

Supplementary

Table S1 Shifts in relative abundance of the dominated bacterial and fungal communities at phylum level at different depths in different stand ages.

Stand age	Depth	%								
		Bacteria					Fungi			
		Acidobacteria	Proteobacteria	Actinobacteria	Chloroflexi	Firmicutes	Ascomycota	Basidiomycota	Zygomycota	
3a	D1	37.72 ±1.96cd	38.80 ±1.58a	13.19 ±0.48b	3.28 ±0.45c	0.59 ±0.09a	65.94 ±0.62b	20.65 ±3.07ab	8.84 ±0.25a	
	D2	44.69 ±2.37a	32.64 ±2.59a	11.11 ±0.50a	5.32 ±0.45b	2.24 ±0.89b	70.28 ±3.44a	9.99 ±2.70b	18.21 ±0.81a	
	D3	45.20 ±2.96b	30.41 ±1.40a	11.04 ±1.05a	7.08 ±0.20c	2.26 ±0.90b	65.21 ±1.71bc	18.73 ±0.50b	14.86 ±1.19bc	
6a	D1	43.42 ±5.22bc	34.99 ±3.56ab	11.79 ±2.47b	3.41 ±0.51c	0.41 ±0.02a	69.06 ±3.94ab	20.65 ±2.23ab	13.51 ±1.68a	
	D2	49.57 ±8.16a	27.05 ±6.17ab	10.45 ±2.1a	6.32 ±0.62b	2.04 ±0.08b	75.69 ±5.03a	11.49 ±1.49b	11.91 ±0.84b	
	D3	53.23 ±6.91a	21.98 ±6.41bc	8.67 ±1.38ab	9.87 ±2.82c	2.18 ±0.76b	78.31 ±5.35a	9.77 ±1.50c	10.53 ±3.02c	
12a	D1	45.40 ±0.56abc	32.73 ±0.74b	11.30 ±0.49b	3.73 ±0.28c	0.28 ±0.02a	60.65 ±1.79bc	25.97 ±0.88a	12.85 ±0.91a	
	D2	53.65 ±5.39a	19.01 ±2.98b	10.37 ±1.84ab	10.31 ±1.40a	1.43 ±0.09b	70.33 ±3.49a	21.14 ±2.93a	5.38 ±0.22b	
	D3	54.79 ±8.74a	17.73 ±1.58d	6.10 ±1.25c	14.32 ±2.72b	2.46 ±0.46b	63.33 ±6.03bc	29.52 ±1.66a	4.55 ±0.18d	
18a	D1	51.55 ±2.23ab	23.69 ±0.63d	8.73 ±0.40c	6.11 ±0.81a	2.18 ±0.05a	59.63 ±2.62bc	25.72 ±1.66a	12.42 ±0.64a	
	D2	51.17 ±4.69a	14.91 ±1.78c	6.12 ±0.87c	8.72 ±0.36b	11.45 ±1.69a	54.51 ±8.29b	18.43 ±2.08a	24.27 ±4.09a	
	D3	56.40 ±1.21a	13.18 ±2.22d	6.34 ±1.38c	8.87 ±0.74c	8.99 ±1.53a	56.38 ±3.68c	12.70 ±3.98bc	28.08 ±2.71a	
25a	D1	53.36 ±2.52a	26.32 ±0.92cd	7.99 ±0.31c	4.34 ±0.17b	1.19 ±0.48a	51.37 ±6.45c	25.34 ±4.59a	10.31 ±3.41a	
	D2	51.12 ±7.06a	16.63 ±1.38c	6.64 ±0.69c	11.79 ±0.48a	7.41 ±0.98a	60.85 ±4.62ab	19.56 ±1.31a	15.21 ±1.16ab	
	D3	49.23 ±6.08ab	13.86 ±1.41d	8.67 ±0.25ab	16.15 ±0.73b	6.21 ±0.32a	67.38 ±4.07bc	11.95 ±2.04b	19.25 ±1.44b	
32a	D1	44.91 ±2.57abc	28.14 ±1.03c	7.99 ±0.21c	4.34 ±0.10b	1.19 ±0.06a	72.94 ±7.44ab	16.08 ±0.42b	9.06 ±0.25a	
	D2	45.73 ±2.06a	23.06 ±3.15b	7.81 ±0.42bc	15.36 ±1.26a	2.54 ±0.84b	66.60 ±5.44ab	25.48 ±7.38a	5.74 ±1.56b	
	D3	43.49 ±3.46b	19.19 ±1.48cd	7.22 ±1.17bc	22.65 ±3.83a	2.64 ±0.84b	70.48 ±4.79ab	18.11 ±4.70b	7.32 ±2.63d	
49a	D1	33.46 ±8.73d	27.88 ±2.69c	15.77 ±1.66a	3.41 ±0.08c	0.41 ±0.04a	75.57 ±2.00a	13.72 ±0.82b	8.58 ±1.15a	
	D2	44.81 ±2.26a	24.69 ±2.06b	9.36 ±0.67ab	11.79 ±0.87a	2.65 ±0.17b	70.43 ±3.22a	19.24 ±1.82a	6.27 ±1.67b	
	D3	45.30 ±2.68b	22.21 ±1.82bc	8.06 ±1.06b	15.70 ±2.11b	2.94 ±0.42b	79.37 ±5.73a	8.59 ±0.67c	11.23 ±2.84cd	

Note: Significant differences between the seven age soils were determined using one-way ANOVA at $P < 0.05$. The data are shown as the

means ±SD (n = 9). Different letters represent significant differences between the same soil depth of different stand ages at $P < 0.05$.