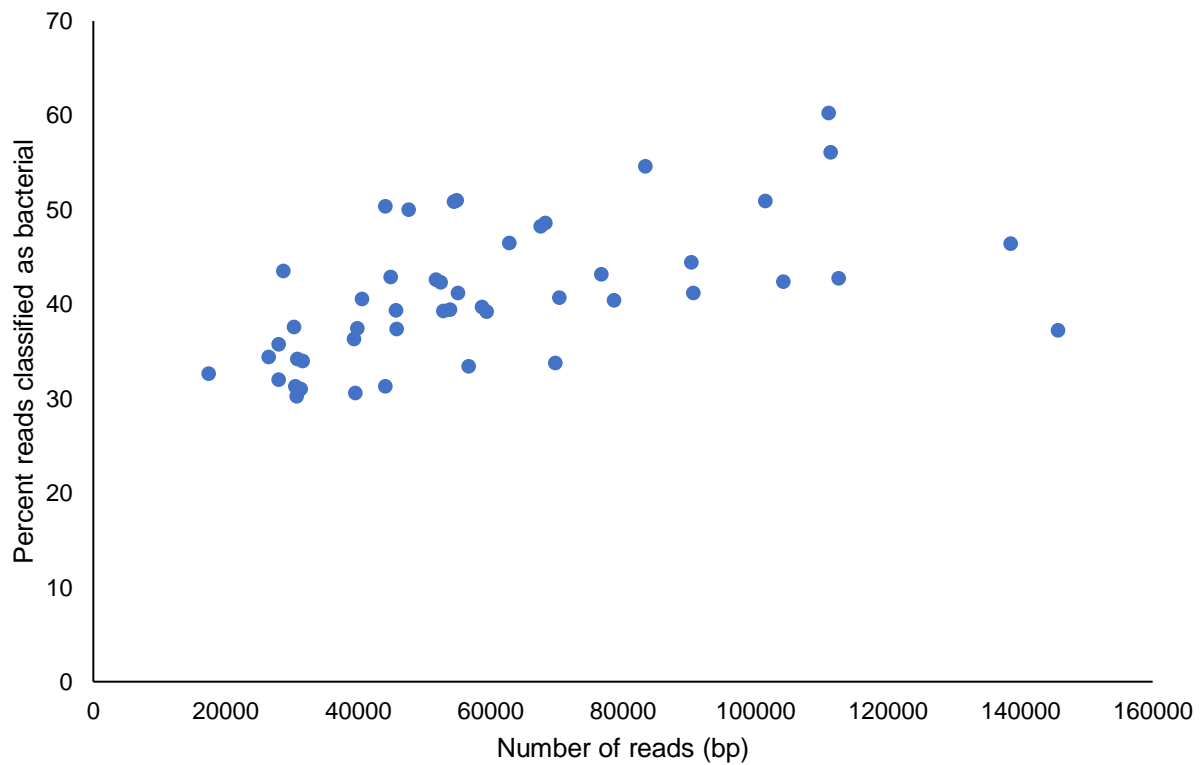


Supplementary File



Supplementary Figure 3. Percentage of metagenomic reads classified as bacterial and the number of reads classified via Kraken2 (n = 48)