

**Discovering the indigenous microbial communities associated with the natural
fermentation of sap from the cider gum *Eucalyptus gunnii***

Cristian Varela, Joanna Sundstrom, Kathleen Cuijvers, Vladimir Jiranek, Anthony Borneman

Table S1. Sequences and abundances for all bacterial OTUs.**Table S2.** Relative abundance for all bacterial phyla (%).

Phylum	Overall	Location			Niche		Date	
		Five Rivers -	Skullbone	Trawtha	Bark	Sap	Soil	Nov
		Serpentine	Plains	Makuminya				Jan
p__ <i>Acidobacteria</i>	0.087	0.032	0.091	0.111	0.116	0.066	0.094	0.079
p__ <i>Actinobacteria</i>	1.952	3.677	0.133	1.558	1.651	2.458	0.545	0.187
p__ <i>Armatimonadetes</i>	0.009	0.000	0.000	0.015	0.002	0.004	0.055	0.013
p__ <i>Bacteroidetes</i>	1.348	0.098	3.347	1.484	1.721	1.000	1.831	1.466
p__ <i>Chlamydiae</i>	0.004	0.000	0.000	0.007	0.000	0.008	0.000	0.000
p__ <i>Cyanobacteria</i>	0.066	0.000	0.000	0.111	0.000	0.090	0.148	0.056
p__ <i>Fibrobacteres</i>	0.003	0.000	0.000	0.005	0.000 ^b	0.000 ^b	0.026 ^a	0.006
p__ <i>Firmicutes</i>	3.199	2.181	1.620	4.014	1.512	3.380	7.276	3.208
p__ <i>Patescibacteria</i>	0.023	0.003	0.051	0.026	0.025 ^b	0.010 ^b	0.075 ^a	0.018
p__ <i>Planctomycetes</i>	0.017	0.019	0.000	0.020	0.024	0.003	0.059	0.025
p__ <i>Proteobacteria</i>	64.722	63.478	62.988	65.675	71.489	59.878	66.895	69.042
p__ <i>Spirochaetes</i>	0.006	0.000	0.000	0.010	0.000 ^b	0.000 ^b	0.052 ^a	0.013
p__ <i>Verrucomicrobia</i>	0.285	0.000	0.021	0.474	0.013	0.440	0.374	0.570
Unassigned	28.279	30.513	31.749	26.489	23.448	32.662	22.570	25.317
								31.104

Different letters indicate statistically significant differences ($p < 0.05$ according to Kruskal test)

Table S3. Sequences and abundances for all fungal OTUs.**Table S4.** Relative abundance for all fungal classes (%).

Class	Overall	Location			Niche		Date	
		Five Rivers - Serpentine	Skullbone	Trawtha	Bark	Sap	Soil	Nov
c__Agaricomycetes	0.201	0.608	0.004	0.058	0.103 ^b	0.288 ^a	0.100 ^b	0.062 ^b
c__Cystobasidiomycetes	0.547	1.083	0.012	0.418	0.008 ^b	1.004 ^a	0.054 ^b	1.062
c__Dothideomycetes	7.279	15.940 ^a	0.842 ^b	4.712 ^b	4.105 ^b	9.509 ^a	6.451 ^b	3.711
c__Eurotiomycetes	0.971	0.702 ^b	1.977 ^a	0.874 ^b	1.044 ^a	1.095 ^a	0.204 ^b	0.703
c__Leotiomycetes	1.459	0.663	2.697	1.553	2.262	1.236	0.140	1.832
c__Microbotryomycetes	0.295	0.688	0.060	0.166	0.026	0.253	1.266	0.418
c__Mortierellomycetes	0.685	2.441	0.005	0.026	0.093 ^a	1.219 ^a	0.000 ^b	0.032
c__Mucoromycetes	0.001	0.000 ^b	0.010 ^a	0.000 ^b	0.004	0.000	0.000	0.000
c__Orbiliomycetes	0.064	0.002	0.006	0.106	0.184	0.001	0.002	0.129
c__Pezizomycetes	0.062	0.000	0.443	0.006	0.179	0.000	0.000	0.008
c__Saccharomycetes	38.177	37.748	31.997	39.735	42.216	38.148	26.599	38.840
c__Sordariomycetes	0.508	0.428 ^a	0.158 ^b	0.622 ^a	0.124 ^b	0.808 ^a	0.273 ^b	0.771
c__Tremellomycetes	2.487	3.322	0.864	2.461	0.789 ^b	3.661 ^a	2.132 ^a	2.202
c__Wallemiomycetes	0.021	0.003	0.067	0.020	0.002 ^b	0.015 ^b	0.107 ^a	0.010
Unassigned	47.240	36.373 ^b	60.859 ^a	49.243 ^b	48.863	42.765	62.673	50.221
								44.398

Different letters indicate statistically significant differences (p < 0.05 according to Kruskal test)

Table S5. Most abundant fungal species (%).

Species	Overall
s__ <i>Kregervanrija delftensis</i>	9.308
s__ <i>Hanseniaspora valbyensis</i>	6.684
s__ <i>Lachancea quebecensis</i>	3.182
s__ <i>Citeromyces hawaiiensis</i>	2.465
s__ <i>Schwanniomyces pseudopolymorphus</i>	2.436
s__ <i>Lachancea cidri</i>	2.204
s__ <i>Torulaspora globosa</i>	2.082
s__ <i>Pichia kluyveri</i>	1.865
s__ <i>Hormonema macrosporum</i>	1.348
s__ <i>Hanseniaspora thailandica</i>	1.285
s__ <i>Candida salmanticensis</i>	1.193
s__ <i>Penidiella carpentariae</i>	0.915
s__ <i>Cladosporium delicatulum</i>	0.709
s__ <i>Bettsia alvei</i>	0.395
s__ <i>Holtermanniella takashimae</i>	0.146
s__ <i>Cercophora vinosa</i>	0.109
Unassigned	47.240
Other ¹	11.770
Ambiguous taxa	4.666

¹ identified species with abundance lower than 0.1%

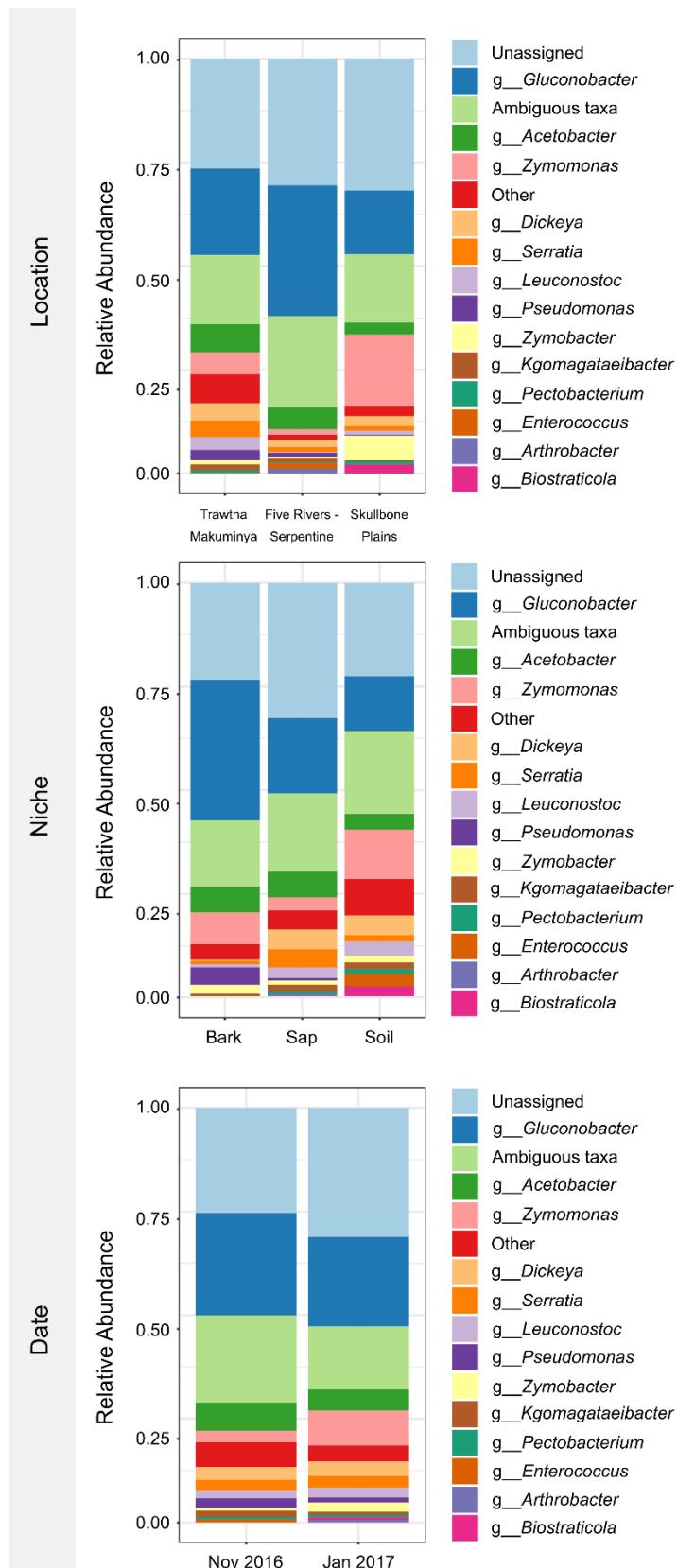


Figure S1. Relative abundance for the most abundant bacterial genera associated with *Eucalyptus gunnii* according to geographical location, ecological niche or sampling date.

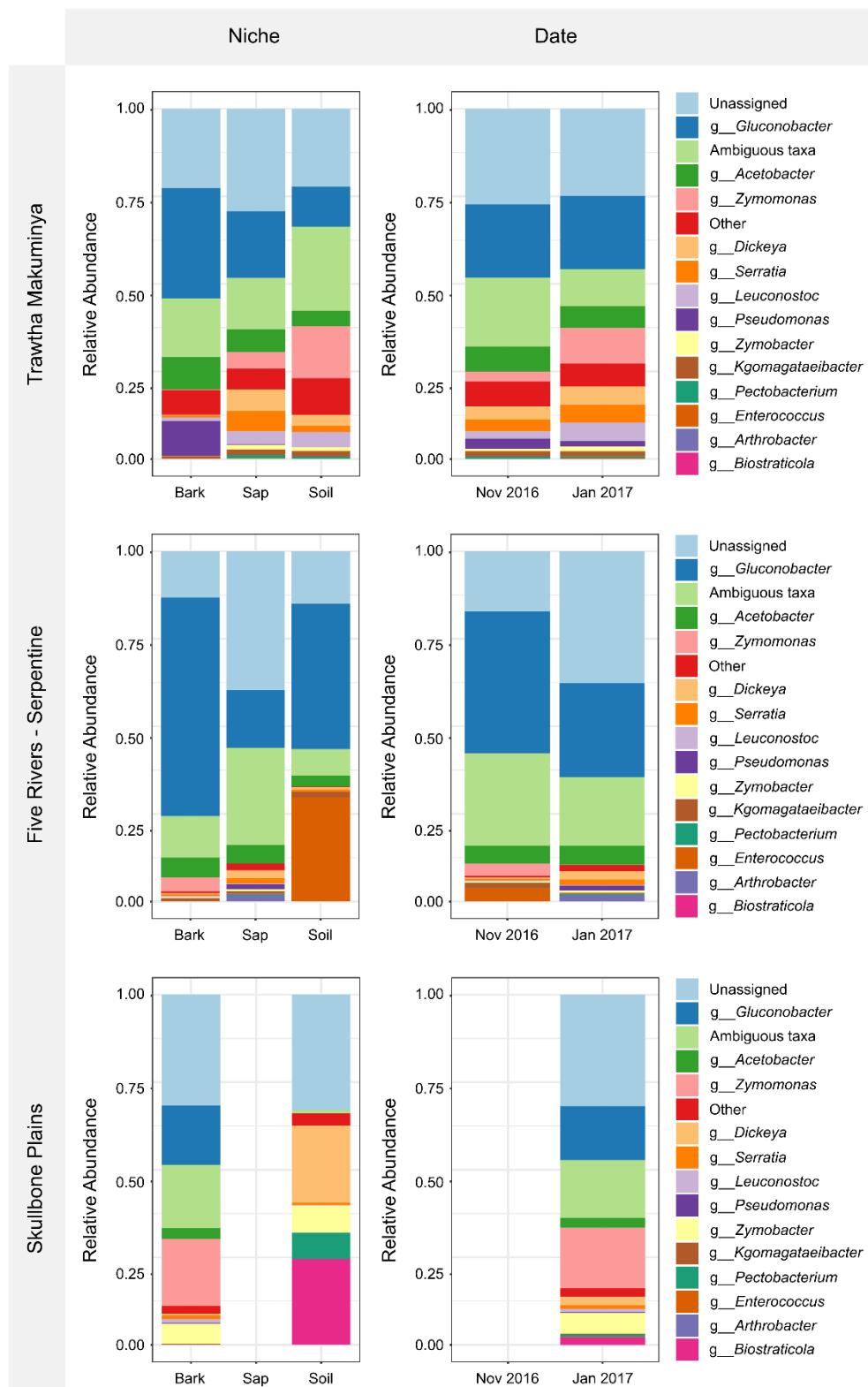


Figure S2. Relative abundance for the most abundant bacterial genera associated with *Eucalyptus gunnii* in Trawtha Makuminya, Skullbone Plains and Five Rivers - Serpentine according to ecological niche, and sampling date.

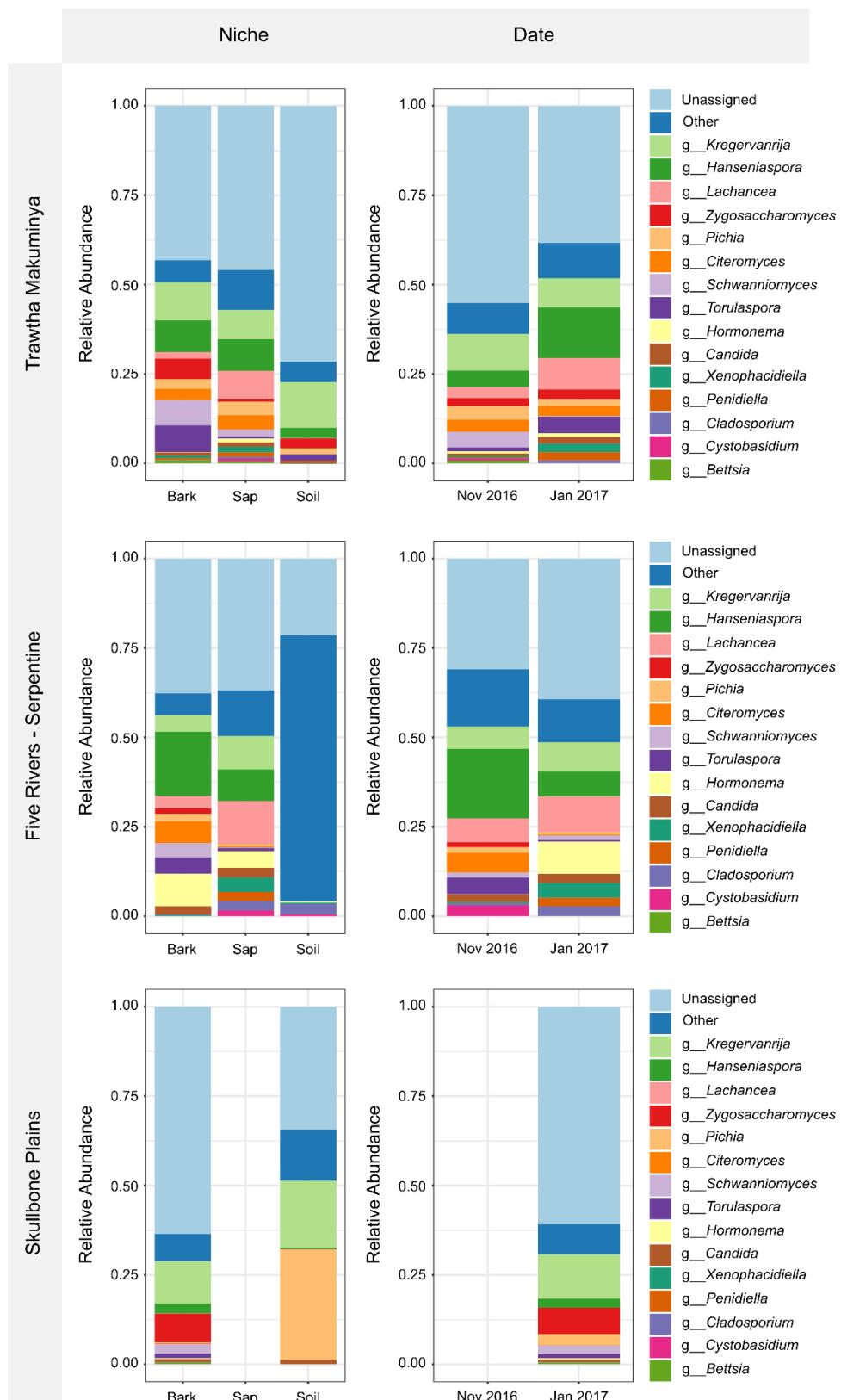


Figure S3. Relative abundance for the 15 most abundant fungal genera associated with *Eucalyptus gunnii* according to geographical location, ecological niche, and sampling date.

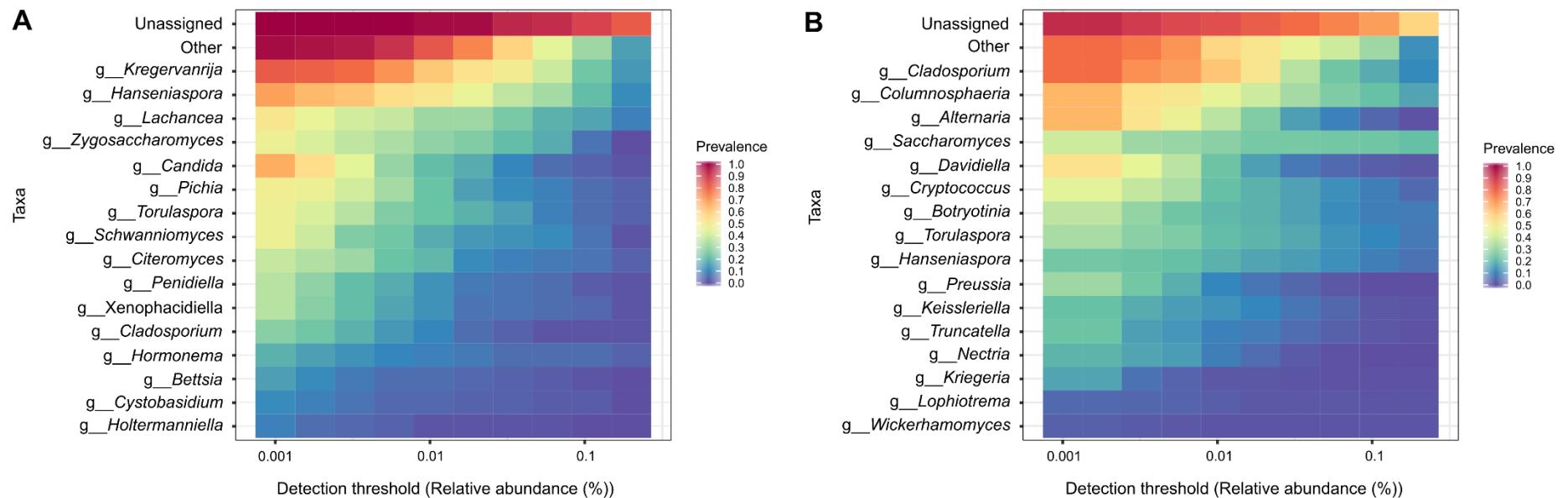


Figure S4. Most prevalent fungal genera associated with *Eucalyptus gunnii* (A) and with *Vitis vinifera* (B). *V. vinifera* data from Morrison-Whittle and Goddard^{33,34}.

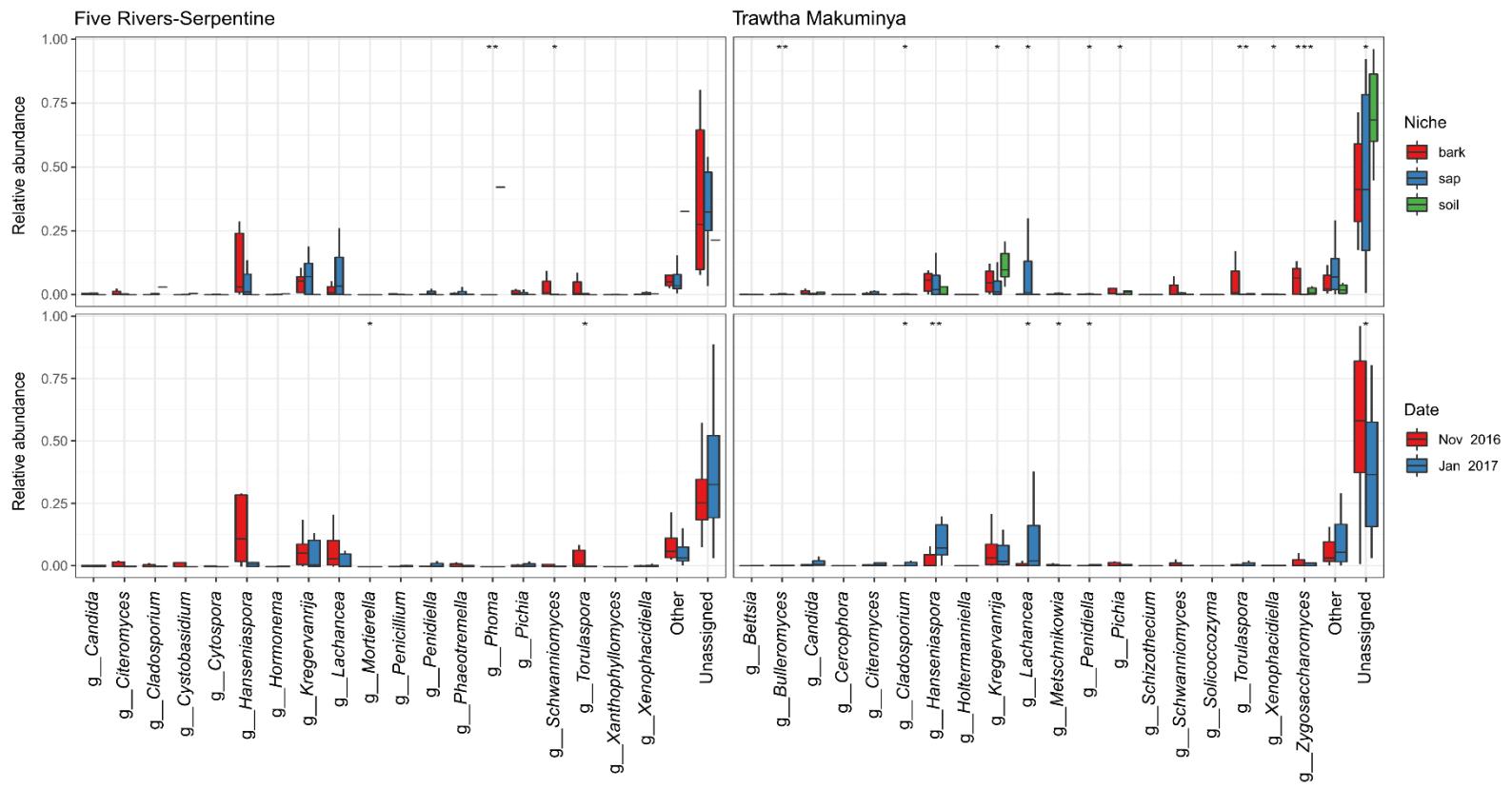


Figure S5. Boxplots showing relative abundance for the 15 most abundant fungal genera in Trawtha Makuminya and Five Rivers - Serpentine according to ecological niche and sampling date. Stars indicate statistically significative differences according to the Kruskal-Wallis test (* p < 0.05; ** p < 0.01; *** p < 0.001).