

Supplementary Fig. S1. Taxonomic classification of Bacteria in two metagenomic of the gall-inducing mite *Fragariocoptes setiger* in Kraken2. There was a total of 49,514,852 and 3,830,185 classified bacterial reads in samples 1 and 2, respectively. OTUs (genera) were filtered based on a normalized abundance threshold of $\geq 0.0338\%$ in either sample, resulting in 83 OTUs (genera) represented by 46,906,755 and 1,895,438 reads (sample 1 and 2, respectively). Data are given in Supplementary Table S2.

