## Shifts in the bacterial diversity of aerobic and anaerobic bentonite microcosms treated with U and G2P

C. Povedano-Priego<sup>1\*</sup>, F. Jroundi<sup>1</sup>, M. Lopez-Fernandez<sup>2</sup>, I. Martín-Sánchez<sup>1</sup>, M. Dopson<sup>3</sup> and M. L. Merroun<sup>1</sup>

- <sup>1</sup> Department of Microbiology, Faculty of Science, University of Granada, 18071 Granada, Spain (\*correspondence: ppriego@ugr.es)
- <sup>2</sup> Institute of Resource Ecology, Helmholtz-Zentrum Dresden-Rossendorf, Bautzner Landstraße 400, 01328 Dresden, Germany
- <sup>3</sup> Centre for Ecology and Evolution in Microbial Model Systems (EEMiS), Linnaeus University, 392 31 Kalmar, Sweden

Microorganisms can potentially disturb the migration of radionuclides through processes such as biosorption, biomineralization, bioaccumulation and biotransformation. These processes would occur in deep geological repository (DGR), if radionuclides were accidentally leaked from nuclear wastes into the environment. To simulate this situation, aerobic and anaerobic microcosms of Spanish bentonite amended with uranium and glycerol-2-phosphate (G2P) were studied.

Changes in the microbial communities after six months of incubation were shown using Next Generation Sequencing (NGS) based on Illumina technology. Under aerobic conditions, uranium enhanced the growth of specific microorganisms with the potential to interact with such radionuclide like Desulfomicrobium, Burkholderia and Bacillus. In addition, in G2P-uranyl-treated microcosms Amycolatopsis was enriched suggesting a U biomineralization process and STEM-EDS analyses showed U-phosphates. In anaerobic microcosms, dominant genera such as Desulfatiglans (a sulfate-reducing bacterium), and sulfuroxidizing bacteria (Sulfurimonas and Thiobacillus) were identified in U microcosms, while Pseudomonas and Desulfovibrio, were abundant in the U-G2P microcosms. Both are described for their ability to immobilize U as U phosphates through biomineralization and by enzymatic reduction of U(VI) to U(IV) (Desulfovibrio). The outputs of this study would help to predict the impact of microbial processes on the DGR long-term performance.

This work has been supported by the project CGL2014-59616-R and the grant FPU 14/04263 from "Ministerio de Educación Cultura y Deporte".