

Supplementary material

Article

Microbial community investigation of wild brambles with root nodulation from a calcareous nitrogen-deficient soil

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Table S1. Amplicon Sequence Variants (ASVs) relative abundances (%) at the phylum level (no cut-off).

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Nodule	Rhizosphere	Domain	Phylum
3.56986101	63.04912244	Bacteria	Actinobacteriota
3.440624238	23.6577007	Bacteria	Bacteroidota
3.089490368	1.290372832	Bacteria	Bacillota
89.5805901	9.396095129	Bacteria	Pseudomonadota

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Table S2. Amplicon Sequence Variants (ASVs) relative abundances (%) at the genus level (cut-off 0.5%).

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Nodule	Rhizosphere	Domain	Phylum	Class	Order	Family	Genus
2.9	1.4	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Rhizobiaceae	ANPR
2.1	1.2	Bacteria	Firmicutes	Bacilli	Bacillales	Bacillaceae	<i>Bacillus</i>
0.8	0.0	Bacteria	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	<i>Brevundimonas</i>
1.4	0.8	Bacteria	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	<i>Caulobacter</i>
1.7	23.3	Bacteria	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	<i>Chitinophaga</i>
1.1	0.0	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Burkholderiaceae	<i>Cupriavidus</i>
0.7	0.0	Bacteria	Bacteroidota	Bacteroidia	Flavobacteriales	Flavobacteriaceae	<i>Flavobacterium</i>
0.0	8.8	Bacteria	Actinobacteriota	Actinobacteria	Glycomycetales	Glycomycetaceae	<i>Glycomyces</i>
0.0	0.5	Bacteria	Proteobacteria	Alphaproteobacteria	Azospirillales	Inquilinaceae	<i>Inquilinus</i>
0.0	1.3	Bacteria	Actinobacteriota	Actinobacteria	Propionibacterales	Nocardioidaceae	<i>Kribbella</i>
0.2	2.7	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Oxalobacteraceae	<i>Massilia</i>
0.0	1.7	Bacteria	Actinobacteriota	Actinobacteria	Corynebacteriales	Nocardiaceae	<i>Nocardia</i>
0.0	3.0	Bacteria	Actinobacteriota	Actinobacteria	Streptosporangiales	Streptosporangiaceae	<i>Nonomuraea</i>
0.8	0.0	Bacteria	Firmicutes	Bacilli	Paenibacillales	Paenibacillaceae	<i>Paenibacillus</i>
2.7	6.5	Bacteria	Actinobacteriota	Actinobacteria	Micrococcales	Promicromonosporaceae	<i>Promicromonospora</i>
21.3	0.0	Bacteria	Proteobacteria	Gammaproteobacteria	Pseudomonadales	Pseudomonadaceae	<i>Pseudomonas</i>
1.3	0.0	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>
55.2	0.0	Bacteria	Proteobacteria	Gammaproteobacteria	Xanthomonadales	Xanthomonadaceae	<i>Stenotrophomonas</i>
0.0	39.3	Bacteria	Actinobacteriota	Actinobacteria	Streptomycetales	Streptomycetaceae	<i>Streptomyces</i>
0.0	2.4	Bacteria	Patescibacteria	Saccharimonadia	Saccharimonadales	Saccharimonadaceae	TM7a
1.4	1.4	Bacteria	Hydrogenedentes	Hydrogenedentia	Hydrogenedentiales	Hydrogenedensaceae	unknown
1.4	0.7	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Comamonadaceae	<i>Variovorax</i>

In the Table ANPR refers to *Allorhizobium*-*Neorhizobium*-*Pararhizobium*-*Rhizobium*.

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Table S3. Biochemical traits of the Gram-positive strains with N₂-fixing ability. The green color indicates positive results, while the grey color negative ones. 5
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	N1	N2B	N6A	N6B	N6C	N7
d-xylose	grey	grey	green	grey	grey	grey
Arginine Dehydrolase 1	green	green	green	green	green	green
α-glucosidase	green	grey	grey	grey	grey	grey
Ala-Phe-Pro arylamidase	grey	grey	green	green	grey	grey
L-aspartate arylamidase	grey	grey	green	green	grey	grey
Phosphatase	grey	grey	green	grey	grey	green
Leucine arylamidase	green	green	green	green	green	green
Proline arylamidase	green	green	green	green	green	green
Pyruvate arylamidase	grey	green	green	green	green	green
Alanine arylamidase	green	green	green	green	green	green
Tyrosine arylamidase	green	green	green	green	green	green
Urease	grey	grey	grey	grey	grey	green
d-ribose	grey	grey	grey	grey	grey	green
L-Lactate alkalinization	grey	green	green	green	green	green
sucrose	green	grey	grey	grey	grey	green
d-trehalose	grey	grey	green	grey	grey	grey
Arginine Dehydrolase 2	grey	green	grey	green	grey	grey

Table S4. Biochemical traits of the Gram-negative strains with N₂-fixing ability. The green color indicates positive results, while the grey color negative ones. 7
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	N2A	N3	N5
L-pyrrolidonyl arylamidase	green	green	green
β-galattosidase	green	grey	grey
d-glucose	green	grey	green
γ-glutammil-transferasi	green	green	green
glucose fermentation	green	grey	green
β-glucosidase	green	grey	green
d-maltose	green	grey	green
d-mannitol	green	grey	green
d-mannose	green	grey	green
L-proline arylamidase	grey	green	grey
Lipase	grey	grey	green
Tyrosine arylamidase	grey	green	green
Urease	grey	green	green
Sucrose	grey	grey	green
d-Tagatose	green	green	green
d-trehalose	green	grey	green
Malonate	green	grey	green
L-Lactate alkalinization	green	grey	green
α-glucosidase	grey	green	grey
Succinate alkalinization	green	grey	green
β-n-acetyl-galattosaminidase	green	grey	green
Phosphatase	grey	green	green
Glycine arylamidase	green	grey	green
L-histidine assimilation	green	grey	green
Coumarate	grey	grey	green
O129 Resistance	green	grey	grey
Ellman	grey	green	grey

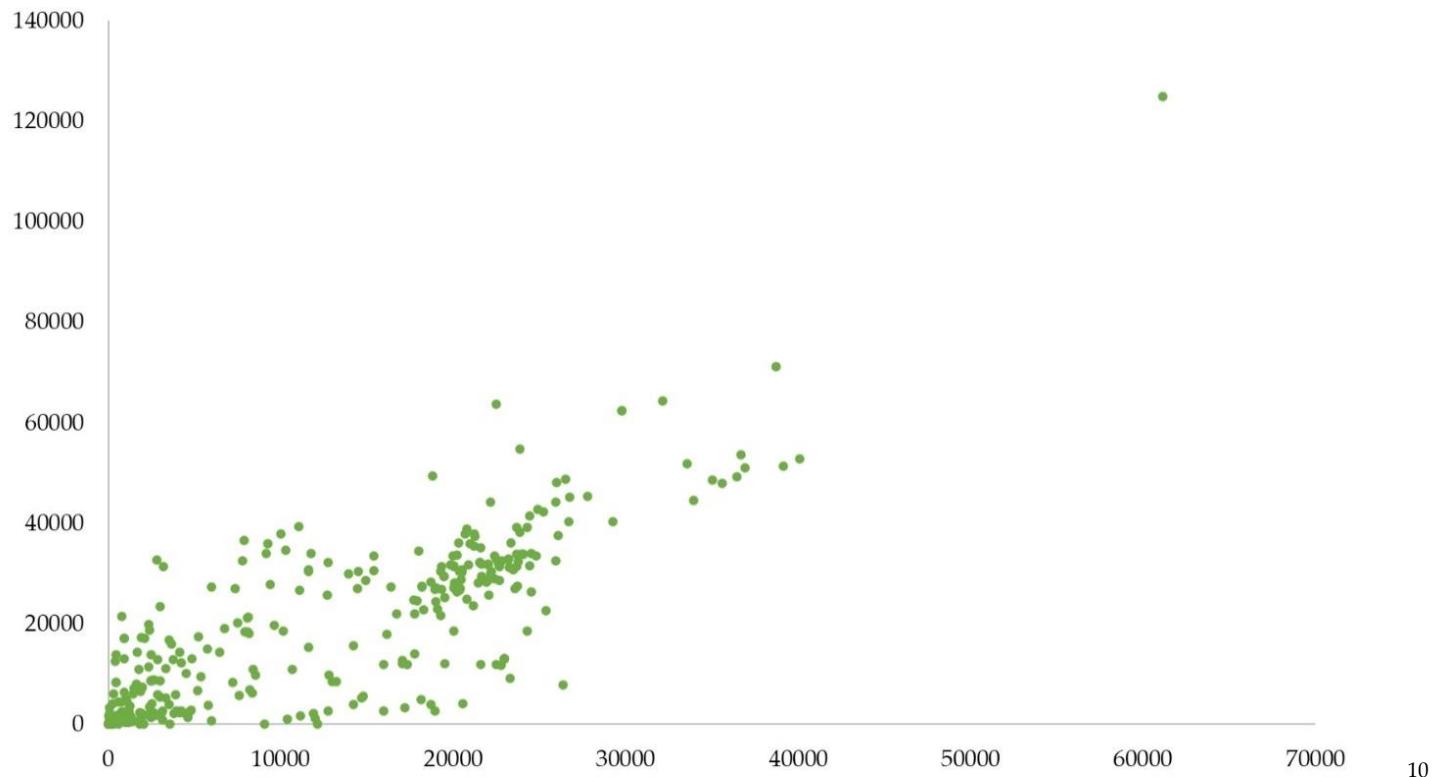


Figure S1. Correlation plot on pathways predicted by PICRUSt 2 software (nodule y axis and rhizosphere x axis). Correlation coefficient: 0.73.