|  |
| --- |
| **Table S2.** **Sequencing summary.** *\** |
| **Parameter** | **Human milk (n=67)** | **Neonatal stool (n=67)** |
| Number of readsMean of reads per sampleLowest sample readsHighest sample reads  | 4’240,314 63,288 5,416 313,468 | 5’335,223 79,630 10,919 342,510 |
| Mean length for sequences\* | 167.16 bases ± 48.35 | 170.94 bases ± 44.14 |
| Total of raw read sequences: 9’575,537; Mean lengths: 169.3 ± 46.1.*\*Summary of sequencing after trimming at 200 nt.*  |