Recovery of microbiological status with organic amendments on soils affected by mining activity in a decadal temporal scale.

PhD. Raúl ortega

Department of Agronomy & CIAIMBITAL, <u>rortega@ual.es</u>. University of Almeria, E-04120, Almería, Spain.





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Centro de Investigación en Agrosistemas Intensivos Mediterraneos a Biotecnología Agroalimentaria (CIAMBITAL) Universidad de Almería (UAL)



ANÁLISIS DE PROPIEDADES MICROBIOLÓGICAS (METAGENÓMICA)



ANÁLISIS DE PROPIEDADES FÍSICAS Y QUÍMICAS



MICROSCOPÍA ÓPTICA Y ELECTRÓNICA







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MIEMBROS DEL LABORATORIO

RNM-934: Agronomía y Medio Ambiente (AGROMA)

Investigadores

Raúl Ortega Pérez (Responsable). e-mail: rortega@ual.es

Isabel Miralles Mellado. e-mail: imiralles@ual.es

Manuel Esteban Lucas Borja. e-mail: manuelesteban.lucas@uclm.es

Técnicos

Natalia Rodríguez Fernández. e-mail: nrodfer@ual.es

Rocio Soria Martínez. e-mail: rocio.soria@ual.es

Víctor Manuel Gallegos Cedillo e-mail: vgc253@ual.es

Paloma Gómez Belmonte e-mail: biopgb@ual.es

Colaboradores

Miguel Ángel Domene Ruiz. e-mail: madomene@fundacioncajamar.com





Almería, SE Spain

Opencast mining for aggregate extraction



Semiarid climate with extreme weather conditions



high solar radiation



loss plant cover



Completely degraded soil with high erosion rates

Limitations on physical, chemical and biological properties

Luna et al., 2016



Source: Isabel Miralles



Introduction Almería, SE Spain

Opencast mining for aggregate extraction





Semiarid climate with extreme weather conditions



high solar radiation



Loss of productivity and soil fertility

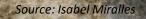
(decreasing organic matter, N, structure, etc)

Loss of soil quality

Loss of ecosystem functionality

DESERTIFICATION!!

2016



Why use organic amendments?

Organic amendments

Restoration

Accelerates recovery

Improve soil properties

Soil microorganims could be excellent indicators to evaluate quality recover in restoring soils

Soil bacteria communities

Contribute to plant establishment

(Zink and Allen, 1998)

Improve biogeochemicals cycles (N, C, P)

(Bastida et al., 2015)



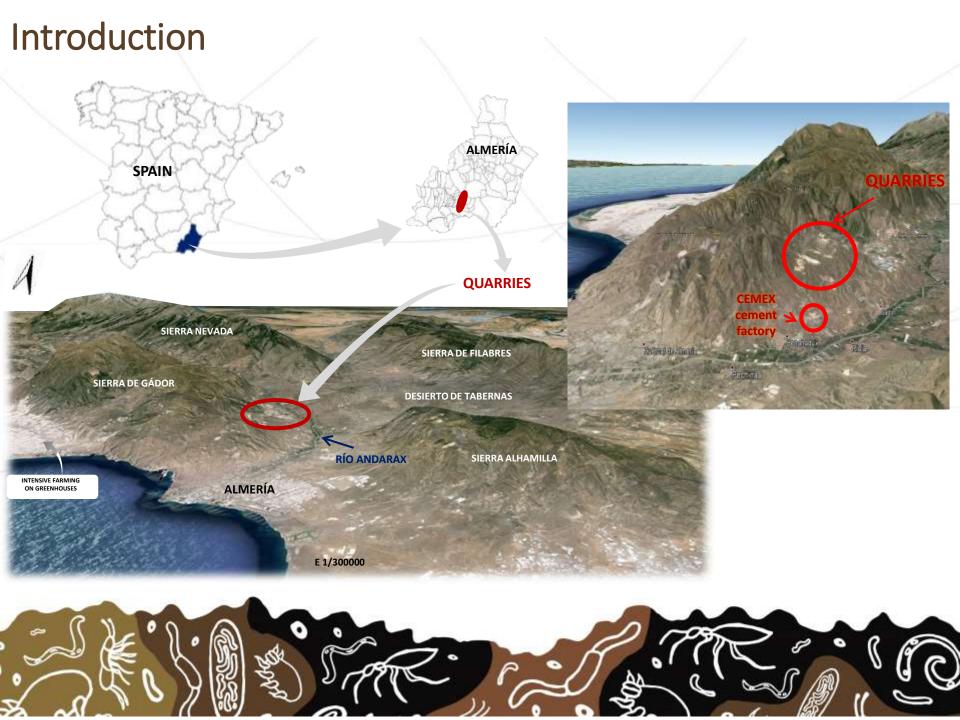
Improve soil structural formation

(Bender et al., 2016)

Organic matter transformation

(Zhao et al., 2019)





Experimental plots 2008



Experimental plots 2018



Long term study

Experimental plots 2008



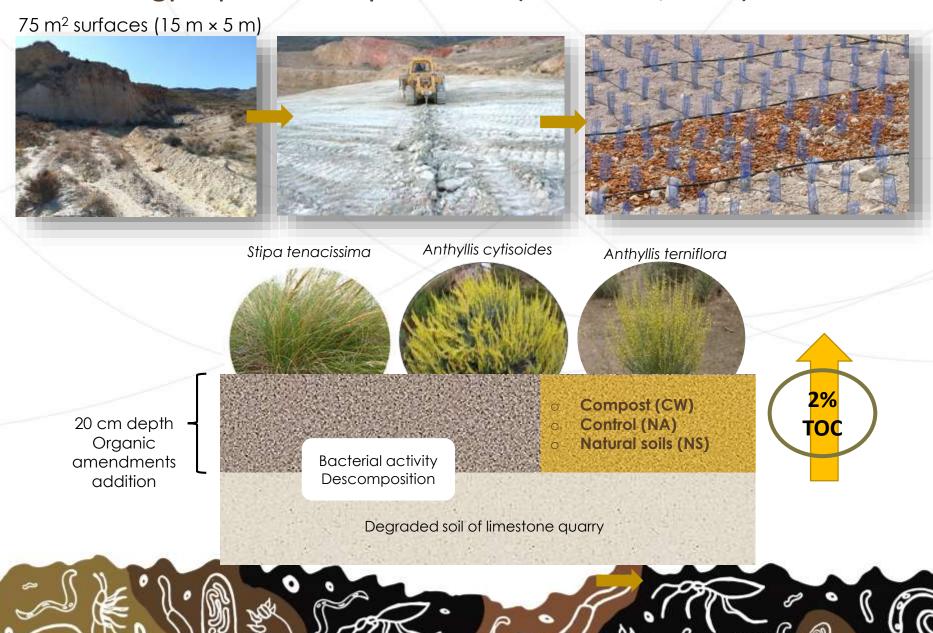
Experimental plots 2018

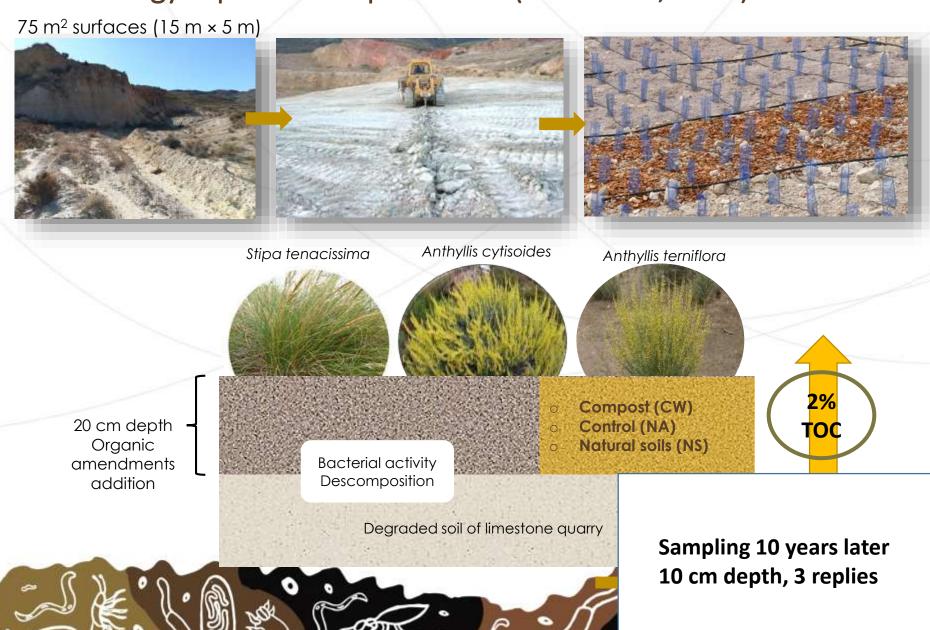


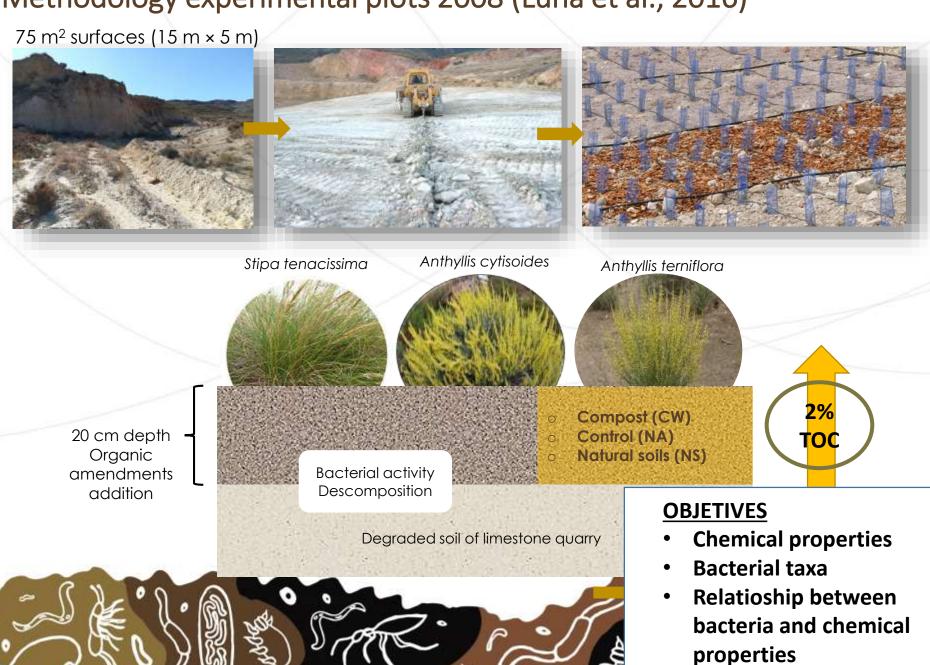


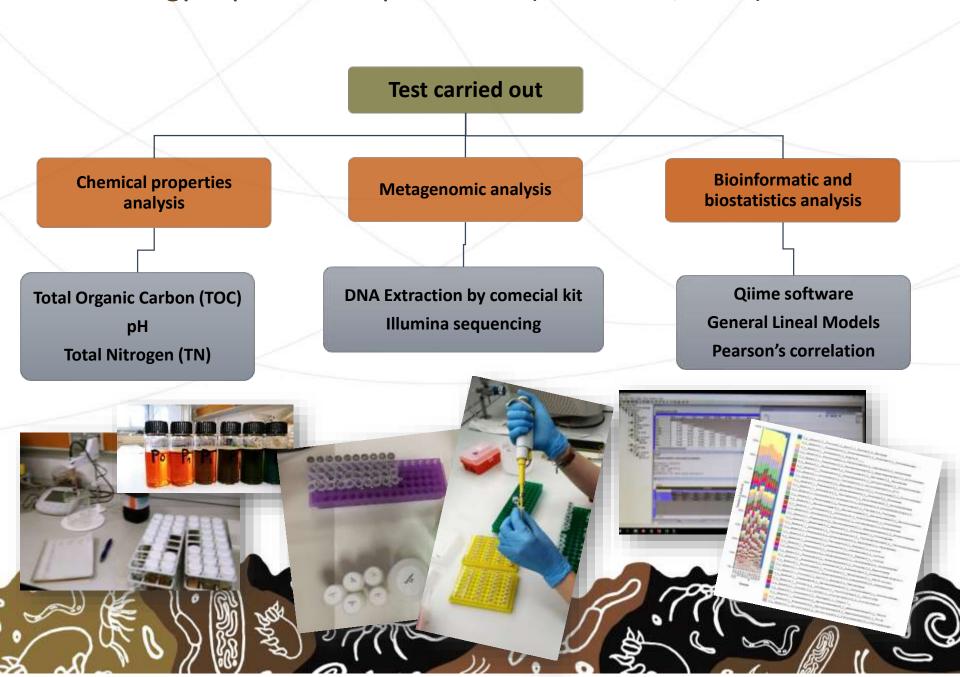
The <u>objective</u> of this work was to study the bacterial communities at genera taxonomic level to determine if application of organic amendment (compost from urban waste) approached in the long-term to reference state (natural soils) after its addition in restored soils of a limestone quarry in a semi-arid climate.





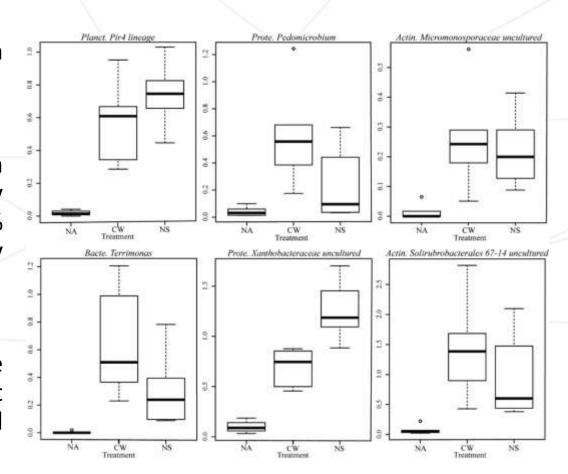






Results

- 162 soil bacteria taxa with abundance upper 0.1%.
- GLMs showed compost from urban waste (CW) significantly influenced (p<0.05) the 59% of bacterial taxa and 14% by natural soils (NS).
- Boxplots revealed some bacteria taxa most abundant in both compost (CW) and natural soils (NS).





Results

Chemical soil properties

	рН	тос	TN
Compost (CW)	8.0 ± 0.1 a	3.4 ± 1.5 a	0.5 ± 0.2 a
Reference Soil (NS)	8.1 ± 0.1 a	3.1 ± 0.5 a	0.3 ± 0.1 c
No Amendment (NA)	8.7 ± 0.1 b	0.2 ± 0.0 b	0.02 ± 0.01 b

Values represent average \pm standard deviation (p<0.05). Letters show statistical differences between treatments.



Results

Pearson's correlation analysis of soil bacteria taxa that were common between CW-treated soils and natural soils and chemical soil properties showed significant (p < 0.05) and positive correlations with TOC and TN and significant negative correlation with soil pH.

	pН	TOC	TN
Actin. Blastococcus	-0,336	0,305	0,416
Actin. Geodermatophilus	-0,498	0,561	0,612
Actin. Micromonosporaceae uncultured	-0,629	0,784	0,717
Actin. Solirubrobacterales 67-14 uncultured	-0,651	0,680	0,747
Bacte. Chryseolinea	-0,414	0,452	0,115
Bacte. Terrimonas	-0,731	0,725	0,770
Planc. Pir4 lineag	-0,761	0,852	0,736
Planc. Singulisphaera	-0,552	0,551	0,351
Prote. Pedomicrobium	-0,615	0,802	0,805
Prote. Xanthobacteraceae uncultured	-0,636	0,709	0,378



Discussion

- Compost treatment (CW) promoted in the long-term soil restoration, showing TOC and TN values similar to natural soils (NS).
- Organic amendments have contributed in different studies to increase and improve the microbial proliferation of the soil (Bastida et al., 2008; Tejada et al., 2006).
- Type of organic matter and plant residues influence the composition of soil microbial communities (Kramer and Gleixner, 2008).
- The soil treated with compost was the most influential in the bacterial taxa filling us to believe that this organic amendment favoured the bacterial proliferation due to the improvement of the chemical properties of the soil which in turn favoured increased plant growth by Luna et al. (2016) five years after restoration.
- Interestingly, these bacterial taxa were present almost exclusively in soils amended with compost and natural soils as showed boxplots.
- May have been the organic amendment of compost was the one that came closest to the reference soils, confirming again that it is possibly due to the vegetal stabilization that favours rhizospheric niches and also to the improvement of the chemical properties of the soil (Bastida et al., 2016).



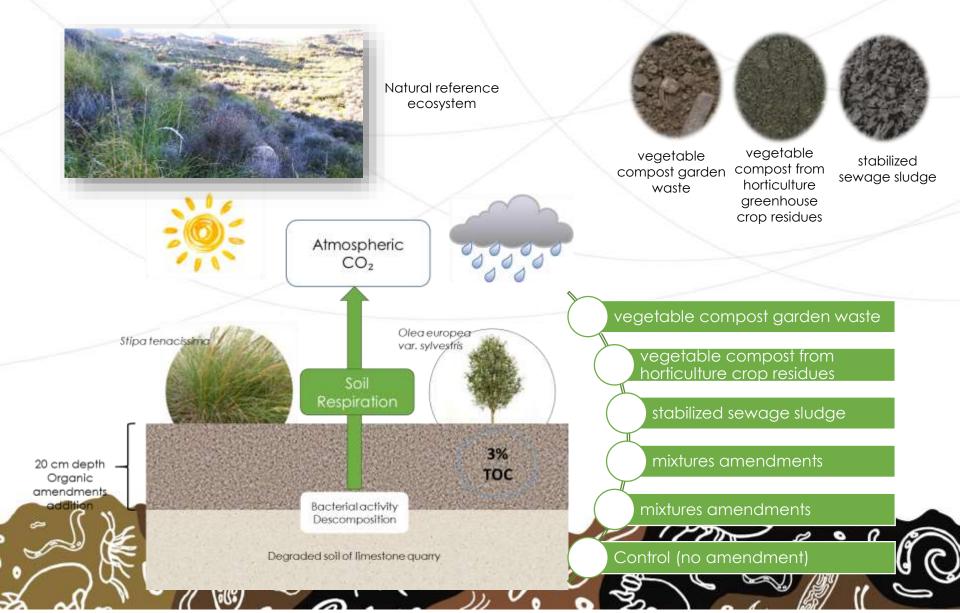
Discussion

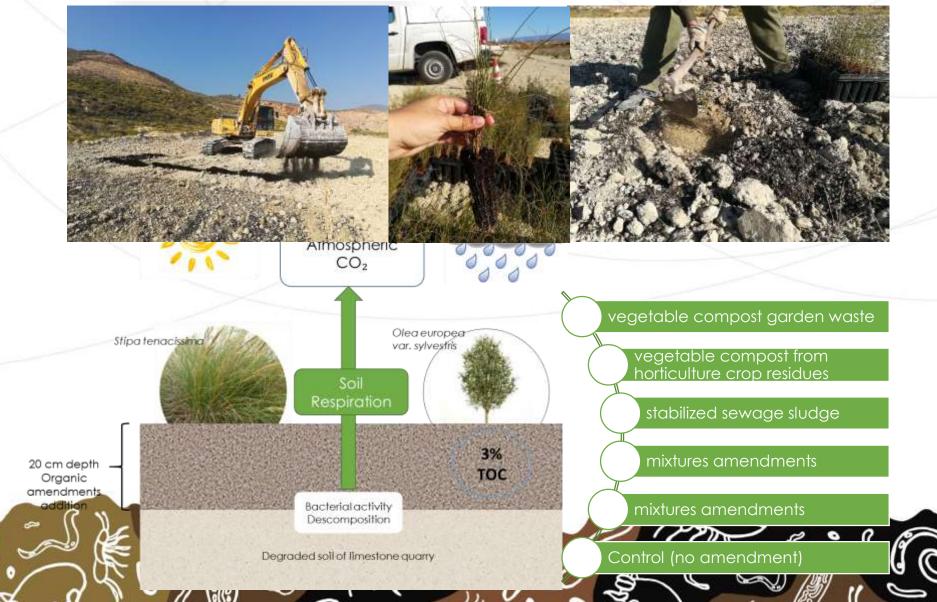
- Several common bacterial taxa among CW-treated and NS soils also showed significant positive correlations with soil TOC and TN (Table 1). Besides, bacterial taxa as Pedomicrobium was observed in soils rich in organic matter (Lima et al., 2015) or as Terrimonas that was found in developed soils near our study area (Sánchez-Marañón et al., 2017), corroborating that compost-treated soils improved their quality in ten years.
- Similarly, these bacterial taxa also strong negative correlations with soil pH (Table 1). Lauber et al. (2009) established that soil pH plays a crucial role on soil microbial composition which drive changes in soil bacterial taxa.

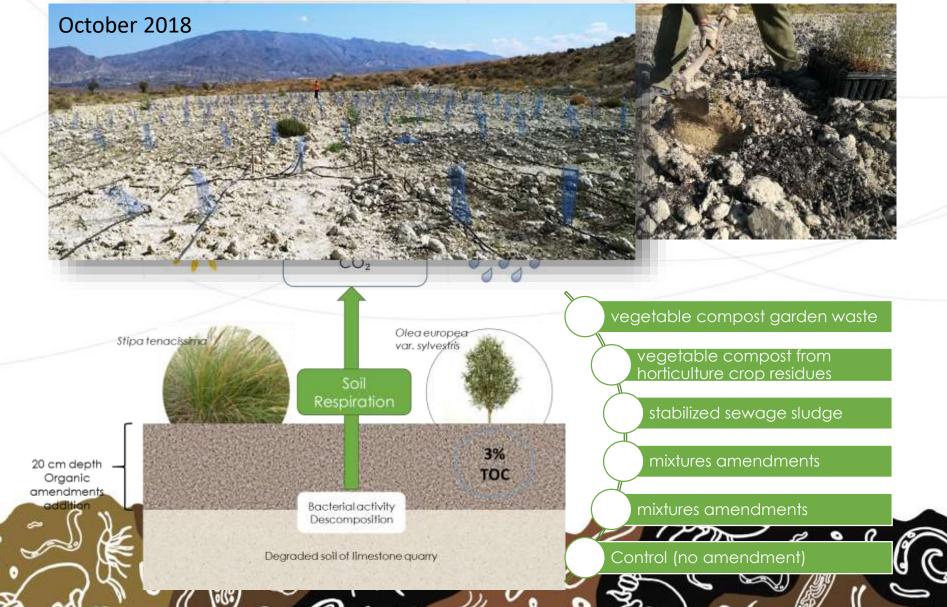
Conclusions

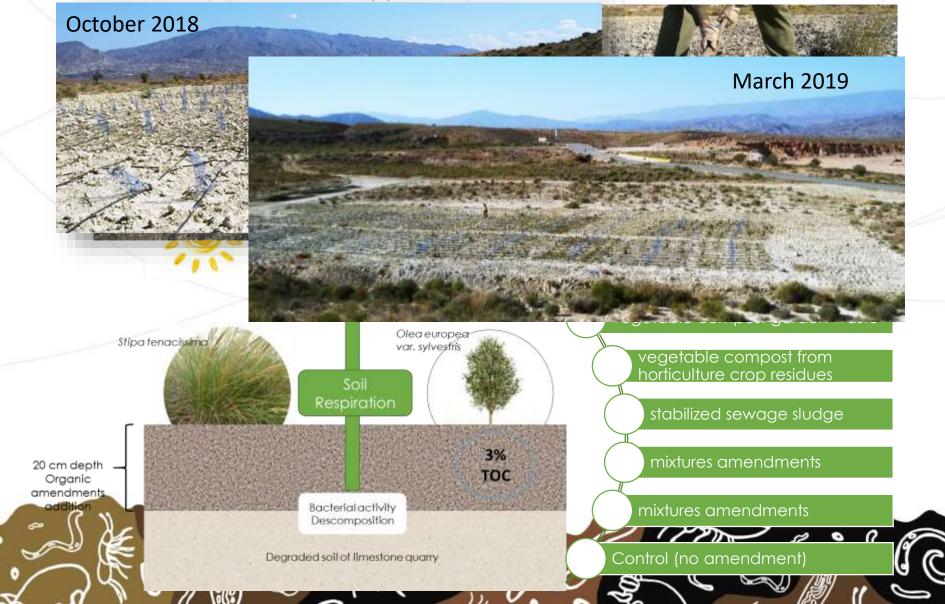
These results suggest that long-term restored soils with compost treatment have established microbial communities similar to surrounding reference natural soils. These results show that compost management of urban waste is a suitable method to recover microbial communities of degraded soils in a temporal scale.

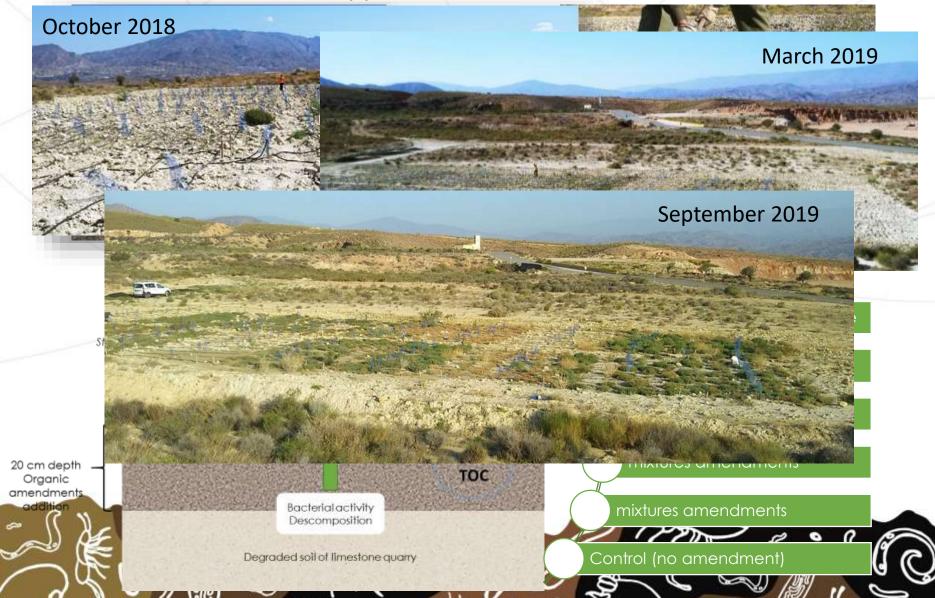


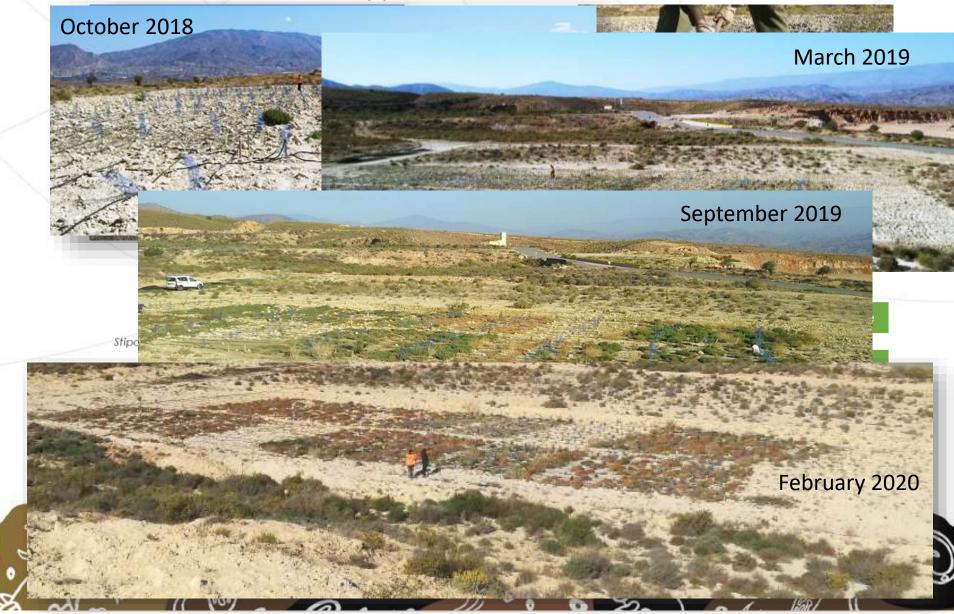


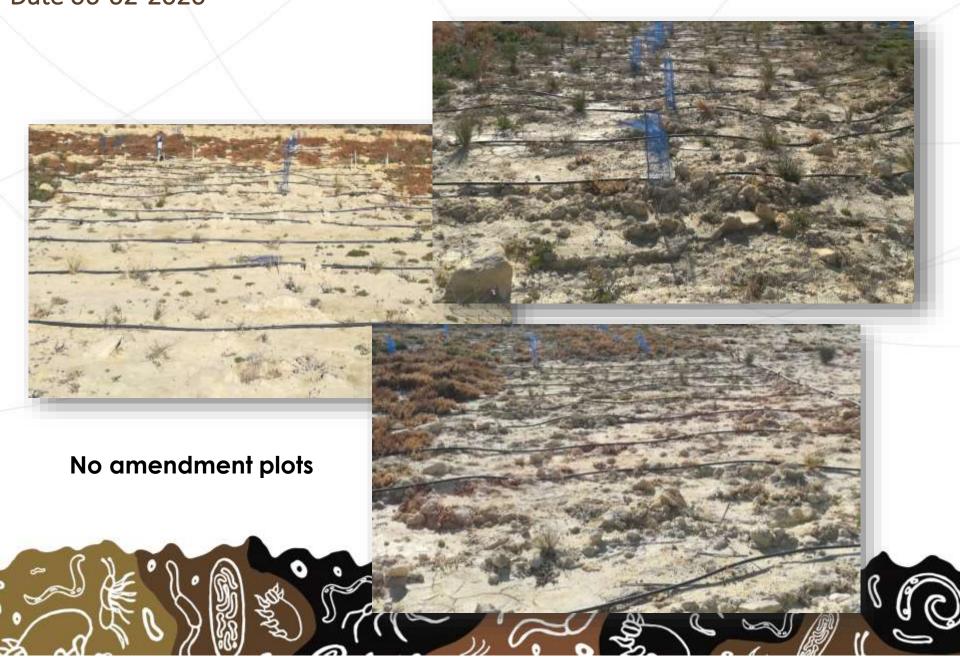


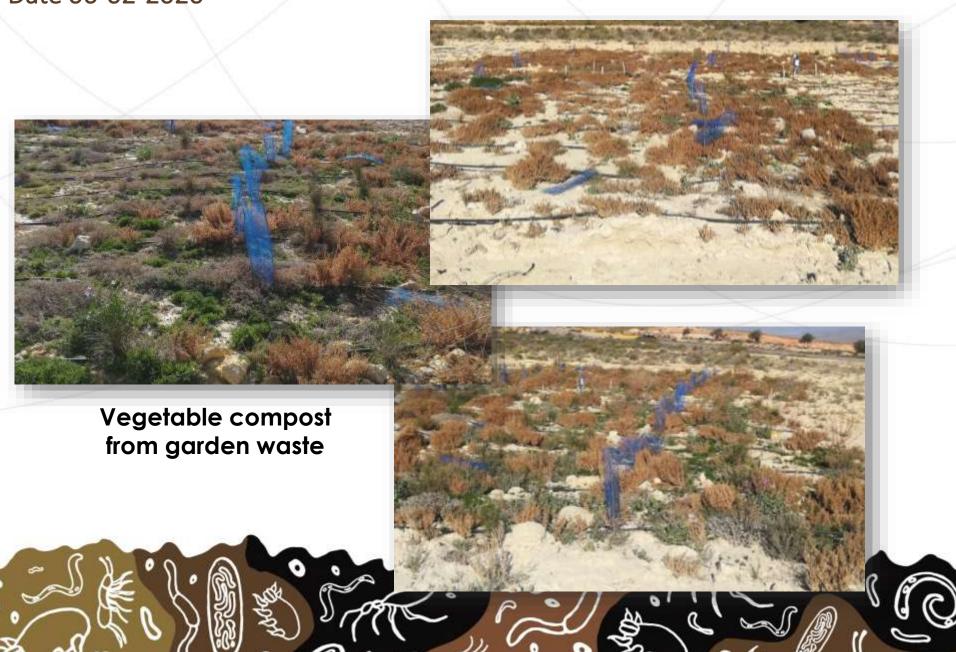


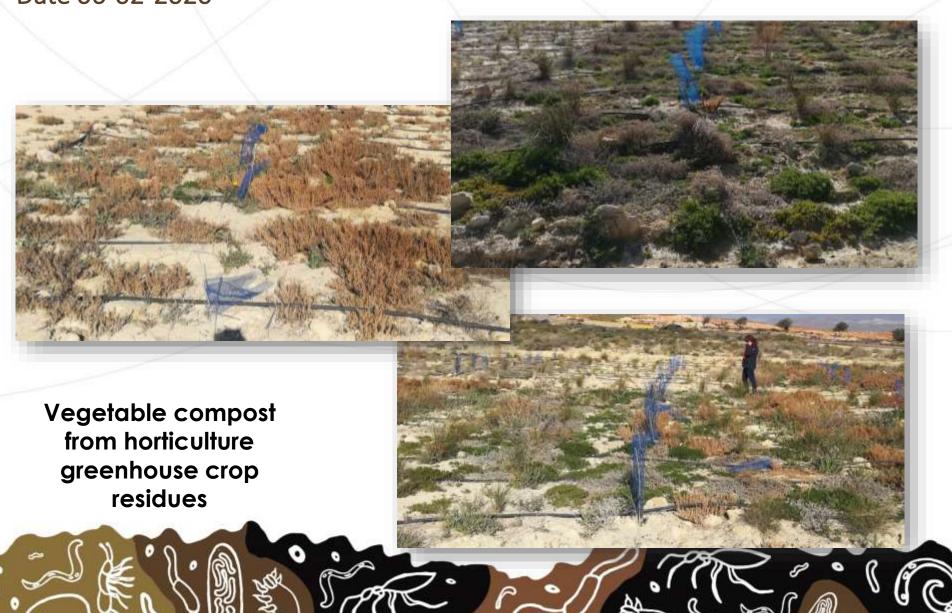




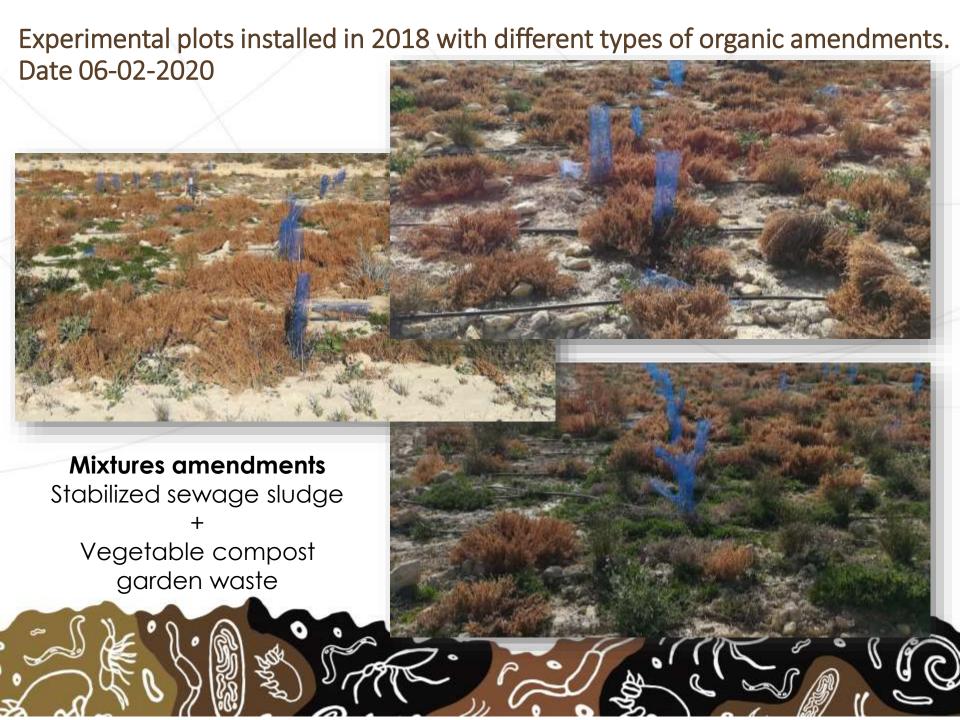








Experimental plots installed in 2018 with different types of organic amendments. Date 06-02-2020 Stabilized sewage sludge







The preliminary results indicate that there are short-term changes in bacterial communities and their relationships with carbon emissions and sequestration, so we think this is an interesting line of thought for restoring fragile arid ecosystems to promote their functionality and improve their global change disease.





Thank you for your attention

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