



Fig. S3. ASVs that significantly differed in abundance in snail microbiota based on reproductive mode or sex. Points represent significantly differentially abundant taxa and are plotted by \log_2 fold change in abundance (ANCOM; adj- $p < 0.01$). Comparisons are asexual/sexual and male/female. There were no significantly differentially abundant taxa between infected and uninfected snails. Figure is labelled by phyla and deepest available taxonomy, where G = genus and F = family. Snails are only from NZ.