

Fig. S7. Functional clustering of bacterial MAGs. Each row represents a MAG, each column is a KEGG module. The bars on the left show taxonomic position of the MAG on three levels (family, order, and phylum) and the cluster it was assigned to by three clustering methods (apcluster, kmeans, and hdbscan). The dendrograms show the hierarchical clustering done using the hclust function in R. For this analysis, we only used MAGs with at least 90% completeness.

