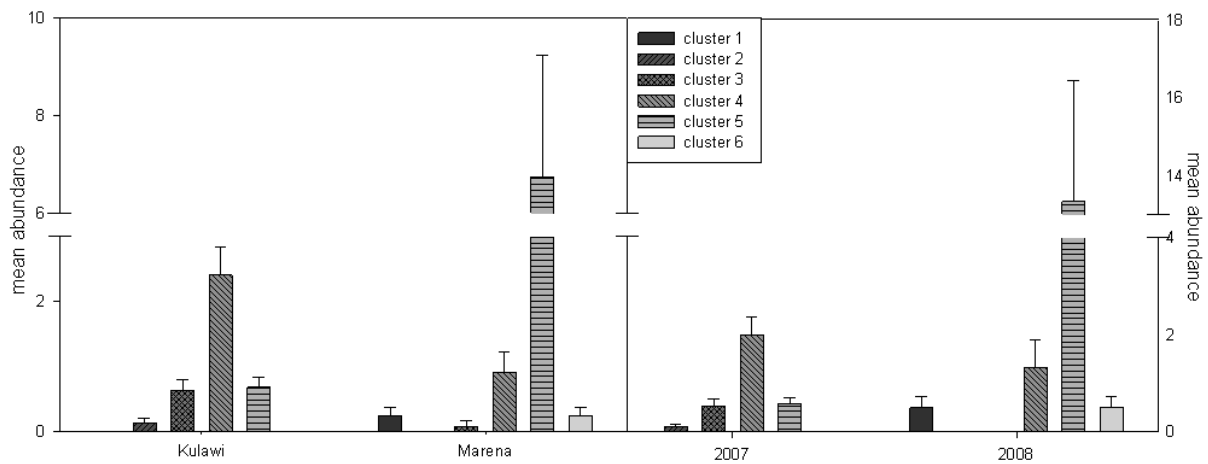


Figure 2 Mean isolate abundance (with standard error) based on LSU clusters and groups separated for two different locations and for two different sampling seasons. The distribution of *Fusarium* endophytes differed significantly by the influence of the locations ($p=0.026$) and the sampling seasons ($p<0.001$).

Correlation analysis

In the species composition of the Kulawi valley, *Fusarium* isolates were located centrally on the first axis of the NMDS model, reflecting the main gradient of distribution (Fig. 3). From the second axis it is however apparent that *Fusarium* isolates are affected by the opposed gradients of minimal annual temperature ($p=0.014$, $r^2=0.379$) and altitude ($p=0.0052$, $r^2=0.449$). The distribution of *Fusarium* clusters showed similar pattern: the abundance of isolates per cluster correlated positive with the minimal annual temperature ($p=0.036$, $r^2=0.233$, Tab. 1), while the total species richness decreased with increasing altitude ($p=0.028$, $r^2=0.207$). In a generalized linear model over all 6 LSU clusters the minimal annual temperature expressed a significant impact on the distribution of clusters over the area ($p=0.007$), while altitude proved no relevance concerning the diversity of *Fusarium* spp.. In a linear regression the general abundance of all *Fusarium* clusters was decreasing with increasing altitude over sea level ($r^2=0.207$, $p=0.028$, Tab. 1), suggesting that besides radiation, altitude might be the the main driver of temperature. Additional the minimal annual temperature correlated significantly with the mean *Fusarium* abundance per LSU cluster ($r^2=0.233$, $p=0.036$) and with the annual cacao yield in total ($r^2=0.210$, $p=0.033$, Tab.1). The cacao yield correlated positively with the diversity of LSU clusters ($r^2=0.425$, $p=0.003$) and the *Fusarium* abundance ($r^2=0.249$, $p=0.029$). There was also a strong correlation between the diversity of *Fusarium* clusters and the infection levels of the cacao pod borer ($r^2=0.491$, $p<0.001$, Tab. 1).

Table 1 Significant correlations of a linear regression between factors along the cacao plantations in the Kulawi valley. The significance level is rated with * for $p < 0.05$, ** for $p < 0.01$ and * for $p < 0.001$. The direction of the correlation is shown by + for positive correlated and - for negative correlated factors.**

factor 1	factor 2	formula	r^2	p		
number of LSU clusters	annual yield	$y = -31.3617 + 53.5482 \cdot x$	0.425	0.003	+	**
number of LSU clusters	cacao pod borer	$y = -0.189 - 0.2602 \cdot x$	0.491	> 0.001	+	***
<i>Fusarium</i> abundance	annual yield	$y = 24.4181 + 13.669 \cdot x$	0.249	0.029	+	*
<i>Fusarium</i> abundance	altitude (asl)	$y = 9.5846 - 0.0082 \cdot x$	0.207	0.028	-	*
mean abundance/cluster	min. temperatur	$y = -5.881 + 0.3259 \cdot x$	0.233	0.036	+	*
annual yield	min. temperatur	$y = -753.7016 + 41.6465 \cdot x$	0.210	0.033	+	*

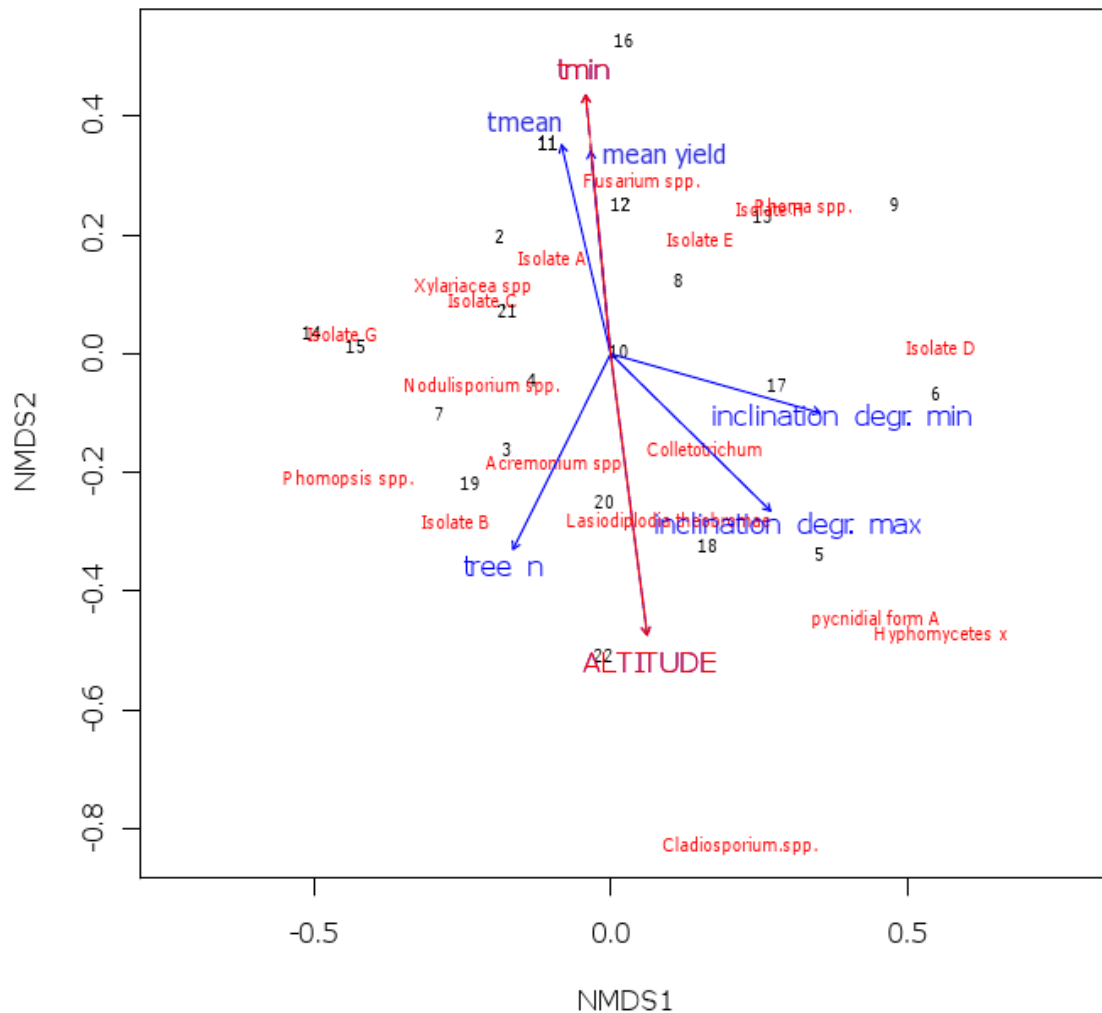


Figure 3 Calculated non-metric multidimensional model of the fungal endophyte diversity in cacao plantations along Kulawi valley, based on morphological data. Altitude ($P = 0.005$) and minimal annual temperature (t_{min} , $P = 0.014$) displayed significant impact on the fungal distribution and diversity. The probability error of the impact by mean temperature (t_{mean}), number of shadetrees ($tree\ n$), maximal and minimal degree of plantation inclination ($inclination\ degr.\ min$ & max) and the mean annually yield lied between 10% and 5%.

Several shade tree species were found to have significant impact on the distribution of *Fusarium spp.* belonging to each LSU cluster respectively (Tab. 2). Some of the species that demonstrated an influence are planted intercrops like the Indonesian endemic mango (*Mangifera mino*) and its introduced relative (*Mangifera indica*), the endemic water apple (*Syzygium aqueum*) and an introduced coffee cultivar (*Coffea robusta*). Other forest trees like gliricidia (*Gliricidia sepium*) and toon (*Toona ciliata*) displayed also a significant correlation to the *Fusarium* cluster distribution.

Table 2 Positive tested tree species, which have an influence on the distribution of the different *Fusarium* groups of the LSU data. P-values smaller than significance level (0.05) were enlisted.

tree species	cluster LSU C	group LSU D1	group LSU D2	all groups
<i>Mangifera indica</i>	ns	0.021	0.033	0.039
<i>Mangifera minor</i>	0.05	ns	0.025	ns
<i>Gliricidia sepium</i>	ns	ns	0.022	ns
<i>Piper anduncum</i>	ns	0.017	ns	ns
<i>Syzygium aqueum</i>	ns	0.003	ns	ns
<i>Toona ciliata</i>	ns	0.013	ns	ns
<i>Coffea robusta</i>	ns	0.049	ns	ns

***Fusarium* diversity based on TEF 1-alpha analysis**

In order to relate representative *Fusarium* isolates of the LSU clusters to known *Fusarium* species, 17 sequences of apparently suitable species were included in the TEF 1-alphacluster analysis. Figure 4 shows the result of a neighbor joining tree based on maximum likelihood similarities of the alignment of the samples. The high bootstrap values allowed a differentiation of six clusters using *G. zeae* as an outgroup, however the focus of the present study stands in four clusters, namely clusters I, II, III and IV (Fig.4). Three of the isolates (183, 187 and 98) arranged in a subcluster of 95% with *Gibberella fujikuroi* within the first cluster (I). Close related to these species are *F. proliferatum* and a subcluster (100%) comprised by *F. mangiferae* and the isolate 243. Furthermore, cluster II shows a close relation between the isolate 66 and *F. incarnatum*, which appears as a subcluster that is moreover related to isolate 272. The third cluster (III) indicates a high affinity of the isolates 16 and 214 to *F. lateritium* evidenced by a bootstrap of 94%. Finally the fourth cluster (IV) demonstrates a relative close relation of *N. haematococca* to a subcluster constituted by the isolates 221 and 234, which show high affinity (bootstrap of 100%) to *N. rigidiuscula*.

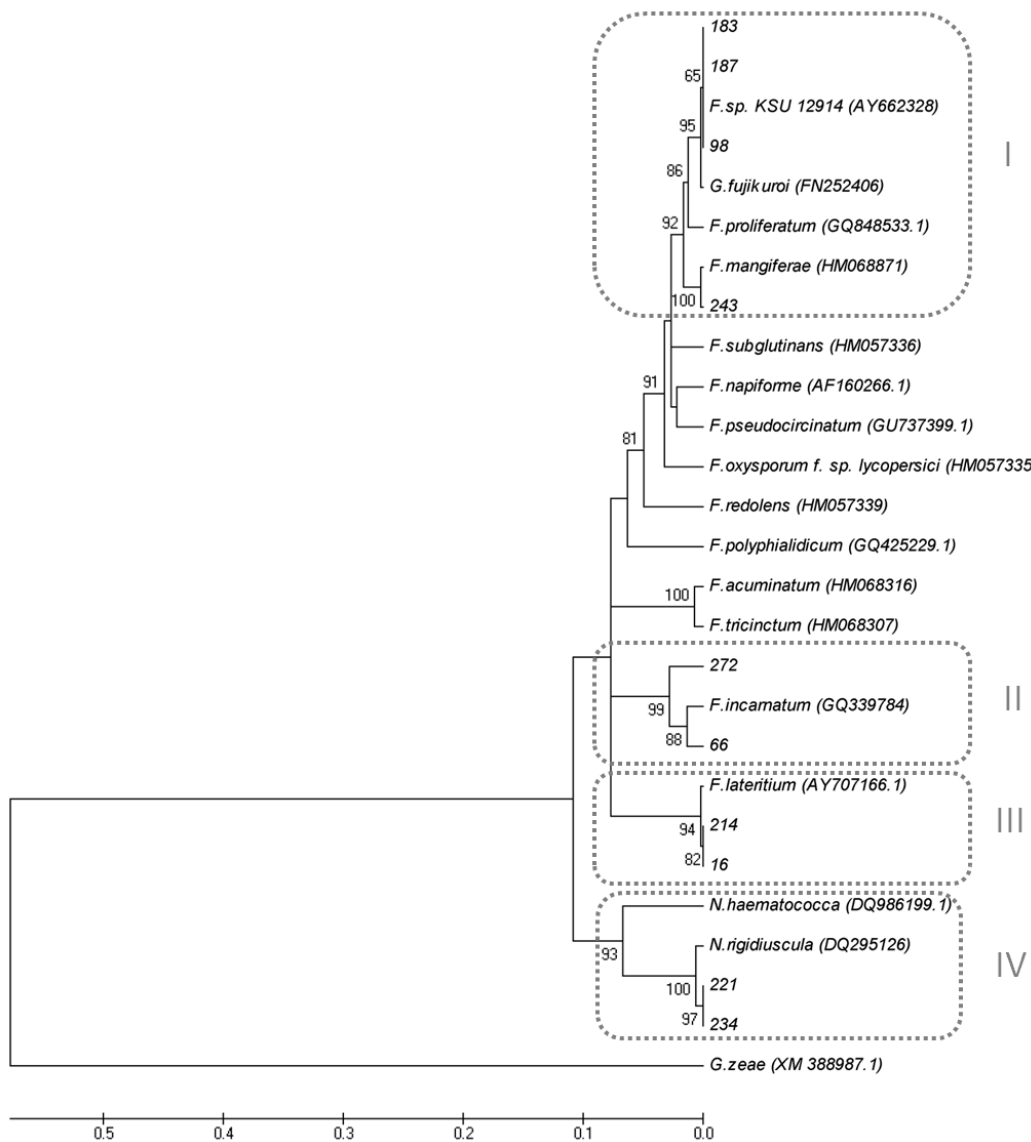


Figure 4 Phylogenetic analysis of 27 *Fusarium* spp.. The evolutionary history was inferred using the Neighbor-Joining method. The optimal tree with the sum of branch length = 1.91101542 is shown. The percentage of replicate trees in which the associated taxa clustered together in the bootstrap test (1000 replicates) is shown next to the branches. The phylogenetic tree was linearized assuming equal evolutionary rates in all lineages. The evolutionary distances were computed using the Maximum Composite Likelihood method and are in the units of the number of base substitutions per site. All positions containing gaps and missing data were eliminated from the dataset. This phylogenetic analysis was conducted in MEGA4.

Discussion

Phylogenetic identification

The first phylogenetic analysis using the LSU sequences of 172 morphospecies confirmed the affiliation of the isolates to the genus *Fusarium*. In addition, it was possible to group the isolates into four main clusters, indicating the presence of different species. Genetic variability in the LSU gene and the insufficient number of published sequences didn't allow sufficient discrimination among our isolates. Further resolution in the identification of the isolates was made possible using TEF-1alpha in the second cluster analysis. This more detailed analysis revealed a non negligible diversity of cacao associated *Fusarium* species in Sulawesi.

The first TEF-1alpha cluster (I) shows that three *Fusarium* isolates (98, 183 and 187) probably (bootstrap 65%) belong to the same species, *Gibberella fujikuroi* (telomorph *F. fujikuroi*). All three isolates are found in cluster D of the LSU-phylogram, though in different groups. *G. fujikuroi* is a known phytopathogenic fungus that overloads its host with gibberellins causing bakanae, seedling rot and grain sterility especially on rice, barley, sugarcane, millet, but also some tropical trees. *G. fujikuroi* is known to be part of the endophytic community in cacao (Rubini et al., 2005). In the same cluster (I), the isolate 243 could be identified as *F. mangiferae*. This species is widely distributed and causes the mango malformation in the tropics (Marasas et al. 2006, Ploetz 2006). The isolate 243 was part of the cluster B in the LSU-phylogram, where it is clearly distinct from three other isolates (bootstrap of 98%). In the same or at least close related to *F. mangiferae* cluster was supposed to be also isolate 221, since this *Fusarium* morphospecies was found in cluster B in the LSU-phylogram as well. Instead, there seems to be a relation between this last isolate and *N. rigidiuscula* and the isolate 234 from cluster D-group 1 (cluster IV, Fig. 4). *N. rigidiuscula* (telomorph *F. decemcellulare*) is known to be a pathogen of tropical trees (Lombard et al 2007) and was sampled in Indonesia before (Ali et al. 1998), but it never played a crucial role as a cacao pathogen (Ploetz 2007). Furthermore, high similarities in the TEF 1α cluster analysis were found between the isolate 214, which was not within the main clusters in the LSU-phylogram, and the isolate 16 from cluster D (LSU). Both isolates could be identified as *F. lateritium* considering their position and bootstrap values in the TEF 1alpha tree. To the best of our knowledge, this is the first report which describes *F. lateritium* in cacao (*Theobroma cacao*). This observation is nevertheless consistent with a previous report by Booth (1981) stating that strains of *F. lateritium* have a worldwide distribution and a wide host range. Recent changes in the classification *F. lateritium* group nevertheless complicate host range and distribution predictions (Kvas et al. 2009). Finally, it was possible to confine the identification of the isolates 66 and 272 by the analysis of the TEF-1alpha to the species *F. incarnatum*. The fungi of the species *F. incarnatum* (syn. *F. pallidoroseum*, *F.*

semitecum) have a widespread host range (e.g. Srenivasa et al. 2006, Belisario et al. 2010). Strains of this species, known for their micotoxin production capacity, have been linked to several plant diseases (Knight et al. 1977, Jiménez et al. 1993, Ploetz 2005, Zaccardelli et al. 2006) such as banana crown rot in tropical regions (Knight et al. 2008). Bananas (*Musa x paradisiaca*) are commonly grown in home gardens and as shade trees within cacao plantation (Kehlenbeck & Maass 2004). Therefore it seems plausible that both isolates 66 and 272 which are related to *F. incarnatum* infested cacao plants as endophytes. Some strains of the *F. incarnatum* group showed entomopathogenic potential against mosquitoes and could therefore be suitable as bio control agents (Mohanty et al. 2008). The pathogenicity of these cultivars was not tested, but the high diversity of bioactive endophytic *Fusarium* spp. isolated from cacao plants in this area increases the likelihood of the occurrence of potential bio control agents.

Regional and temporal distribution

Even though our study was limited in time and space, and thus did not cover *Fusarium* spp. occurrence exhaustively, it clearly revealed distribution patterns in *Fusarium* spp.. *Fusarium* distribution was significantly influenced by geographical and temporal differences (Fig. 2). According to the analysis, Isolates of the LSU D1 group formed a dominant cluster in all sites, except in the Marena plantation where the drought experiment was conducted. The Marena plantation has been managed more intensively in the past compared to the other plantations (Schmidt & Vidal unpublished). Contrary to the other plantations the cacao plants in Marena have been fertilized annually. An additional nitrogen source could have influenced the competitiveness of single strains or species of *Fusarium* (Candau et al. 1992) and affect the diversity and abundances of the *Fusarium* spp.. Because the *Fusarium* diversity of the Kulawi valley was consistent with the *Fusarium* diversity sampled in 2007 Marena plantation before the drought was conducted, an influence due to plantations history can be excluded. Indeed there is a considerable difference between both sampling periods in the Marena plantation (Fig. 2), suggesting the geographical pattern might have been masked by temporal pattern. During the drought experiment in the Marena plantation the same shift in dominant species diversity observed in the treatment plots also occurred in the control plots (Schmidt & Vidal unpublished). A dry spell, that occurred at once before the second sampling period in 2008 was held responsible for the shift in abundances of the dominant species on the one hand. On the other hand the induced drought stress of the host plants decreased the endophyte composition significantly (Schmidt & Vidal unpublished). The drought might increase the evolutionary pressure on *Fusarium* spp., rebounding to the advantage of *Fusarium* spp. in the LSU D2 group, which might have benefited from the lack of precipitation (Suryanarayanan et al. 2002). Beside precipitation, other environmental factors were also affecting the distribution of LSU *Fusarium* clusters.

Environmental gradients

The distribution of fungal endophytes in general (Arnold 2007, Arnold & Herre 2003) and *Fusarium spp.* in particular (Blackhouse & Burgess 2002, Summerell et al. 2003, Vujanovic et al. 2006) are influenced by temperature. In this investigation *Fusarium spp.* were distributed along a local temperature gradient (Fig. 3). The minimal annual temperature correlated positively with the abundance per group, which was caused mainly by the lower abundance of the group LSU D1 in plantations with low temperatures (Tab. 1). Reasons for low abundances in this group might be: 1) some species or strains were sensitive to lower temperatures. 2) Some species or strains were limited in dispersal to host plants, which are only planted in lower altitudes and therefore higher temperatures (Tab. 1). 3) Or most likely, the main group of this cluster had (similar to its relative *G. fujikuroi*) optimal growing conditions from 29 to 32°C (Borrow et al. 1964), which would result in a decrease of abundances in temperatures below 25°C. Further evidence was given from the Marena drought experiment, where *Fusarium spp.* that were clustering mainly within group LSU D2 were most abundant after a drought spill. Though the air temperature in the shade was remaining constant (Köhler et al. 2009), it is likely that the temperature on the canopy surface might have increased. Further correlations between altitude over sea level and general *Fusarium* abundance were most likely caused by the link between altitude and temperature. Except cacao plantations with a slope, that may have higher mean temperatures as a result of increased solar exposition (Schmidt et al. unpublished).

Fusarium, its influence on cacao yield and pathogenicity

The results of this investigation showed that the *Fusarium* diversity is strong correlated to the cacao yield (Tab. 1), suggesting that some of the isolated *Fusarium spp.* could increase the yield. As discussed before, *G. fujikuroi* is known as causer of the Kabake disease in rice, but also as an industrial producer of gibberellins (Phinney 1983). Some gibberellins produced by bacteria have been shown to promote plant growth and increase the yield of many crop plants (Okon & Labandera-González 1994, Bottini et al. 2004). In our study after the impact of drought, cacao plantations displayed an increased abundance of *Fusarium* group LSU D2, even though a decrease in yield was recorded. Because of the inhibition of gibberellins production in *G. fujikuroi* by nitrogen (Candau et al. 1992) and high nitrogen availability in the soil of the rainforest margins (Siebert 2002, Schwendenmann et al. 2010) it is very unlikely that gibberellins were produced and accounted for the increase yield. Yield increases by biocontrol activities of *Fusarium spp.* against insect pest, like expressed by some *F. oxysporum* strains (Barahona 2010), was excluded by the strong positively correlation of *Fusarium* diversity with the cacao pod borer (*Conopomorpha cramerella*, Tab. 1). An explanation for this correlation could be an increased *Fusarium* infestation of the cacao plant resulting from additional plant wounds by the cacao pod borer. Most reported wounds by the

cacao pod borer are caused by the larvae tunneling cacao shells (Day 1989). The resting sites of the adults were found to be located at the undersides of branches (Day et al. 1994), suggesting occasional feeding on leaves, which could support the infestation of leaves by *Fusarium spp.*. But the most plausible hypothesis might be that the infestation rates of the cacao pod borer and *Fusarium spp.* follow the same temperature gradient as the cacao yield. Surprisingly, independently of the *Fusarium* infestation rates, no symptoms of *Fusarium* induced diseases had been observed in the cacao plantations during our investigation.

Alternative host trees

The number of observed fungal species in a habitat depends on the diversity of plant taxa (Hyde 2001, Cannon & Simmons 2002). In this investigation, the neighbourhood of seven tree species were shown to have an influence on the distribution of endophytic *Fusarium spp.* in cacao plants (Tab. 2). The occurrence of the intercrops *Coffea* and *Mangiferae* in cacao plantations correlated with *Fusarium* diversity, explaining the isolation of typical associated *Fusarium* species. Both genera are known to host various *Fusarium spp.* (Ploetz 2006, Vega et al. 2009) and both genera have been commonly grown in cacao plantations in Sulawesi (Kehlenbeck & Maass 2004). Additionally the most common planted shade tree *G. sepium* (Clough et al.) correlated positively with the abundance of group LSU D2. All cacao trees of plantations with numerous *G. sepium* stands showed high abundances of group LSU D2, except the cacao plantation with the highest *G. sepium* density, where *Fusarium spp.* of group LSU D2 were missing. The absence of one group might have been caused by the random selection of isolates for DNA identification, even though it is unlikely as it concerned only one group. Interestingly group LSU D1 correlated in the same plantation negatively with *P. anduncum*, which occurred rarely, except in this plantation where 23 individuals occurred. There is no literature evidence that *P. anduncum* might produce antifungal agents active against *Fusarium* (specifically of group LSU D2). In the contrary, *T. ciliata*, which was also negatively correlated with *Fusarium* group LSU D2, has previously been proven to contain antifungal metabolites (Govindachari et al. 2000, Chowdhury et al. 2003). Trees like *T. ciliata* with antifungal activities, might indeed reduce the endophyte composition in an area by three mechanisms 1) occupying the stand of an endophyte rich tree, or 2) by direct effects on the fungal diversity of a canopy located below the tree 3) or effects on the fungal community of the leaf litter below, as antifungal compounds remain temporary preserved in leaf litter (Chapuis-Lardy et al. 2002). Changes in the fungal leaf litter community due to different in susceptibility to antifungal compounds could change the ejected spore diversity and therefore affect the endophyte composition within cacao leaves (Herre et al. 2007, Suryanarayanan et al. 2009).

Conclusion

In the present investigation we have shown how environmental gradients, such as temperature, drought or habitat plant composition, change the *Fusarium* composition within host plants and influence the distribution of phylogenetically distinct *Fusarium* groups. We conclude that the variety and the diverse combinations of distinct drivers makes tropical plant communities a hotspot for new endophytic, antagonistic or pathogenic *Fusarium* species and strains. The results of this investigation contribute to the knowledge about the development of *Fusarium* diversity and ecology.

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General Discussion

Endophytes: existence in a tight spot

Fungal endophytes are one part of giant fungal continuum, which contains: soil, leaf litter and water fungi, as well as fungal epiphytes, phellophytes, mycorrhiza or fungi associated with plants, lichens, and animals (Hawksworth 1991). Endophytes are neither a phylogenetic group nor consistent in their definition. Fungi or bacteria are termed endophytes based on their lifecycle characteristic. For the definition it is sufficient if the organism occurs in living tissues of plants in some of his developmental stages without causing negative affects (Wilson 1995). Besides there are pathogens that occur endophytic in other host plants and endophytes that are regarded as latent pathogens, with the capability to change their life cycles (Müller & Kraus 2005, Schulz & Boyle 2005, Kogel et al. 2006, Arnold & Engelbrecht 2007, Arnold 2007). Hence, the term endophyte is independent of the phylogenetic relationships among fungi and the kind of detriment free interactions that might occur between fungi and the host plants. Based on several studies that have shown the important role of the ubiquitous fungal endophytes in influencing the diversity, structure and dynamics of plant communities, fungal endophytes can be used as models investigating plant communities and their ecology (Cubit 1974). The close association to their host frees fungal endophytes of nearly all artificial selection pressure. This qualifies them as ideal model subjects for population biology analyses (Petrini 1998). Not least fungal endophytes can be cultivated on common malt extract or potato dextrose media, with the exception of most of the biotrophic endophytes, this makes them easily accessible for scientific investigations. In this study the foliar fungal endophyte community was investigated, because the foliage comprises the biggest surface of the tree and therefore reflects the diversity of fungal spores that have been ejected into the air surrounding the trees. Consequently the diversity obtained guaranties conclusions about the fungal diversity plant community present.

Variations in diversity

The amount of fungal endophytes species of a tree reported by previous studies differs from low diversities of 50 and less isolated species (e.g. Brown et al 1998, Suryanarayanan et al. 2002, Rubini et al. 2005) up to a high diversity of 200 and more species (e.g. Arnold et al. 2000, Vega et al. 2009b). Thereby, the total number of species depends on the sample size, the tree species, the tree's environment and the endophyte classification method. Comparing rare faction analysis, which normalize sampling size and estimates the total number of species, the number of endophyte species for a tree in temperate zones is limited to about 30 species (Unterseher et al. 2007, Hoffmann & Arnold 2008). Interestingly, the estimated

fungal endophyte diversity of algae is also below 30 species with the exception of brown algae which are expected to host more than 50 species. Host trees of tropical areas were proven to host more than several hundred of fungal endophytes (e.g. Arnold et al. 2000, Arnold et al. 2001, Vega et al. 2009b). According to the species estimation of Arnold et al. (2009) this high fungal endophyte diversity of tropical trees is comparable to the high fungal endophyte diversity of tropical lichens, suggesting that fungal endophyte diversity follows climatic conditions (Suryanarayanan et al. 2002, Hoffman & Arnold 2007). Thus the same plant species in comparable climatic zones with a comparably diverse plant community should host a comparable amount of fungal endophytes. The results of the present study showed a decreased in diversity of endophytic fungi inhabiting cacao plants in Indonesia compared to Latin America. On one hand this suggests the high diversity of fungal endophytes Arnold et al. (2001) isolated from cacao that cacao plants are highly susceptible to fungal endophytes. On the other hand several studies showed some evidence for host preferences of endophytes to other host plants (Arnold et al 2000). This argument could explain the reduced diversity in Indonesia by the insusceptibility of cacao for the Indonesian fungal community, combined with the loss of cacao adapted fungal endophytes during the introduction of cacao in Indonesia 25 years ago. Conversely found Cannon & Simmons (2002) no clear evidence for host preferences in Guyana, which leads to the suggestion that the fungal endophyte diversity in Indonesia might be decreased altogether. The molecular identification of the second most abundant taxa (*Fusarium spp.*) in the fungal endophyte community sampled in Indonesia revealed that the real diversity of species and strains might be much higher. Since the fungal taxa as well as the molecular differenced groups follows several environmental gradients, this provides various opportunities for speciation by isolation and evolutionary pressure (Wuenschel 1969, Dynesius & Jansson 2000).

Environmental drivers of diversity and host preferences

The endophyte communities within the same host plant species are not the same, if they are located in different sites (Arnold 2000). For example, Arnold (2003) showed a continuously decreasing similarity of fungal endophyte communities in cacao plants with increasing distance between the locations of the communities compared. Other studies found differences in plant associated fungi of different areas respective to climatic differences (Blackhause & Burgess 2002). Suryanarayanan et al. (2002) compared endophytes of a tropical dry thorn, a dry deciduous, a moist deciduous and a semi-evergreen forest resulting in a fungal diversity gradient correlating with precipitation rates. Contrary to that was the fungal endophyte composition of cacao plants shown in the present study which changed gradually under the impact several opposed gradients (Schmidt et al. unpublished). For example was the similarity of fungal endophyte composition influenced by the minimal

temperature, which was dependent on the altitude (Schmidt et al. unpublished). This was reflected by the abundance of morphologically distinguished *Fusarium* groups, which was more abundant in lower altitudes and therefore warmer cacao plantations. Additionally the degree of shade, or rather the intensity of exposure of solar radiation of the cacao canopy showed an impact on the fungal diversity (Schmidt et al. unpublished). Besides temperature and intensity of solar radiation, the fungal endophyte community was heavily influenced by the surrounding plant community. Similarities in the plant community of a cacao plantation resulted in very similar cacao inhabiting fungal endophyte compositions. Further correlations between molecularly distinguished *Fusarium* groups and single shade tree species planted in between cacao plants outline the interactions of fungal communities in a habitat (Schmidt et al. unpublished). These correlations between fungal endophytes and tree species, give clear evidence for host preferences within the fungal endophyte community (Arnold et al. 2000). Further support comes from drought experiment, where the fungal endophyte diversity was decreased, while the host plant was exposed to an artificial drought (Schmidt & Vidal unpublished). Because the cacao canopy was exposed during the experiment to the same precipitation and humidity as the control, it was suggested that the fungal endophyte composition reacted to plants drought stress induced secondary metabolites (Bartels & Sunkar 2005). If plants secondary metabolites can alter the fungal endophyte community (Saunders & Kohn 2009), host preferences will be established. In this study, the influence of plant community on the fungal endophyte community showed that fungal endophyte assemblages of a host tree species can not be seen as being independent. The interactions between the fungal communities in the foliage and decaying leaf litter (Herre et al. 2007, Suryanarayanan et al. 2009) connects plant communities in dependency of the leaf fall range and spore distribution by wind. This characteristic of fungal endophytes communities qualifies them as ideal models in ecological research and as indicators of stable ecological systems.

Climate Change

One aspect of the climate change is the increased frequency and duration of El Niño Southern Oscillation (ENSO) events (IPCC 2001), tropical plant communities in Asia are affected by more frequent and more severe ENSO related drought events (Sheffield & Wood 2008). As fungal distributions have been shown to be influenced by minimal temperature, climate change concerns fungal endophyte communities and therefore also all kinds of host plant endophyte relations. In this study an artificial drought was conducted independent of temperature, which allowed the investigation of fungal endophyte community in regards to the plants drought stress reactions. The decreased fungal endophyte diversity as well as higher similarity of fungal endophyte composition between cacao trees under the impact of

drought compared to control, suggest a disturbance of interactions between host plant endophytes. While most of the fungal endophytes were affected by the changes within the host plant conditions, one endophyte taxa took benefit from the new situation. Molecular methods confined the characterization of this taxa to a group in close relation to *Gibberella fujikuroi*, the pathogen causes of kabake disease in rice. The increased of abundance in this group of *G. fujikuroi* could be seen as 1) competitive success of a latent pathogen, producing antifungal metabolites (Müller & Kraus 2005. Schulz & Boyle 2005), 2) benefit of a species resistant to secondary metabolites, as *F. culmorum* protects of reactive oxygen species conferring salt tolerance (Rodriguez et al. 2008), or 3) benefit of a species conferring resistance against drought e.g. by increasing the water use efficiency (Rodriguez et al. 2008). Anyhow, the decreased fungal endophyte diversity is alarming, because a reduced fungal community will support pathogens (see Arnold et al. 2003). Future ENSO related drought events might decrease the endophyte diversity of South-East Asia even more rapidly (Thomas et al. 2004). Further investigations of fungal endophyte diversity and pathogen interactions respectively climate change, could help creating fungal pathogen prediction models and agricultural management models.

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Publications

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Poster

Schmidt C, Clough Y, Vidal S, (2008) Diversity of endophytic fungi in *Theobroma cacao* leaves in Central Sulawesi, Tropical Rainforest and Agroforests under Global Change, Bali, Indonesia

Schmidt C, Vidal S (2009) Distribution pattern of foliar fungal endophytic fungi in *Theobroma cacao* in Central Sulawesi, German Society for General and Applied Entomology (DGAAE), Goettingen, Germany

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Eidesstattliche Erklärung

Hiermit erkläre ich eidesstattlich, dass diese Dissertation selbständig und ohne unerlaubte Hilfe angefertigt wurde

Göttingen, den 30. September 2010