

Biofertilizers: Unraveling the *in vivo* potential of PGPB traits

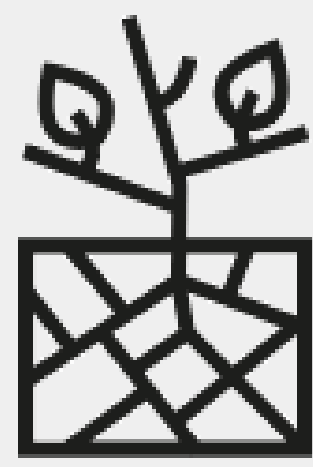
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Root-Benefit

Beneficial root-associated micro-organisms for sustainable agriculture



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Scope and Objectives

Soil degradation is a worldwide concern that results mostly from anthropogenic activities such as industry and the use of non-sustainable farming practices, like the overuse of agrochemicals. This problem severely impairs the quality and productivity of crops.

The use of formulations (**biofertilizers**) including living microorganisms, such as **phosphate-solubilizing bacteria** and **non-symbiotic nitrogen-fixing bacteria**, has attracted distinct attention due to their potential to help ameliorate soil functions while enhancing crops' growth under stress conditions.



However, **bacteria selection** is often **based exclusively on *in vitro* traits**, which coupled with a **lack of long-term quality control and cell viability assessment**, has resulted in **biofertilizer's low efficiency under field application**.

Formulations should be tailored, as far as possible, to climatic conditions, and the plant to be inoculated. **A single formulation may not be suitable for such diverse and complex ecosystems.**

This work (ongoing) aims to add value to this area of knowledge by validating, *in vivo* (greenhouse), the performance of different bacterial strains that have demonstrated the best ability to solubilize phosphorus and fix nitrogen in laboratory tests.

Methods

Soil Sampling- Mértola, Alentejo, Portugal

Four composite samples were collected from three different locations:

Alvares

Two fields: one sown with triticale and the other sown with oat.

Corte Gafo de Cima

Sheep grazing field with no specifically defined vegetation.

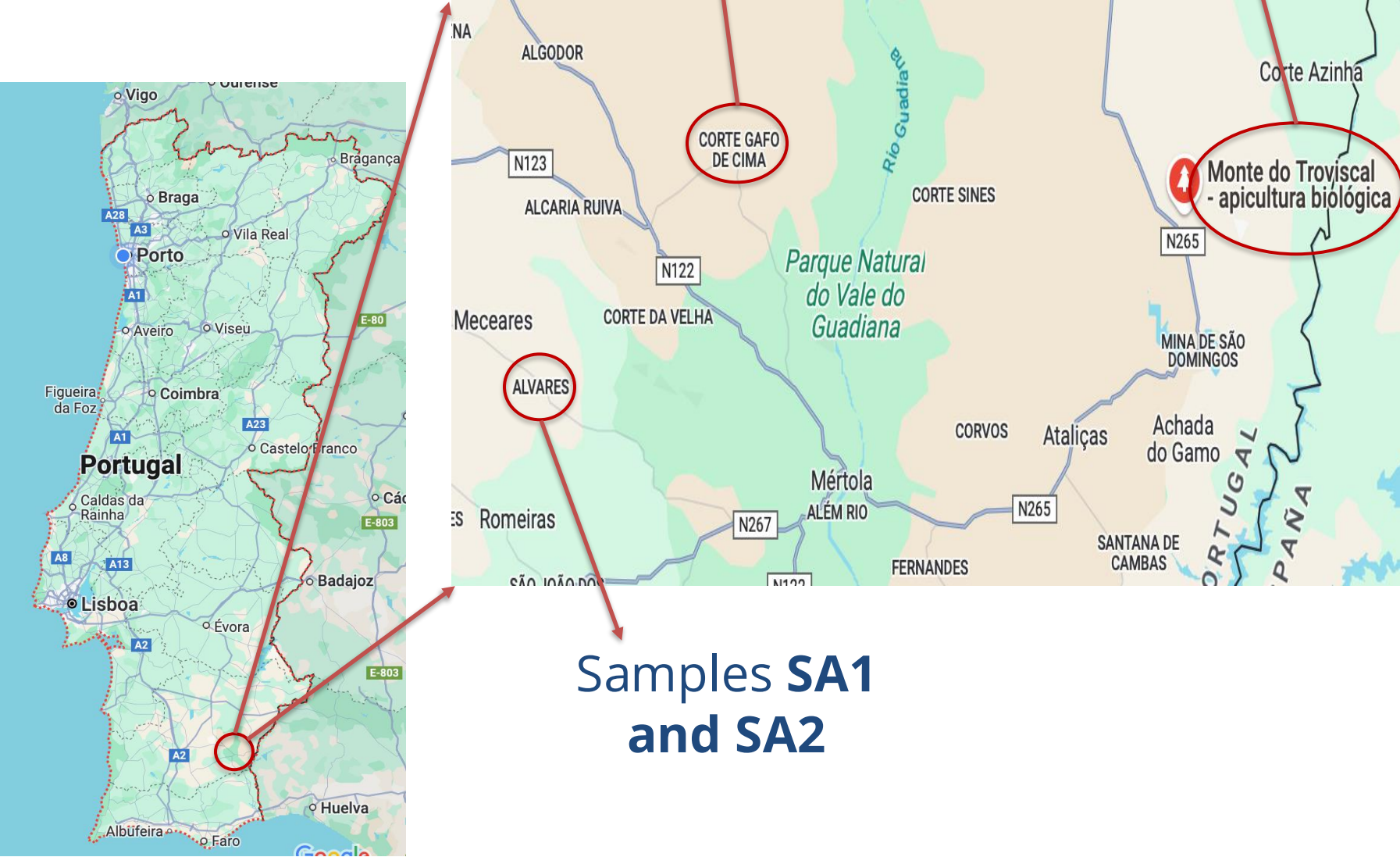
Troviscal

Field sown with oat and lupin in the soil stand-by season.



Sample SCGC

Sample ST

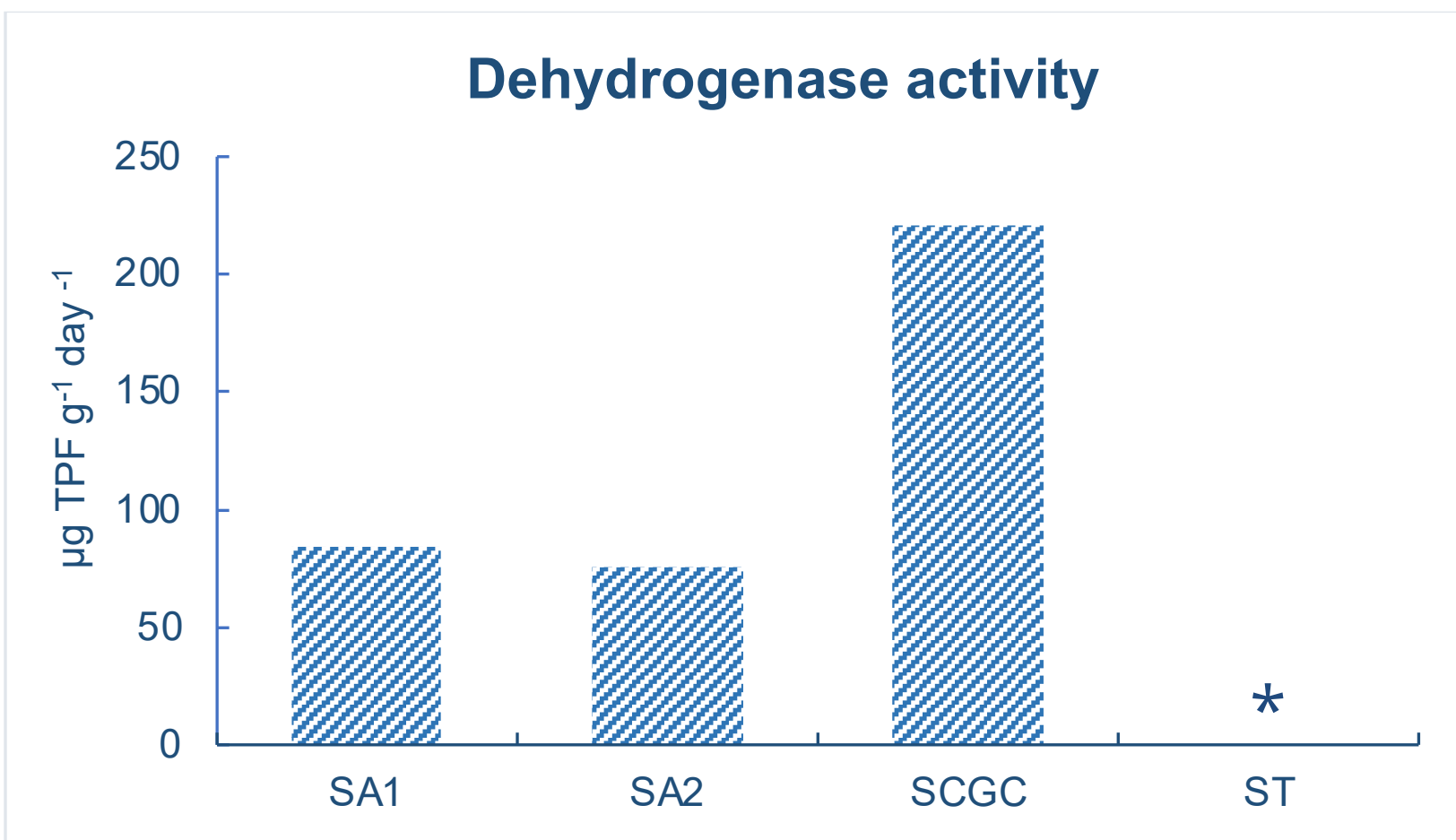


After soil physicochemical, biochemical characterization and microbial functional diversity assessment, **54 representative bacterial strains** were isolated and **P solubilization and N₂ fixation abilities** were verified.

Results

Soil Biochemical Characterization

Soil dehydrogenase enzymatic activity enables to infer about the viable microbial community present in the soil.

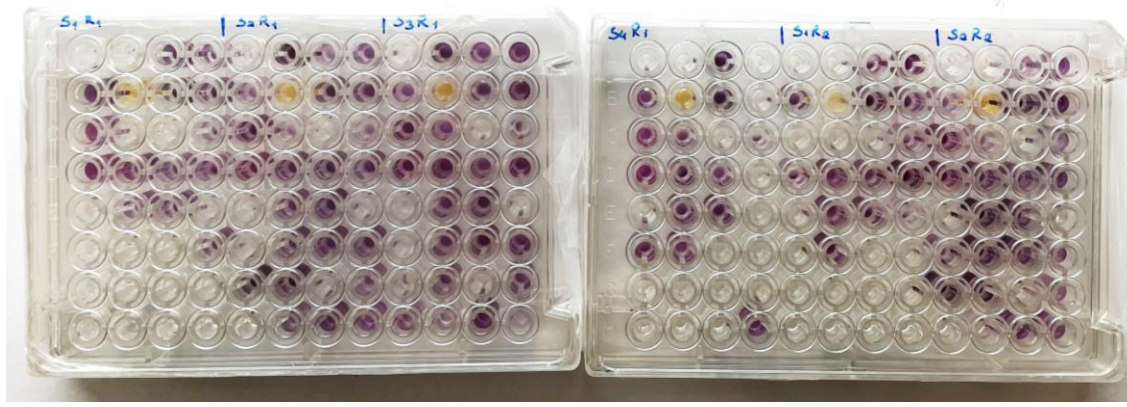


*below the limit of quantification

Soil Microbial Functional Diversity

Assessment of the physiological profile of the microbial community through the ability to oxidize different C substrates in a 96-well commercial plate.

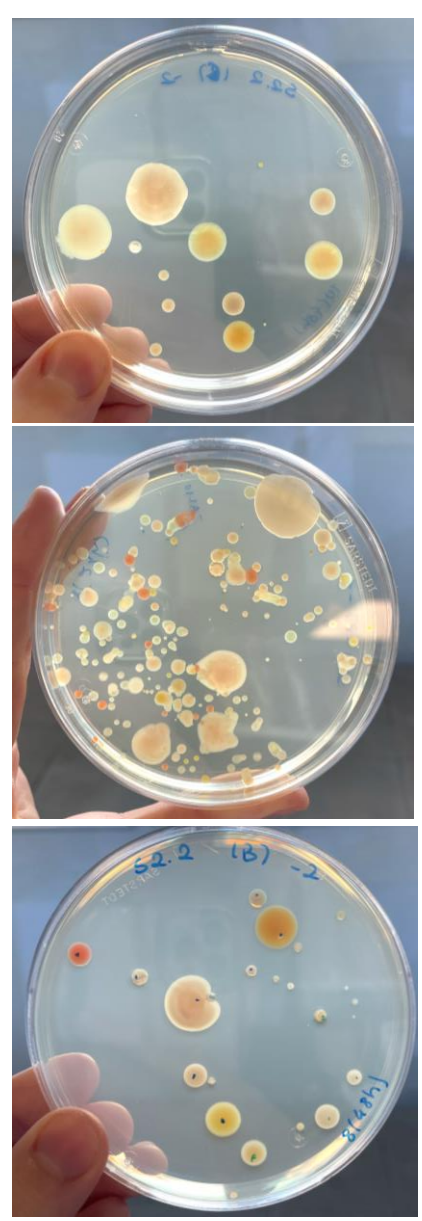
Biolog®
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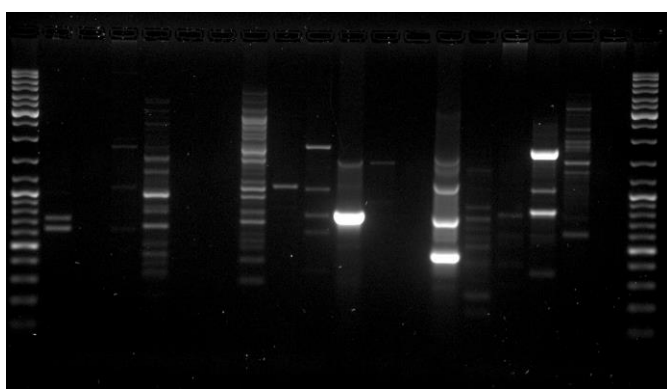
	Richness / NUS	Shannon's Diversity Index (H')	Shannon's Evenness Index (E)
SA1	16.17 ± 2.47	2.74 ± 0.17	0.99 ± 0.01
SA2	24.17 ± 0.76	3.13 ± 0.01	0.98 ± 0.01
SCGC	22.5 ± 1.0	3.06 ± 0.04	0.98 ± 0.01
ST	BLQ	BLQ	BLQ

BLQ: below the limit of quantification

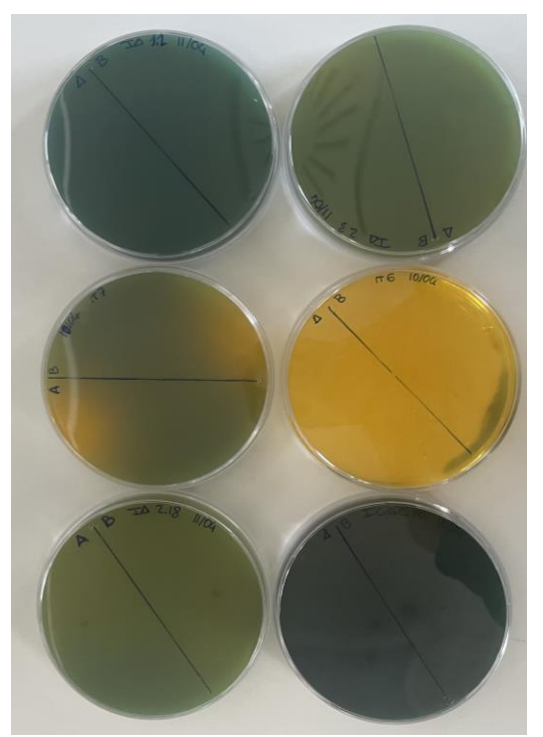
Bacteria isolation, and PGP traits assessment



54 representative strains (BOX-PCR fingerprinting analysis through BioNumerics software)



BioNumerics



P solubilization test

≈ 30% of the isolates solubilized Ca₃(PO₄)₂ (with higher and lower solubilization indexes)

N₂ fixation test

≈ 54.5% of the isolates exhibited nitrogen fixation (qualitative analysis)

Ongoing Work

- ✓ Several PGPB traits are being tested: **osmotic stress resistance, siderophores, lytic enzymes, and EPS production.**
- ✓ Selection of top ten best-performing bacterial strains.
- ✓ **PGPB testing in a pot experiment** using soil from Mértola's region and two different crops (wheat and oat).
- ✓ Plants will be analyzed for biomass, elongation, and nutritional composition.
- ✓ Rhizospheric soil samples will be collected at sowing, after bacterial inoculation, and at the end of the experiment for microbial community analysis and physicochemical characterization.

Acknowledgements

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