

Supplementary

Effects of amendments on soil microbial diversity, enzyme activity and nutrient accumulation after assisted phytostabilization of an extremely acidic metalliferous mine soil

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Table 1. Results of canonical correspondence analysis (CCA).

Axis	1	2
Eigenvalue	0.826	0.096
Cumulative percentage variance		
Of species data	48.2	54.7
Of species-environment correlations	84.4	95.5
species-environment correlations	0.819	0.578
Monte Carlo significance test	For all axes	
F-ratio	36.32	6.50
P-value	0.002	0.002
Intraset correlations ($100 \times r$)		
OM	-62.4	-8.1
TN	-54.9	-1.0
TP	-46.8	22.6
TK	-46.5	7.3
NH ₄ ⁺ -N	-60.1	14.6
NO ₃ ⁻ -N	-54.7	22.7
AP	-51.1	14.0
AK	-51.6	9.3



Figure S1. A photo showing the pot experiment.

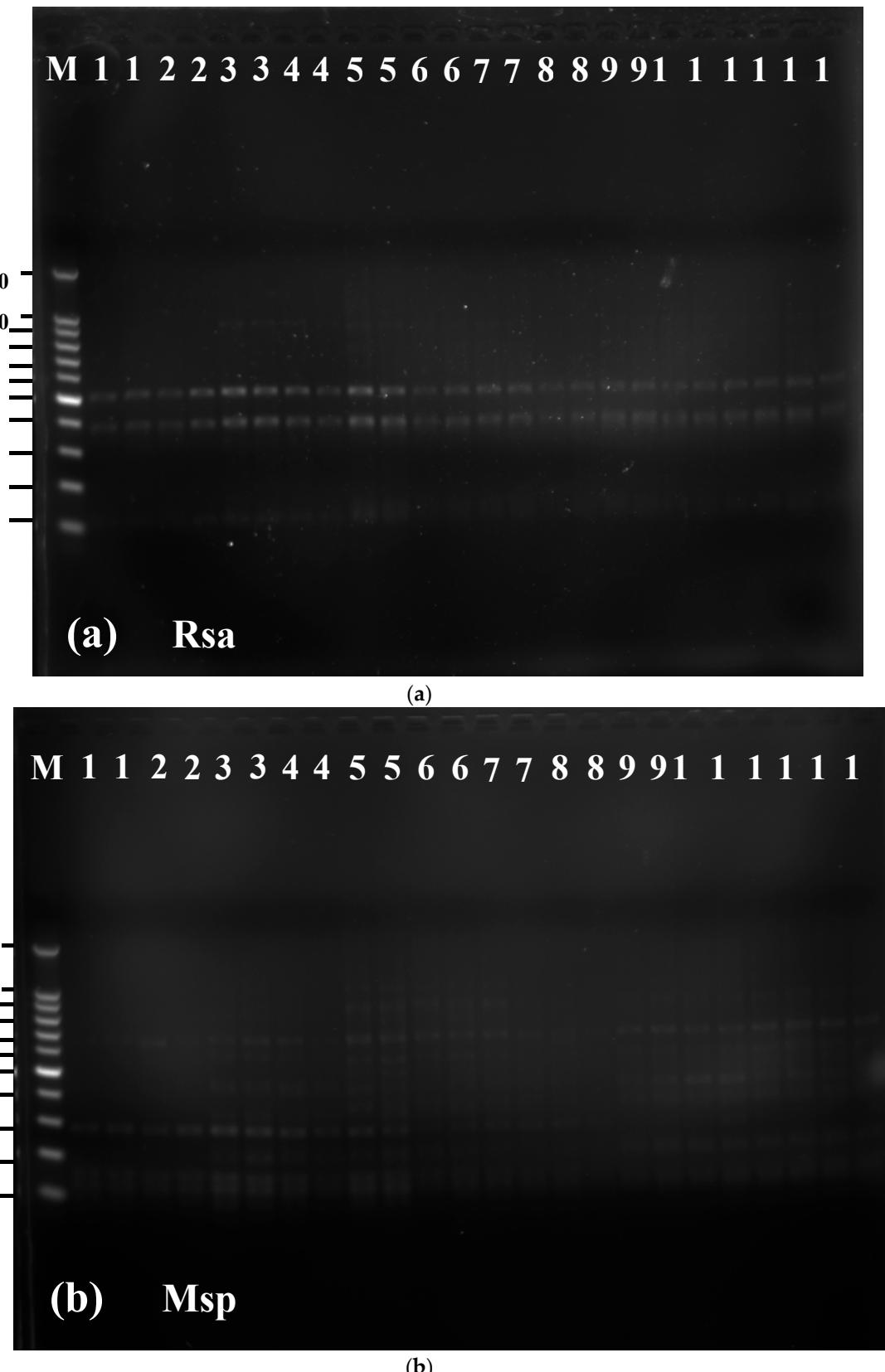


Figure S2. Bacterial community profiles based on T-RFs of the 16S rRNA gene digested with Rsa I and Msp I. Lanes 1–12 stand for 12 treatments with double lanes per treatment. Lane 1: control1, Lane 2: control 2, Lane 3: lime1, Lane 4: lime1 + NPK, Lane 5: lime1 + P, Lane 6: lime1 + Rs, Lane 7: lime1 + NPK + Rs, Lane 8: lime2, Lane 9: lime2 + NPK, Lane 10: lime2 + P, Lane 11: lime2 + Rs, Lane 12: lime2 + NPK + Rs.