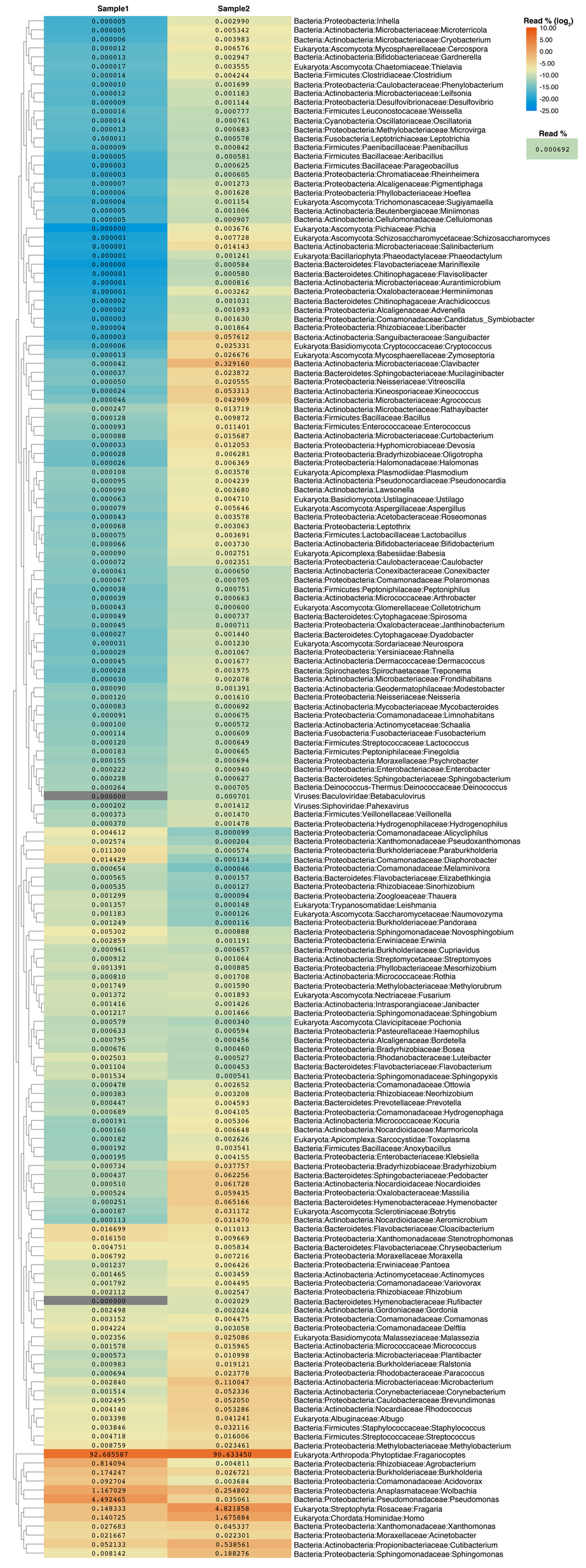Supplementary Fig. S2. Taxonomic classification of two metagenomic samples of the gall-inducing mite Fragariocoptes setiger in Kraken2. There was a total of 708,046,814 and 82,009,061 classified reads in samples 1 and 2, respectively. OTUs (genera) were filtered based on a normalized abundance threshold of ≥0.0005% in either sample, resulting in 171 OTUs represented by 670,717,361 and 72,439,919 reads (sample 1 and 2, respectively). Data are given in Supplementary Table S3.