

Ironborn: Fe distribution in geothermal fluids and its influence on the biosphere

BERNARDO BAROSA¹, LUCA TONIETTI^{1,2}, ALESSIA BASTIANONI¹, MATTEO SELCI¹, MARTINA CASCONI¹, FELICIANA OLIVA¹, DEBORAH BASTONI¹, FRANCESCO MONTEMAGNO¹, ANNARITA RICCIARDELLI¹, DAVIDE CORSO¹, MONICA CORREGGIA¹, LUCIANO DI IORIO¹, PETER H BARRY³, DAVID V BEKAERT³, SÆMUNDUR A. HALLDORSSON⁴, ANDRI STEFÁNSSON⁵, KAREN G LLOYD⁶, GERDARD JESSEN⁷, AGOSTINA LAURA CHIODI⁸, MAARTEN DE MOOR^{9,10}, CARLOS RAMÍREZ-UMAÑA¹¹, MUSTAFA YUCEL¹², MATT SCHRENK¹³, ALESSANDRA ROTUNDI^{2,14}, ANGELINA CORDONE¹ AND DONATO GIOVANNELLI^{1,15,16,17,18}

¹University of Naples Federico II

²Parthenope University of Naples

³Woods Hole Oceanographic Institution

⁴Nordic Volcanological Center, Institute of Earth Sciences, University of Iceland

⁵University of Iceland

⁶The University of Tennessee, Knoxville

⁷Austral University of Chile

⁸Instituto de Bio y Geociencias del Noroeste Argentino (IBIGEO, CONICET-UNSA), Salta, Argentina

⁹National University

¹⁰University of New Mexico

¹¹Servicio Geológico Ambiental de Costa Rica (SeGeoAm), San José, Costa Rica

¹²Middle East Technical University, Institute of Marine Sciences

¹³Michigan State University

¹⁴Istituto di Astrofisica e Planetologia Spaziali di Roma

¹⁵Woods Hole Oceanographic Institution

¹⁶CNR-IRBIM, National Research Council

¹⁷Rutgers University

¹⁸Earth-Life Science Institute

Presenting Author: bernard.barosa@gmail.com

In geothermal environments, geochemistry often determines the boundary conditions within which life operates¹. Recently, studies have demonstrated how different geochemical environments, sampled through deeply-sourced seeps across various geological settings, constrain the distribution of microbial taxa, as well as the genes associated with relevant metabolic functions^{2,3}. However, the role of geothermal environments and trace element availability in driving protein distribution remains poorly understood⁴. Iron is the 4th most abundant element in Earth's crust and one of the most fundamental elements for biological systems, as it is associated with numerous metabolic reactions across all the various domains of life. For this reason, iron is tightly coupled to the evolution of the biosphere over deep time⁵. Additionally, the

availability of iron in geothermal environments within different geological settings (e.g., variable rock types and tectonic contexts) is still poorly constrained. In the present study, we apply a novel sampling strategy across large-scale geological gradients⁶ encompassing geothermal environments in different tectonic settings, including Costa Rica, Argentina, Chile, and Italy arc/back-arc systems, as well as the Iceland spreading center. Through high resolution geochemical analysis, coupled with metagenomics on more than 50 deeply-sourced springs, we constrain the presence, diversity, and abundance of iron-related proteins to geothermal iron availability within different geological settings.

References

1. Reysenbach, A.-L. & Shock, E. Merging Genomes with Geochemistry in Hydrothermal Ecosystems. *Sci*, 2002
2. Fullerton, K. M. *et al.* Effect of tectonic processes on biosphere–geosphere feedbacks across a convergent margin. *Nat. Geosc.* (2021).
3. Rogers, T. J. *et al.* Chemolithoautotroph distributions across the subsurface of a convergent margin. *ISME J*, 2023
4. Srivastava, S., Dong, H. & Briggs, B. R. The Effect of Spring Water Geochemistry on Copper Proteins in Tengchong Hot Springs, China. *Appl. Env. Micro* 2020
5. Wade, J., Byrne, D. J., Ballentine, C. J. & Drakesmith, H. Temporal variation of planetary iron as a driver of evolution. *Proc. Natl. Acad. Sci.* 2021
6. Giovannelli, D. *et al.* Sampling across large-scale geological gradients to study geosphere–biosphere interactions. *Front. Micro* 2022