

# **Different responses of rhizosphere and non-rhizosphere soil microbial communities to consecutive *Piper nigrum* L. monoculture**

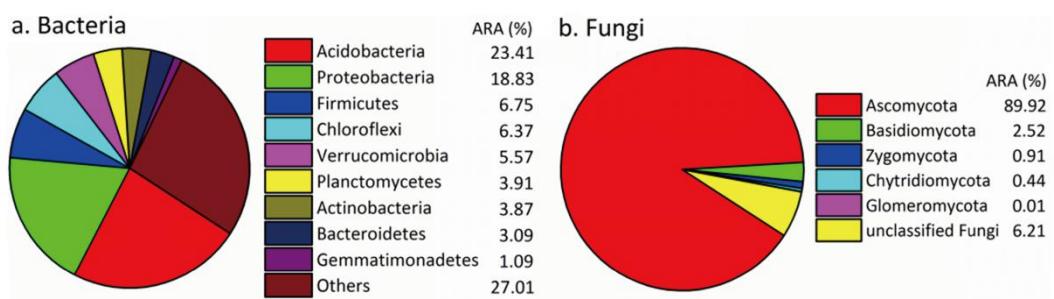
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## Supplementary Information

**Supplementary Figure S1. Overall bacterial (a) and fungal (b) communities in the 24 soil samples from four consecutive monoculture of black pepper fields.** Average relative abundances (ARAs) were calculated as the proportion of individual bacterial and fungal taxa to the total number of 16S rRNA or ITS gene sequence reads.



**Supplementary Table S1. Spearman's rank correlation ( $r$ ) between abundant taxa and soil properties.**

		OM	AN	AP	AK	pH
Rhizosphere soil	<i>Acidobacteria</i>	0.673*	-0.694*	-0.563	0.816**	0.107
	<i>Proteobacteria</i>	-0.636*	0.660*	0.622*	-0.413	0.320
	<i>Firmicutes</i>	0.417	-0.482	-0.353	0.643*	0.343
	<i>Chloroflexi</i>	-0.140	0.005	0.137	-0.047	0.390
	<i>Verrucomicrobia</i>	-0.570	0.400	0.546	-0.694*	-0.450
	<i>Actinobacteria</i>	-0.884**	0.813**	0.859**	-0.679*	-0.470
	<i>Planctomycetes</i>	-0.149	0.218	0.041	-0.531	-0.274
	<i>Bacteroidetes</i>	-0.302	0.175	0.212	-0.617*	-0.112
	<i>Gemmatimonadetes</i>	-0.618*	0.585*	0.676*	-0.187	0.016
Fungi	<i>Nitrospira</i>	0.631*	-0.643*	-0.507	0.725**	-0.077
	<i>Ascomycota</i>	-0.774**	0.675*	0.837**	-0.286	-0.489
	<i>Basidiomycota</i>	-0.098	0.317	-0.053	-0.540	-0.034
	<i>Chytridiomycota</i>	0.044	0.021	-0.005	0.342	0.310
	<i>Glomeromycota</i>	0.459	-0.448	-0.534	0.291	-0.142
Non-rhizosphere soil	<i>Zygomycota</i>	-0.132	0.140	-0.005	-0.599*	0.255
	<i>Acidobacteria</i>	-0.713**	0.743**	-0.056	-0.717*	-0.843**
	<i>Proteobacteria</i>	-0.575	0.535	-0.518	-0.485	-0.397
	<i>Firmicutes</i>	0.777*	-0.783**	-0.091	0.877**	0.890**
	<i>Chloroflexi</i>	0.494	0.086	-0.032	-0.328	0.178
	<i>Verrucomicrobia</i>	-0.472	0.374	0.302	-0.465	-0.548
	<i>Actinobacteria</i>	-0.885**	0.534	0.212	-0.408	-0.695*
	<i>Planctomycetes</i>	-0.647*	0.666*	0.237	-0.608*	-0.815**
	<i>Bacteroidetes</i>	-0.382	-0.265	0.257	0.095	-0.070
Fungi	<i>Gemmatimonadetes</i>	-0.134	0.473	0.133	-0.669*	-0.623*
	<i>Nitrospira</i>	0.246	-0.385	-0.140	0.522	0.524
	<i>Ascomycota</i>	-0.588	0.570	-0.016	-0.269	-0.597*
	<i>Basidiomycota</i>	0.463	0.231	-0.074	-0.385	-0.136
	<i>Chytridiomycota</i>	0.372	-0.227	0.122	0.068	0.460
	<i>Glomeromycota</i>	0.457	0.194	0.061	-0.279	-0.240
	<i>Zygomycota</i>	-0.137	-0.197	-0.157	-0.095	-0.060

\*P<0.05; \*\*P<0.01