

Soil microbial resource limitations and community assembly along an age gradient of a subtropical planted *Camellia oleifera* forest

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Supplementary information

Table S1. Enzymes included in this study.

Enzyme	Abbreviations	EC ^a	Function
β -D-cellulobiosidase	CBH	3.2.1.91	Cellulose degradation: hydrolyses cellobiose dimers from non-reducing ends of cellulose molecules
β -1, 4-glucosidase	BG	3.2.1.21	Cellulose degradation: hydrolyses glucose from cellobiose
L-leucine aminopeptidase	LAP	3.4.11.1	Proteolysis: hydrolyses leucine and other hydrophobic amino acids from the N terminus of polypeptides
β -N-acetylglucosaminidase	NAG	3.2.1.14	Chitin and peptidoglycan degradation: hydrolyses glucosamine from chitobiose
Acid phosphatase	AP	3.1.3.2	Hydrolyses phosphate from phosphosaccharides and phospholipids

Note: ^a Enzyme commission classification (Sinsabaugh et al., 2008, 2009).

Table S2. Significant differences of soil bacterial community between stand age.

	<i>P</i>
< 9 years vs 9-20 years	0.519
< 9 years vs 21-60 years	0.005
< 9 years vs > 60 years	0.001
9-20 years vs 21-60 years	0.046
9-20 years vs > 60 years	0.008
21-60 years vs > 60 years	0.852

Table S3. Mantel tests with Spearman's rank correlations between environmental factors (Euclidean distance) and β NTI of bacterial communities with 999 permutations. SOC: soil organic carbon, TN: total nitrogen, TP: total phosphorous, Olsen-P: Olsen phosphorous. ($P < 0.05$).

Variables	ρ (Mantel test)	P
pH	0.130	0.017
SOC	-0.021	0.573
TN	-0.080	0.785
TP	0.142	0.117
Olsen-P	0.084	0.247

Table S4. Topological properties of networks at different stand ages in *C. oleifera* plantation. Different letters denote significant difference among stand ages ($P < 0.05$).

Stand age	Edges	Nodes	Transitivity (Trans)	Average Degree (AD)	Betweenness centrality (BC)	Average path Length (GD)	Density (DS)
< 9 years	245.13±63.77a	131.00±28.37a	0.37±0.04b	3.72±0.25b	0.19±0.06a	4.37±0.33a	0.03±0.01b
9 – 20 years	263.20±42.34a	143.70±19.80ab	0.35±0.05ab	3.67±0.31b	0.18±0.05a	4.54±0.28ab	0.03±0.01ab
21 – 60 years	269.80±45.71a	156.80±18.62b	0.35±0.03ab	3.43±0.28ab	0.17±0.05a	4.79±0.17b	0.02±0.00a
> 60 years	251.29±28.78a	152.57±9.64ab	0.32±0.02a	3.29±0.24a	0.19±0.049a	4.82±0.36b	0.02±0.00a

Different lowercase letters in the same column indicated the significance between different stand age ($P < 0.05$).

Table S5. Variations of soil physicochemical properties at different stand ages in *C. oleifera* plantation.

Different letters denote significant difference among stand ages.

Stand age	pH	SOC(g/Kg)	TN(g/Kg)	TP(g/Kg)	Olsen-P(mg/Kg)
< 9 years	4.39±0.20a	10.52±3.97a	1.14±0.29a	0.35±0.21a	6.54±5.78a
9 – 20 years	4.40±0.15a	17.68±5.83b	1.50±0.37ab	0.40±0.19a	5.70±3.42a
21 – 60 years	4.32±0.20a	20.71±2.67b	1.75±0.28b	0.30±0.10a	4.21±0.95a
> 60 years	4.24±0.16a	23.11±5.77b	1.85±0.50b	0.34±0.14a	5.10±1.39a

SOC, soil organic carbon; TN, total nitrogen; TP, total phosphorus; Olsen-P, Olsen phosphorous.

Different lowercase letters in the same column indicated the significance between different stand age ($P < 0.05$).

Fig. S1. P value for the KS, SSE, and NNSD tests of the exponential distribution generated by random matrix theory.

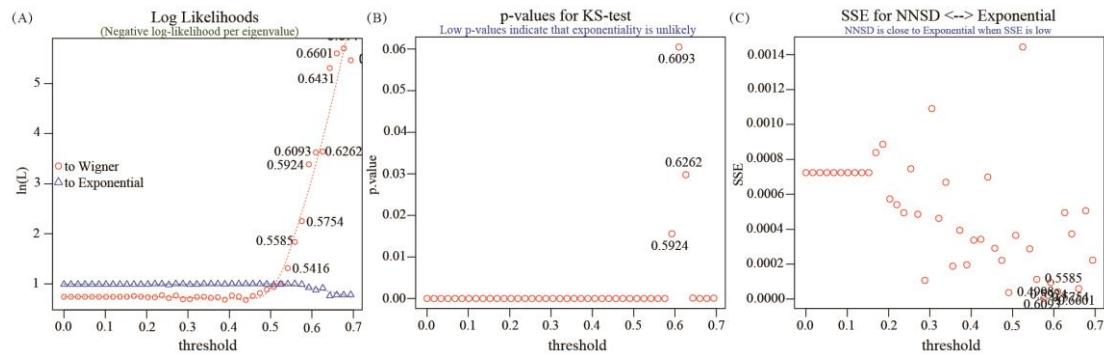


Figure S2. Variations of soil enzyme activities at different stand ages in *C. oleifera* plantation. Note: (A) CBH; (B) BG; (C) LAP; (D) NAG; (E) AP. Different letters denote significant difference among stand ages ($P < 0.05$).

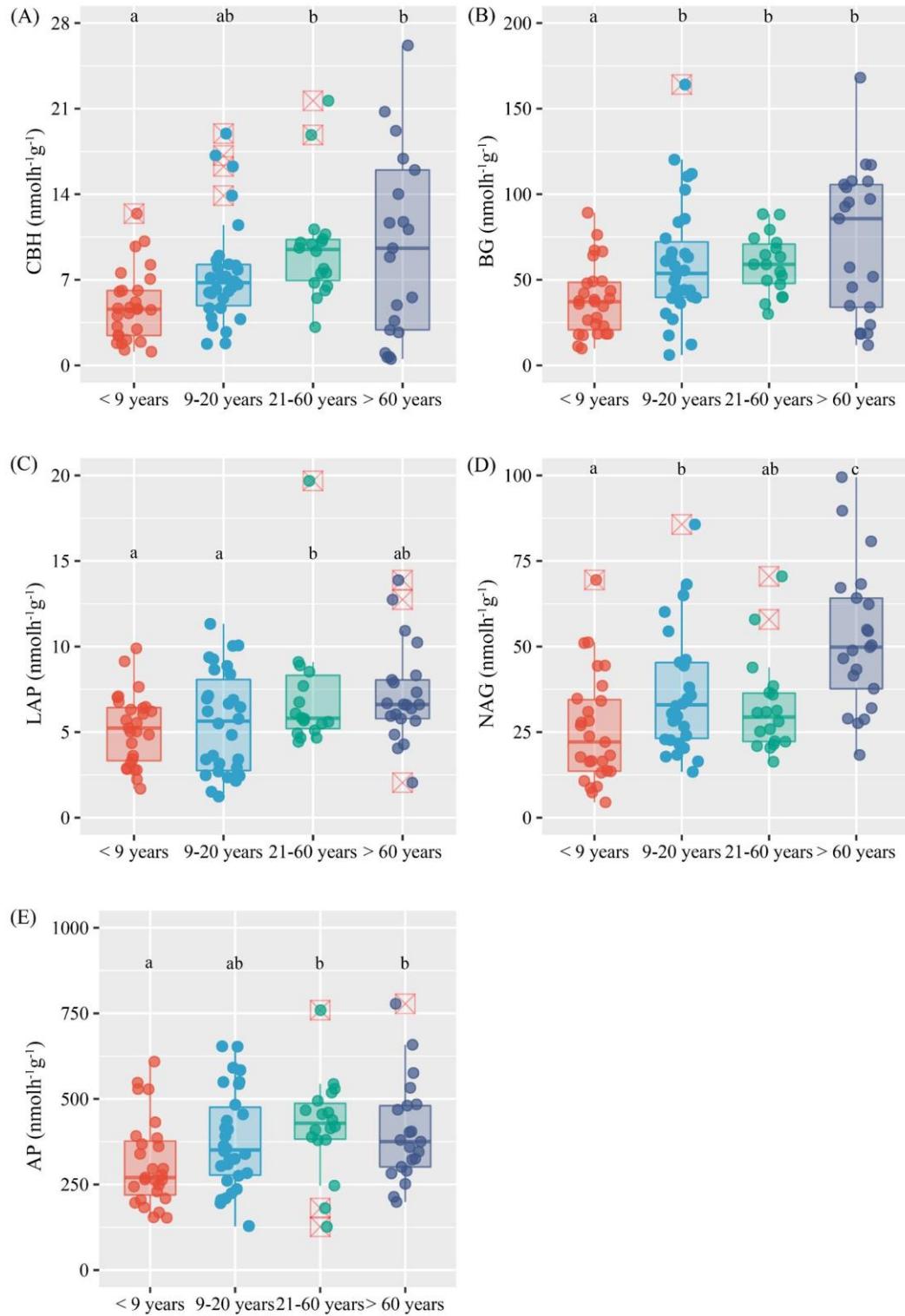


Fig. S3. Variations of soil enzymatic stoichiometry at different stand ages in *C. oleifera* plantation Note:
 (A), (CBH+BG)/(NAG+LAP); (B), (CBH+ BG)/AP; (C), (NAG + LAP)/AP. Different letters denote significant difference among stand ages ($P < 0.05$).

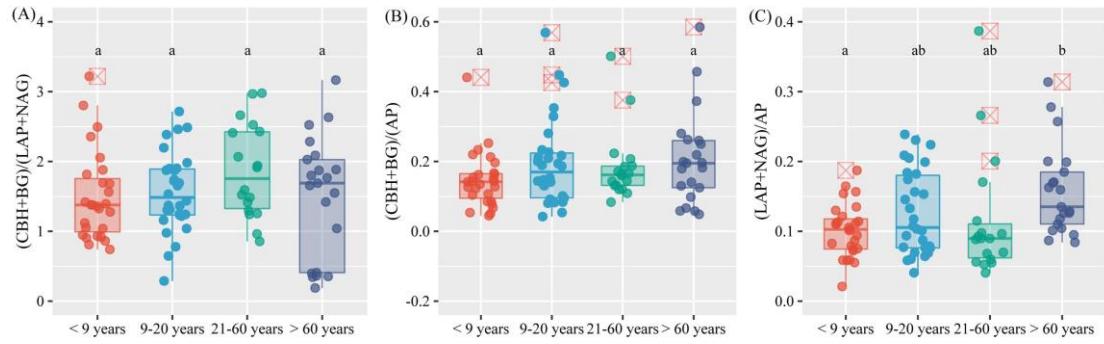


Fig. S4. Relationships between (A-C) vector A and soil organic carbon, total nitrogen or total phosphorus, and between (D-F) vector L and soil organic carbon, total nitrogen or total phosphorus. All P values of the regression analysis were < 0.05 . SOC: soil organic carbon, TN: total nitrogen, TP: total phosphorous.

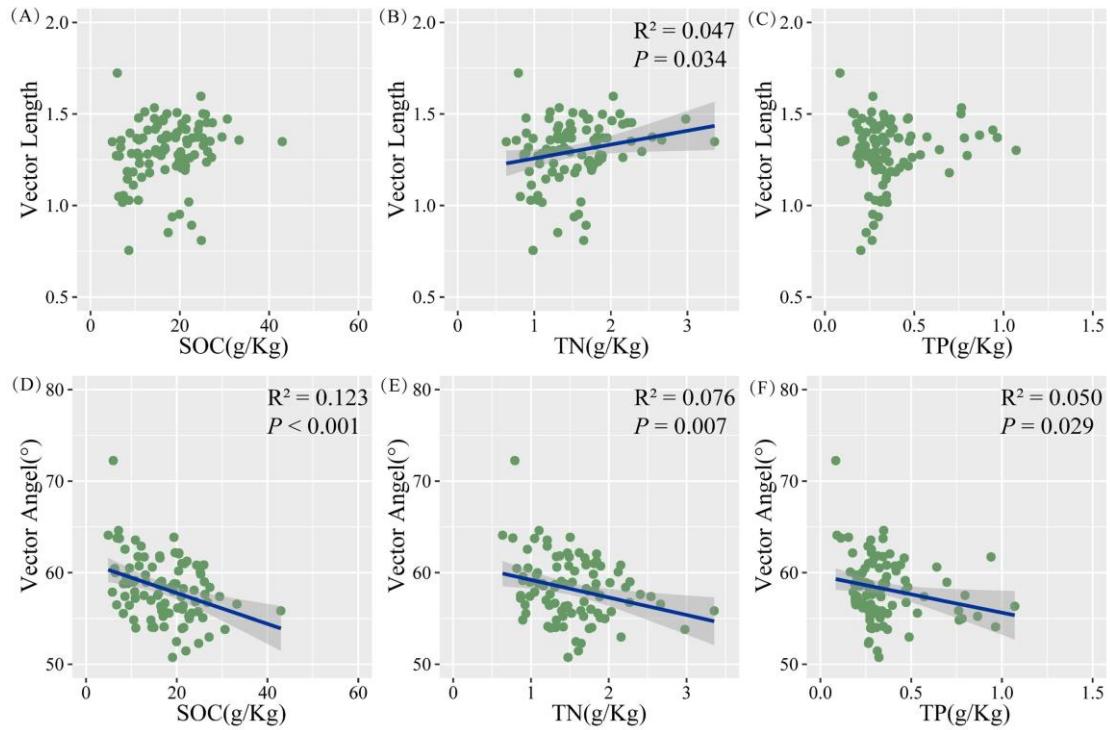


Fig. S5. Variations of soil bacterial alpha diversity at different stand ages in *C. oleifera* plantation Note:
(A), richness; (B), Shannon diversity.

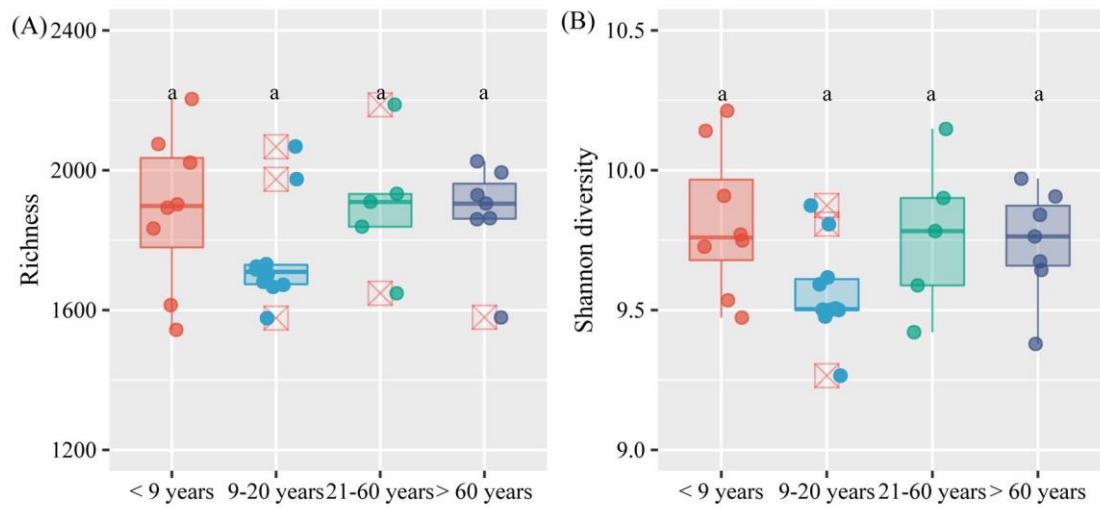


Fig. S6. Phylogenetic tree, taxonomic composition, and distribution patterns of *C. oleifera* dominant taxa.

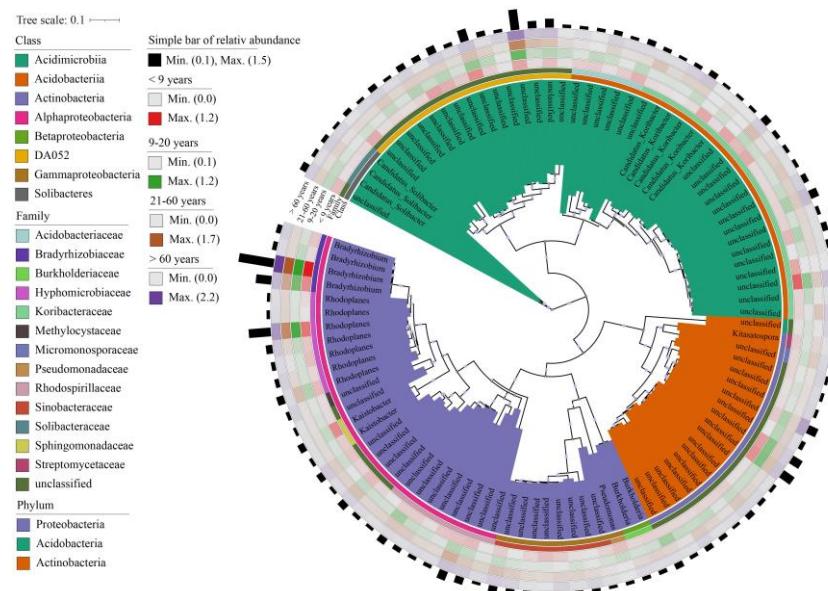


Fig. S7. Boxplot of the abundance of the 30 most abundant OTUs.

