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Table S2 – mRNA raw read count attributed to *Lactobacillus* spp. genomes in carious root surfaces

| Genomes | Samples | | | | | | | | |
|-----------------------|---------|------|----------|--------|--------|------|--------|--------|--------|
| | RC_7 | RC_8 | RC_A | RC_B | RC_D | RC_E | RC_F | RC_G | RC_H |
| <i>L. acidophilus</i> | 0 | 0 | 1.78 | 308.77 | 3.13 | 0 | 4.73 | 529.72 | 3.28 |
| <i>L. brevis</i> | 0 | 0 | 2.10 | 5.02 | 1.23 | 0 | 0 | 2.41 | 2.91 |
| <i>L. buchneri</i> | 0 | 0 | 48.41 | 5.10 | 10.08 | 4.23 | 1.13 | 0 | 4.95 |
| <i>L. casei</i> | 0 | 0 | 23.45 | 2.11 | 6.86 | 0 | 0 | 0 | 4.99 |
| <i>L. crispatus</i> | 0 | 0 | 6.97 | 226.55 | 120.56 | 0 | 153.44 | 466.70 | 75.02 |
| <i>L. curvatus</i> | 3.56 | 0 | 13.75 | 9.35 | 6.73 | 0 | 1.22 | 0 | 38.03 |
| <i>L. delbrueckii</i> | 0 | 0 | 3,582.72 | 4.51 | 0 | 0 | 53.89 | 2.06 | 9.13 |
| <i>L. fermentum</i> | 51.29 | 0 | 1,664.80 | 310.76 | 282.33 | 8.18 | 5.79 | 5.85 | 258.08 |
| <i>L. gasseri</i> | 1.63 | 0 | 1,166.59 | 135.77 | 70.52 | 7.57 | 142.89 | 33.04 | 474.36 |
| <i>L. jensenii</i> | 0 | 0 | 6.00 | 7.69 | 0 | 0 | 0 | 6.73 | 0 |
| <i>L. johnsonii</i> | 0 | 0 | 133.04 | 21.5 | 2.68 | 0 | 7.49 | 15.98 | 5.08 |
| <i>L. paracasei</i> | 0 | 0 | 17.73 | 4.00 | 64.06 | 6.08 | 0 | 0 | 6.61 |
| <i>L. plantarum</i> | 0 | 0 | 2.78 | 0 | 0 | 6.64 | 0 | 0 | 0 |
| <i>L. rhamnosus</i> | 0 | 0 | 600.01 | 0 | 139.55 | 0 | 7.88 | 0 | 203.08 |
| <i>L. salivarius</i> | 39.42 | 0 | 50.72 | 50.72 | 34.33 | 0 | 17.69 | 17.07 | 162.72 |

Data normalized considering total number of transcripts divided by the total number of genes in each sample.