Relative abundance of analyzed samples Euryarchaeota Gemmatimonadota Halobacterota Methylomirabilota Verrucomicrobiota Acidobacteriota Actinobacteriota Bacteroidota Chloroflexi Desulfobacterota Myxococcota Proteobacteria 10.0 7.5 5.0 2.5 10.0 Relative abundance SITE_ID 🛱 Ag1 Ag2 Ag3 10.0 7.5 5.0 Ag2-Ag3-Site Ag3 Ag2 Ag3 Ag3 Ag3 Ag2 Ag3 Ag1

Ag1

Ag1

Ag2

Ag1

Ag2

Ag3

Ag3

Ag3

Ag2

Ag1

Ag3

Ag1

Ag2

Ag1

Ag2

Ag1

Ag3

Ag1

Ag3

Ag2

Ag1

Ag2

Ag1

Ag1