

Insights: Publications

# Draft Genome Sequence of *Salinivibrio* sp. Strain EAGSL, a Biotechnologically Relevant Halophilic Microorganism

*Microbiology Resource Announcements* 9 (43), e01020-20

October 22, 2020

Written by

---

The halophilic bacterium *Salinivibrio* sp. strain EAGSL was isolated from the Great Salt Lake (Utah) for use in microbial electrochemical technologies experiencing fluctuating salt concentrations. Genome sequencing was performed with Ion Torrent technology, and the assembled genome reported here is 3,234,770 bp with a GC content of 49.41%.