CORRIGENDA

Microbial ecology of an Antarctic hypersaline lake: genomic assessment of ecophysiology among dominant haloarchaea

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Correction to: *The ISME Journal* (2014) 8, 1645–1658; doi:10.1038/ismej.2014.18; published online 20 February 2014

Since the publication of this article, the authors have noticed an error in Supplementary Table S2. The correct table is shown below.

The error has now been rectified, and the correct Supplementary Information accompanies the online version of the original article.

The authors apologize for any inconvenience this may have caused.

Supplementary Table S2 Concentrations of analytes in Deep Lake at four different depths

Analyte	Depth			
	5 m	13 m	24 m	36 m
Ammonia (mg-NL ⁻¹)	0.616*	0.305	1.12	0.338
Nitrate (mg-NL ⁻¹)	0.066*	0.073	0.057	0.082
Nitrite (mg-NL ⁻¹)	0.002*	0.002	0.002	0.003
Nitrogen, total (mg-NL ⁻¹)	5.3*	5.3	5.1	6.8
Phosphorus, total (mg-PL ⁻¹)	0.656*	0.678	0.657	0.972
Phosphorus, dissolved reactive (mg-PL ⁻¹)	0.124*	0.049	0.222	0.055
DOC (mg L ⁻¹)	65*	57	57	63
TOC (mg L ⁻¹)	59*	57	53	59
Sulfur, dissolved (mg L ⁻¹)	963*	903	862	1080
Sulfur, total (mg L ⁻¹)	904*	899	974	1110

*Average of two samples.

A novel approach, based on BLSOMs (Batch Learning Self-Organizing Maps), to the microbiome analysis of ticks

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Since the publication of this article, it has come to the attention of the authors that there were errors contained within the ‘Results’

and the ‘Discussion’ sections, namely that sequencing data derived from two tick species (IO and IP) were incorrectly transposed. In these two sections, IO should be read as IP and *vice versa*.

The Authors apologise for any inconvenience this may have caused.