

Additional File 2: Metagenomic sequence read statistical quality parameters.

Object Name	Buhera Soda Pan Metagenomic Sequence
Number of Reads	35,876,873
Total Number of Bases	3,587,687,300
Mean Read Length	100.0
Base quality score (range)	34 – 36
Mean base quality score	35.45
Read quality score (range)	24 – 38
Mean read quality score	36.0
GC Percentage	52.08%