

[10.1071/CP24156](https://doi.org/10.1071/CP24156)

Crop & Pasture Science

Supplementary Material

Microbial genes highlight different trends in short term for N cycling in historical alpine pastures

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Supplementary Materials

Table S1. ANOVA and post-hoc test based on permutation test analyzing the abundance of 16S, *nifH*, *nirK*, *nosZ*, archaeal *amoA* (AOA) and eubacterial *amoA* (AOB) genes.

	<i>16S</i>			<i>nifH</i>			<i>nirK</i>				
	Df	Sum Sq	Pr(Prob)	Df	Sum Sq	Pr(Prob)	Df	Sum Sq	Pr(Prob)		
grazing	1	0.08	0.75	1	0.025	0.7451	1	0.21	0.51		
sampling time	3	0.67	0.09	3	8.33	<2e-16	***	3	4.48	0.030	*
grazing:sampling time	3	0.08	0.77	3	0.15	0.9433		3	0.15	0.99	
pasture area	2	0.80	0.10	2	0.90	0.4533		2	0.85	0.47	
grazing:pasture area	2	0.30	0.21	2	0.010	1		2	0.21	0.96	
Sampling time:pasture area	6	0.18	1.00	6	2.12	0.632		6	1.49	0.67	
Residuals	54	7.68		54	21.62			54	25.99		

	<i>nosZ</i>			<i>amoA</i> AOA			<i>amoA</i> AOB			
	Df	Sum Sq	Pr(Prob)	Df	Sum Sq	Pr(Prob)	Df	Sum Sq	Pr(Prob)	
grazing	1	5.27	0.027	*	1	0.79	0.56	1	0.93	0.58
sampling time	3	5.83	0.14		3	4.94	0.60	3	8.84	0.58
grazing:sampling time	3	0.69	1		3	3.07	0.95	3	2.83	0.73
pasture area	2	9.18	0.0088	**	2	3.21	0.74	2	9.93	0.19
grazing:pasture area	2	2.69	0.16		2	0.12	0.90	2	14.16	0.12
Sampling time:pasture area	6	5.95	0.33		6	10.69	0.93	6	7.23	0.91
Residuals	54	51.60			54	191.07		54	182.97	

<i>nosZ</i>			<i>nifH</i>			<i>nirK</i>		
Pasture area	means	groups	Sampling time	means	groups	Sampling time	means	groups
Juribello	4.72	ab	T1	4.71	a	T1	4.97	ab
Vallazza	4.08	b	T2	5.02	a	T2	5.27	ab
Venegiota	4.91	a	T3	4.15	b	T3	4.77	b
			T4	4.95	a	T4	5.41	a
Grazing	means	groups						
Yes	4.84	a						
No	4.30	b						

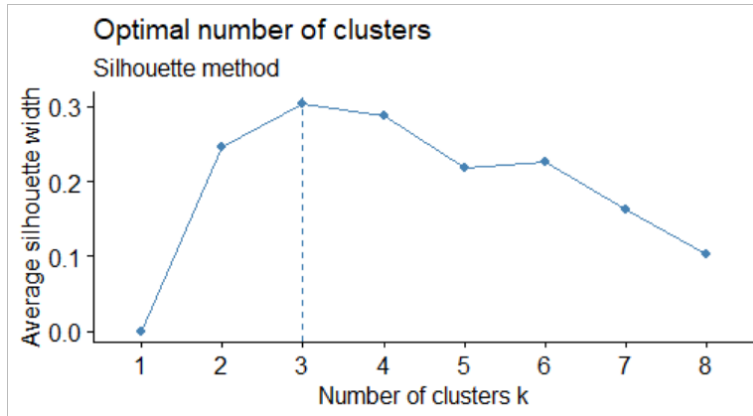


Fig. S1. Optimal number of clusters among sampling sites estimated with the Silhouette method.

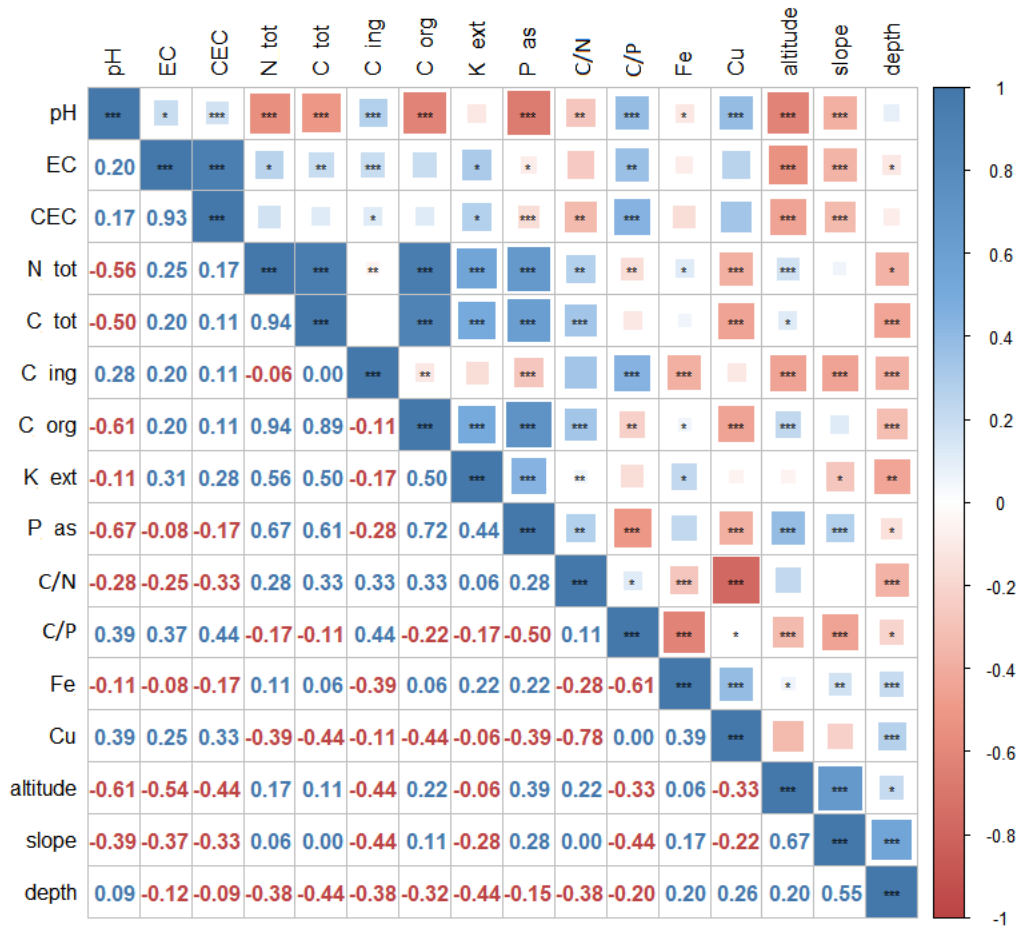


Fig. S2. Corrplot of pedological and morphological variables based on Kendall rank's coefficient coefficient (* = $P < 0.05$; ** = $P < 0.01$; *** = $P < 0.001$).