

Long-Term Analysis of Resilience of the Oral Microbiome in Allogeneic Stem Cell Transplant Recipients

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Table S1. Sequencing output control samples.

| mean ctrl reads per sample | mean exp reads per sample | ratio control/total control reads | ratio exp/total exp reads | taxonomy |
|----------------------------|---------------------------|-----------------------------------|---------------------------|---|
| 3.72 | 0.16 | 0.165 | 9.52E-06 | f__Comamonadaceae |
| 2.69 | 0.07 | 0.119 | 4.21E-06 | f__Comamonadaceae; g__Leptothrix; s__sp_oral_taxon_025 |
| 1.69 | 0 | 0.075 | 0 | c__Alphaproteobacteria |
| 1.45 | 0.08 | 0.064 | 4.39E-06 | f__Caulobacteraceae; g__Caulobacter; s__sp_oral_taxon_002 |

| | | | | |
|------|------|-------|----------|---|
| 1.41 | 0.08 | 0.063 | 4.58E-06 | f__Bradyrhizobiacea e; g__Bosea; s__vestrisii |
| 1.14 | 0.06 | 0.050 | 3.30E-06 | f__Burkholderiaceae ; g__Ralstonia; s__sp._oral_taxon_4 06 |
| 1.03 | 0.05 | 0.046 | 2.75E-06 | c__Alphaproteobact eria |
| 1.03 | 0.02 | 0.046 | 9.15E-07 | f__Burkholderiaceae |
