Distinct spatiotemporal patterns between fungal alpha and beta diversity of soil-plant continuum in rubber tree

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The following supporting information is available for this article:

Method S1

Sampling

In each sampling plot, three rubber trees, with a distance about 100 meters from each other, were selected as our study objectives. Top soil (0-20 cm) samples were collected from at the middle of two rubber trees and the center of three rubber trees (Figure S1B). After sieving, they were mixed to form a composite soil sample. Root samples were selected from the four directions (i.e., north, south, east, and west) of each rubber tree, then mixed to form a single composite sample (Figure S1B). Rhizosphere soil was manually separated from the roots by shaking, while leaving rhizoplane soil (~1 mm thick layer of soil) still attached to the roots (1-3). Leaves were collected in the same time from branches 2 m away from the center of the trunk and 12 m above the ground (Figure S1B). Leaves were selected from each of the four cardinal directions and all sampled leaves from the three trees were mixed for a single composite sample.

A total of 36 leaf samples, 36 soil samples and 36 root samples were collected in the two seasons (rainy season and dry season), and all samples were transported to the laboratory on ice and placed in sterile PBS solution to isolate epiphytic and endophytic microbes within 24h.

To collect the rhizoplane soil sample, place 5 g of roots root in a 50 ml sterile centrifuge tube containing 35 mL of sterile PBS and subjected to sonication using a sonication bath for 1 min, then shaken at 200 rpm for 4 min, this process is repeated 2-3 times. The buffer with microbial cells is then passed through a 0.22 μ m filter membrane, and the resulting filter membrane was stored in a sterile centrifuge tube at -80 °C in preparation for microbial sequencing analyses. To collect the phyllosphere

sample, place 15 g of leaves in a 250 ml sterile glass vial containing 200 mL of sterile PBS and subjected to sonication using a sonication bath for 1 min, then shaken at 200 rpm for 4 min, this process is repeated 2-3 times. The buffer with microbial cells is then passed through a $0.22~\mu m$ filter membrane, and the resulting filter membrane was stored in a sterile centrifuge tube at -80° C in preparation for microbial sequencing analyses.

For leaf and root endophytic sample, most of the microbial cells on the plant surface were removed by treatment as described above, and the leaves or roots obtained were washed with sterile water for the next step of surface disinfection. The main steps are: soak in 70% alcohol for 5 minutes, then in 5.25% sodium hypochlorite solution for 5 minutes, then in 70% alcohol for 30 seconds, and finally wash with sterile water at least 5 times to avoid alcohol residue. The leaves and roots samples were stored at -80 °C in preparation for microbial sequencing analyses.

Therefore, we obtained 216 samples, including those from the phyllosphere (36 samples), leaf endosphere (36 samples), soil (36 samples), rhizosphere (36 samples), rhizosphere (36 samples), and root endosphere (36 samples) compartments.

For leaf sample, TN was determined using micro-Kjeldahl digestion followed by steam distillation. TP and TK were assessed using NaOH digestion. WC content and LOM content were assessed using the gravimetric method. Leaf pH was measured in a leaf/water suspension (1:2.5, w/w) using a pH meter. Soil pH was measured in a 1:1 soil: water mixture. Soil moisture was measured gravimetrically. Soil TN was determined using a micro-Kjeldahl digestion followed by steam distillation. Total phosphorus (TP) and total potassium (TK) were digested with NaOH. Nitrate nitrogen (NN) and ammonium nitrogen (AN) were determined by steam distillation and indophenol-blue colorimetry, respectively. Soil samples were extracted with NaHCO3, and the extracts then used to measure the available soil phosphorus (AP) using molybdate-blue colorimetry. For soil potassium (AK), soil samples were first extracted with ammonium acetate, before loading the extracts onto an atomic absorption spectrometer with ascorbic acid as a reductant (4)

Supplementary figures

Figure S1 Study sites on Hainan Island and Xishuangbanna. Red solid circles indicate study sites on the maps. Each site was further divided into 3 plots, totaling 18 sampling plots.

Figure S2 Spatiotemporal changes in fungal community composition (class level) in different compartments of rubber trees.

Figure S3 Environmental variables in dry and rainy seasons between two locations of rubber tree leaves. The significant differences between seasons and locations were detected by two-way ANOVA. N: Total nitrogen, P: Total phosphorus, K: Total potassium, WC: Water content, LOM: Leaf organic matter.

Figure S4 Environmental variables in dry and rainy seasons among the six sampling sites of rubber tree leaves. The significant differences between seasons and locations were detected by two-way ANOVA. WC: Water content, SOM: Soil organic matter, pH: Soil pH, AN: Ammonium nitrogen, NN: Nitrate nitrogen AP: Available phosphorus, AK: Available potassium, TN: Total nitrogen, TP: Total phosphorus, TK: Total potassium.

Figure S5 Boxplots showing the number of OTUs between the two locations (a) and seasons (b). Leaf.WC: Leaf water content.

Figure S6 Shannon (a) and evenness (b) diversity of rubber tree fungal community.

Figure S7 Boxplots showing the number of OTUs between the two locations (a) and seasons (b).

Figure S7 Principal coordinates analysis (PCoA) of taxonomic similarity based on Bray-Curtis distances (OTU level). Six compartments in Hainan (a) and Banna (b). LD: Ledong, DZ: Danzhou, WN: Wanning; MP: Mengpeng, JH: Jinghong, ML: Menglun.

Supplementary tables

Table S1 The OTUs table.

Table S2 The significant effect of location and season on fungal alpha diversity was detected by two-way ANOVA.

Table S3 Mantel test results for the correlation between community similarity and environmental.

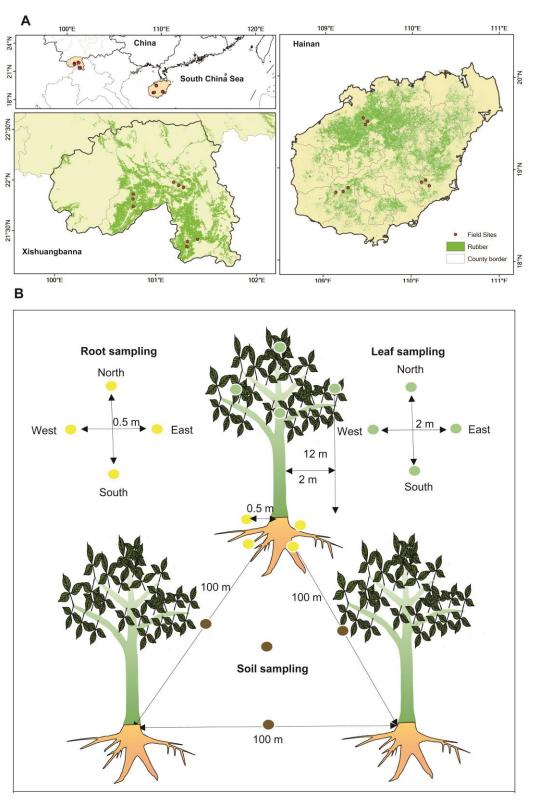


Figure S1 Study sites (A) and sampling design (B) on Hainan Island and in Xishuangbanna. Red solid circles indicate study sites on the maps. Each site was further divided into 3 plots, totaling 18 sampling plots. For each plot, we selected three trees and sampled root (yellow solid circles) and leaves (green solid circles) from all four cardinal directions. The brown solid circles indicate sampling locations of soil.

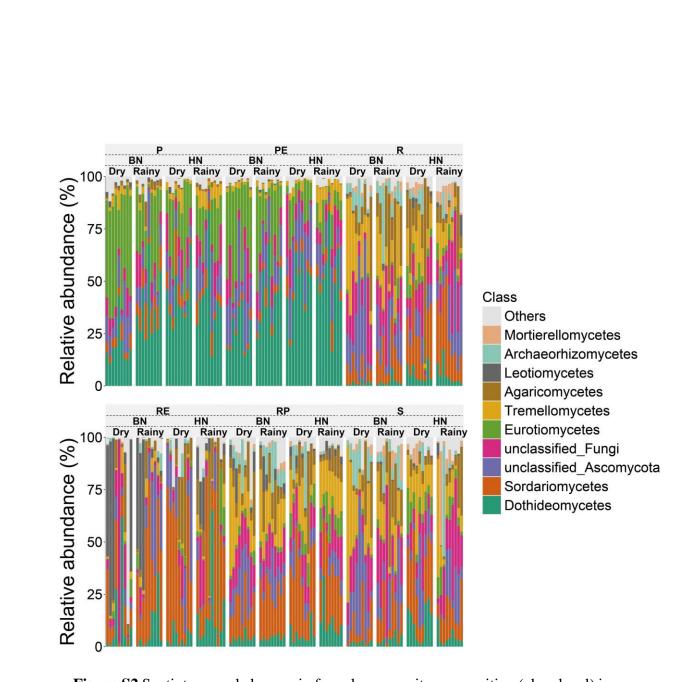


Figure S2 Spatiotemporal changes in fungal community composition (class level) in different compartments of rubber trees.

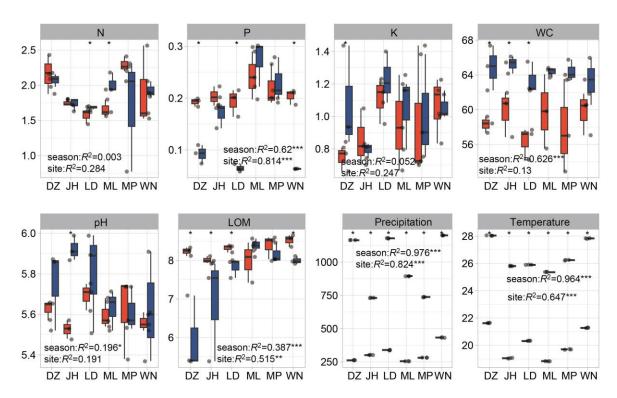


Figure S3 Environmental variables in dry and rainy seasons among the six sampling sites of rubber tree leaves. The significant differences between seasons and locations were detected by two-way ANOVA. N: Total nitrogen, P: Total phosphorus, K: Total potassium, WC: Water content, LOM: Leaf organic matter. *, P < 0.05; **, P < 0.01, ***, P < 0.001

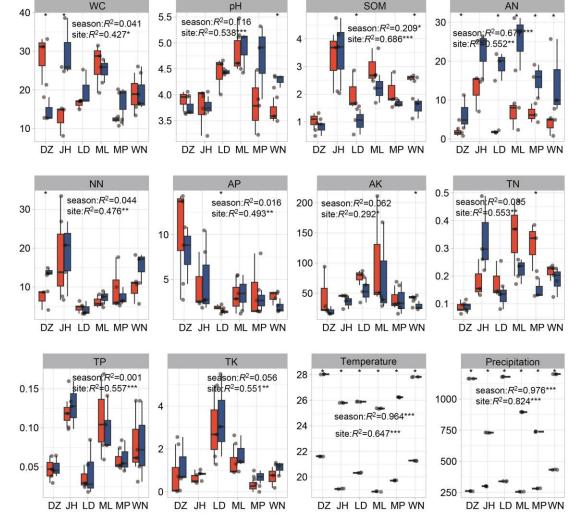


Figure S4 Environmental variables in dry and rainy seasons among the six sampling sites of rubber tree soils. The significant differences between seasons and locations were detected by two-way ANOVA. WC: Water content, SOM: Soil organic matter, pH: Soil pH, AN: Ammonium nitrogen, NN: Nitrate nitrogen AP: Available phosphorus, AK: Available potassium, TN: Total nitrogen, TP: Total phosphorus, TK: Total potassium. *, P < 0.05; **, P < 0.01, ***, P < 0.001

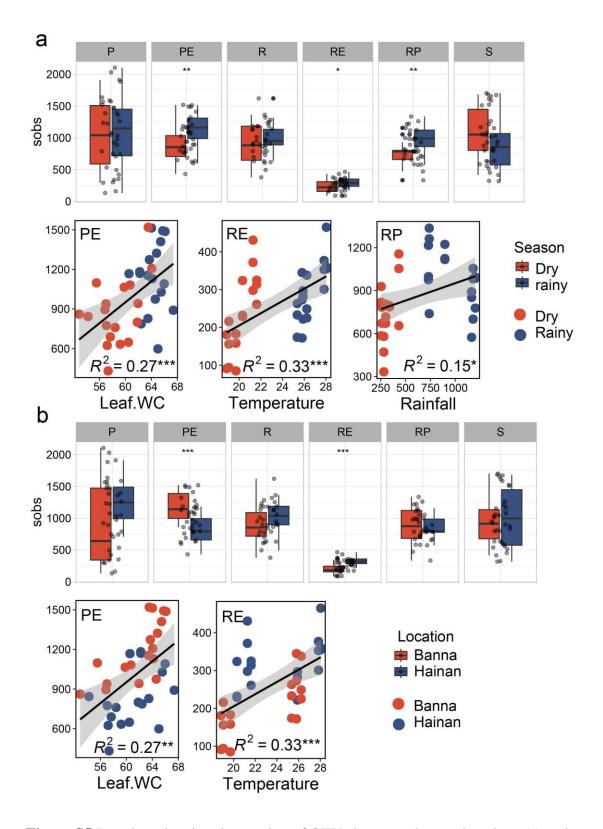


Figure S5 Boxplots showing the number of OTUs between the two locations (a) and seasons (b). ***P < 0.001; **P < 0.01; *P < 0.05. Leaf.WC: Leaf water content.

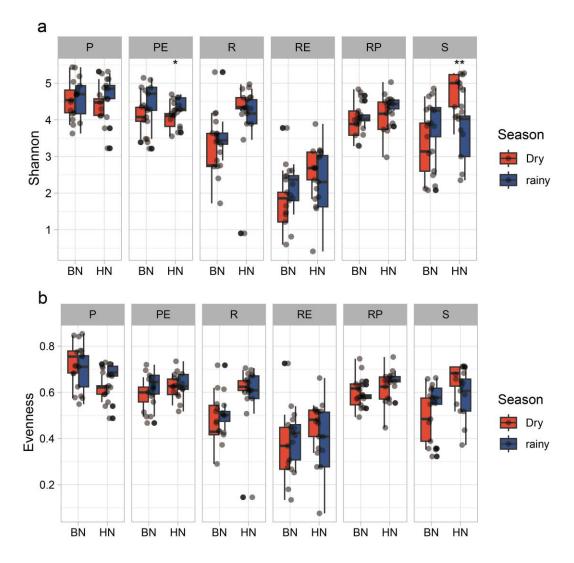


Figure S6 Shannon (a) and evenness (b) diversity of rubber tree fungal community. Significant differences between seasons in each sampling area are marked by stars. **P < 0.01; *P < 0.05.

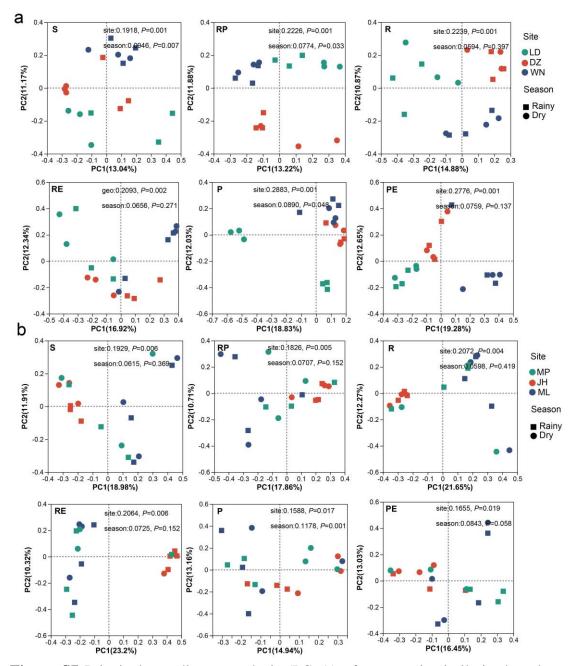


Figure S7 Principal coordinates analysis (PCoA) of taxonomic similarity based on Bray-Curtis distances (OTU level). Six compartments in Hainan (a) and Banna (b). LD: Ledong, DZ: Danzhou, WN: Wanning; MP: Mengpeng, JH: Jinghong, ML: Menglun. ***P < 0.001; **P < 0.01; *P < 0.05

Table S2 The significant effect of location and season on fungal alpha diversity was detected by two-way ANOVA.

		Df		Sum Sq	Н	Pr(>F)
Phyllosphere						
	Location		1	373.78	3.3674	0.066
	Season		1	4.00	0.0360	0.849
	Location: Season		1	625.00	5.6306	0.017
	Residuals		32	2882.22		
Leaf						
endosphere						
	Location		1	1573.4	14.1752	0.00017
	Season		1	841.0	7.5766	0.00591
	Location: Season		1	25.0	0.2252	0.63509
	Residuals		32	1445.6		
Soil						
	Location		1	9.00	0.0811	0.77582
	Season		1	205.44	1.8511	0.17366
	Location: Season		1	2146.78	19.3428	0.00001
	Residuals		32	1523.28		
Rhizosphere						
	Location		1	342.25	3.0841	0.07906
	Season		1	75.11	0.6769	0.41067
	Location: Season		1	1284.03	11.5708	0.00067
	Residuals		32	2182.61		
Rrhizoplane						
	Location		1	26.69	0.2405	0.62383
	Season		1	1024.00	9.2264	0.00239
	Location: Season		1	633.36	5.7067	0.01690
	Residuals		32	2200.44		
Root						
endosphere						
	Location		1	1736.11	15.6427	0.000077
	Season		1	484.00	4.3609	0.036772
	Location: Season		1	177.78	1.6018	0.205647
	Residuals		32	1486.61		

Table S3 Mantel test results for the correlation between community similarity and environmental. WC: Water content, LOM: Leaf organic matter, SOM: Soil organic matter, pH: Soil pH, AN: Ammonium nitrogen, NN: Nitrate nitrogen AP: Available phosphorus, AK: Available potassium, TN: Total nitrogen, TP: Total phosphorus, TK: Total potassium.

Compartment	Factor	r	p
	Leaf.P	0.254	0.001
	Temperature	0.200	0.001
	Rainfall	0.190	0.001
DI 11 1	Leaf.K	0.045	0.193
Phyllosphere	WC	0.041	0.217
	Leaf.N	-0.039	0.717
	pН	-0.044	0.756
	LOM	-0.125	0.955
	Leaf.P	0.167	0.001
	Rainfall	0.140	0.009
	Temperature	0.126	0.008
If 11	WC	0.053	0.148
Leaf endosphere	Leaf.K	0.051	0.135
	pН	-0.027	0.667
	LOM	-0.066	0.81
	Leaf.N	-0.101	0.956
	AK	0.312	0.005
	Rainfall	0.225	0.001
	pН	0.195	0.019
	TN	0.127	0.12
	Temperature	0.121	0.008
Soil	TK	0.108	0.162
5011	AN	0.106	0.092
	TP	0.051	0.273
	AP	0.015	0.419
	SOM	0.004	0.461
	WC	-0.078	0.823
	NN	-0.103	0.83
	AK	0.313	0.001
	pН	0.261	0.003
	TK	0.191	0.043
Rhizosphere	TN	0.125	0.109
Killzosphere	Rainfall	0.105	0.032
	TP	0.090	0.148
	AN	0.067	0.179
	Temperature	0.022	0.301

NN -0.019 0.531 AP -0.043 0.631 WC -0.079 0.824 AK 0.329 0.002 pH 0.293 0.001 TN 0.271 0.006 WC 0.109 0.079 Rainfall 0.108 0.034 Temperature 0.100 0.023 TK 0.093 0.189 TP 0.091 0.13 SOM 0.076 0.19 AP -0.011 0.519 NN -0.060 0.794 AN -0.060 0.795 AK 0.190 0.001 TN 0.177 0.001 pH 0.154 0.003 TP 0.128 0.015 TP 0.128 0.016 T				
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Rainfall 0.091 0.029 Temperature 0.085 0.035 WC 0.066 0.099 AN 0.045 0.195 TK 0.021 0.349		TP	0.128	0.015
Temperature 0.085 0.035 WC 0.066 0.099 AN 0.045 0.195 TK 0.021 0.349		SOM	0.099	0.035
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WC 0.066 0.099 AN 0.045 0.195 TK 0.021 0.349		Temperature	0.085	0.035
TK 0.021 0.349			0.066	0.099
		AN	0.045	0.195
AP 0.010 0.432		TK	0.021	0.349
		AP	0.010	0.432
NN -0.008 0.531		NN	-0.008	0.531

References

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