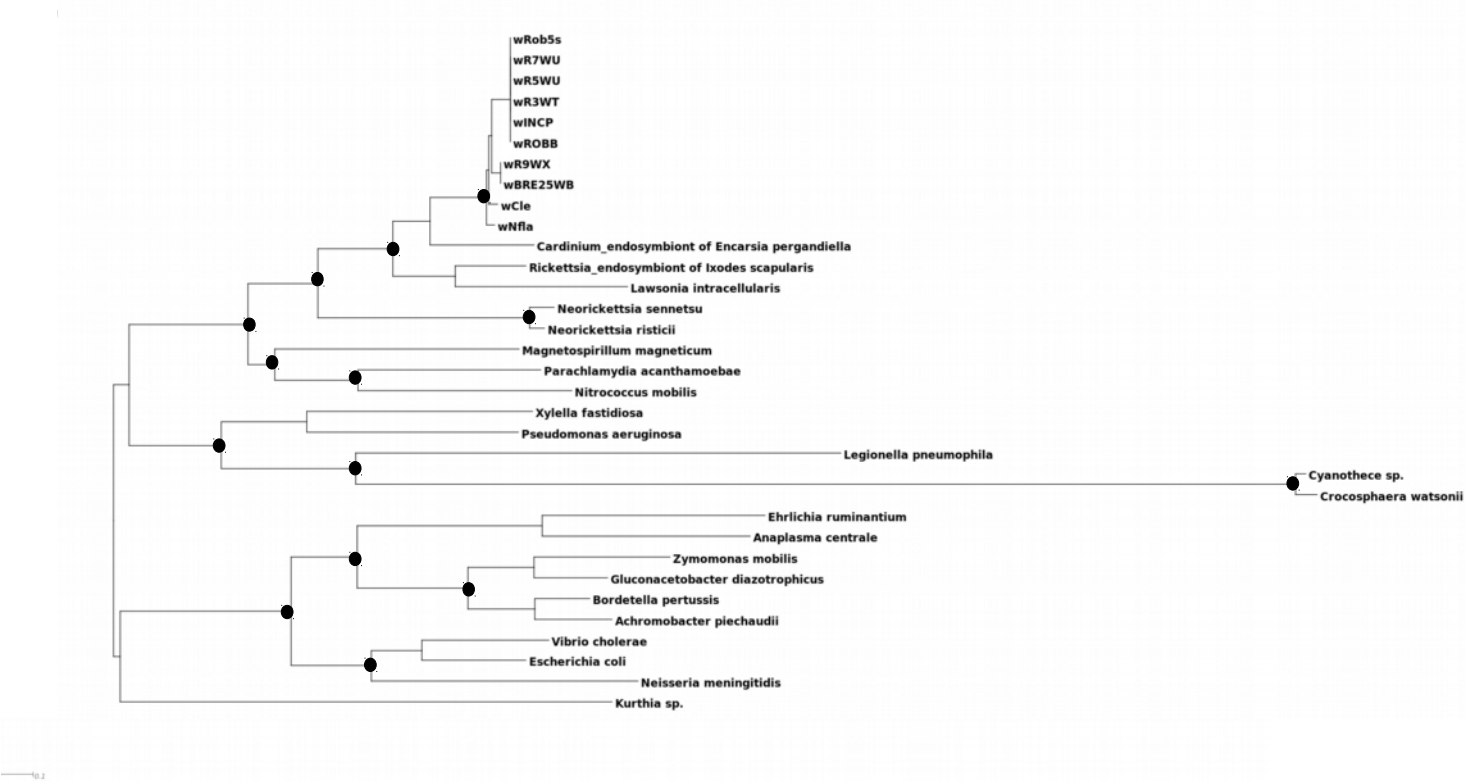
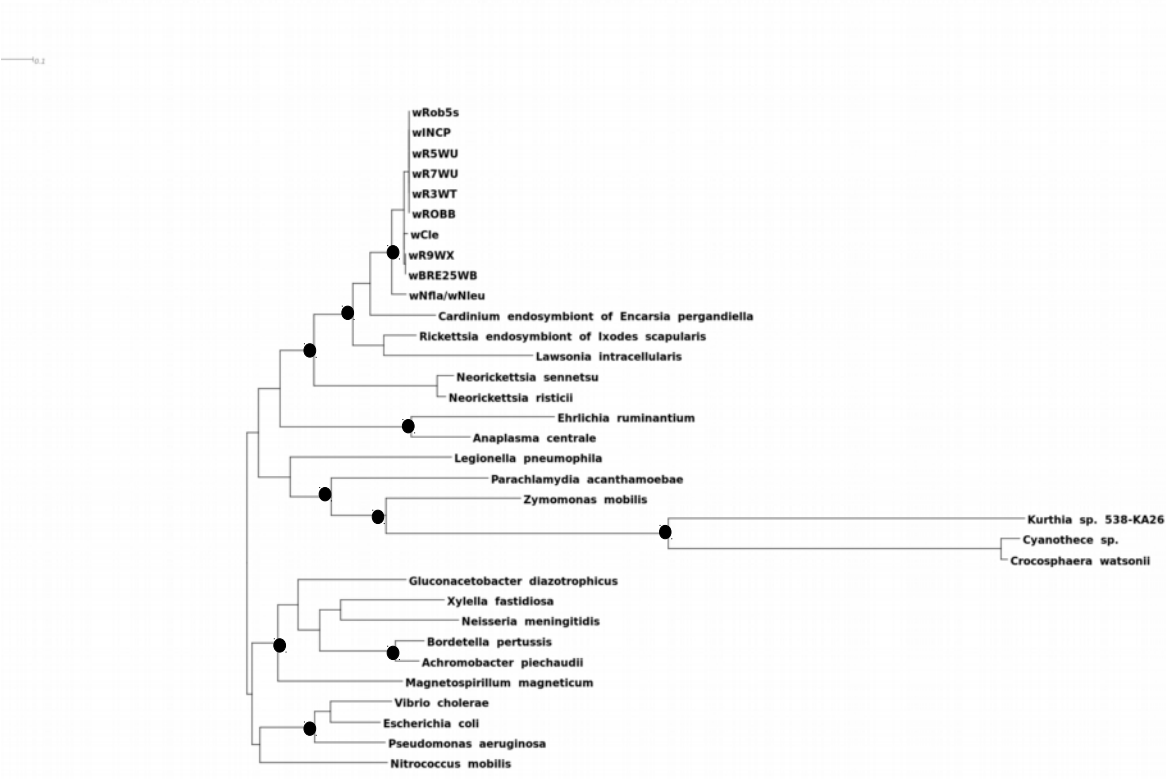


Supplementary Figure 3 : Maximum-likelihood phylogenies of the biotin genes. Black circles at nodes in the phylogeny indicate support values >99%

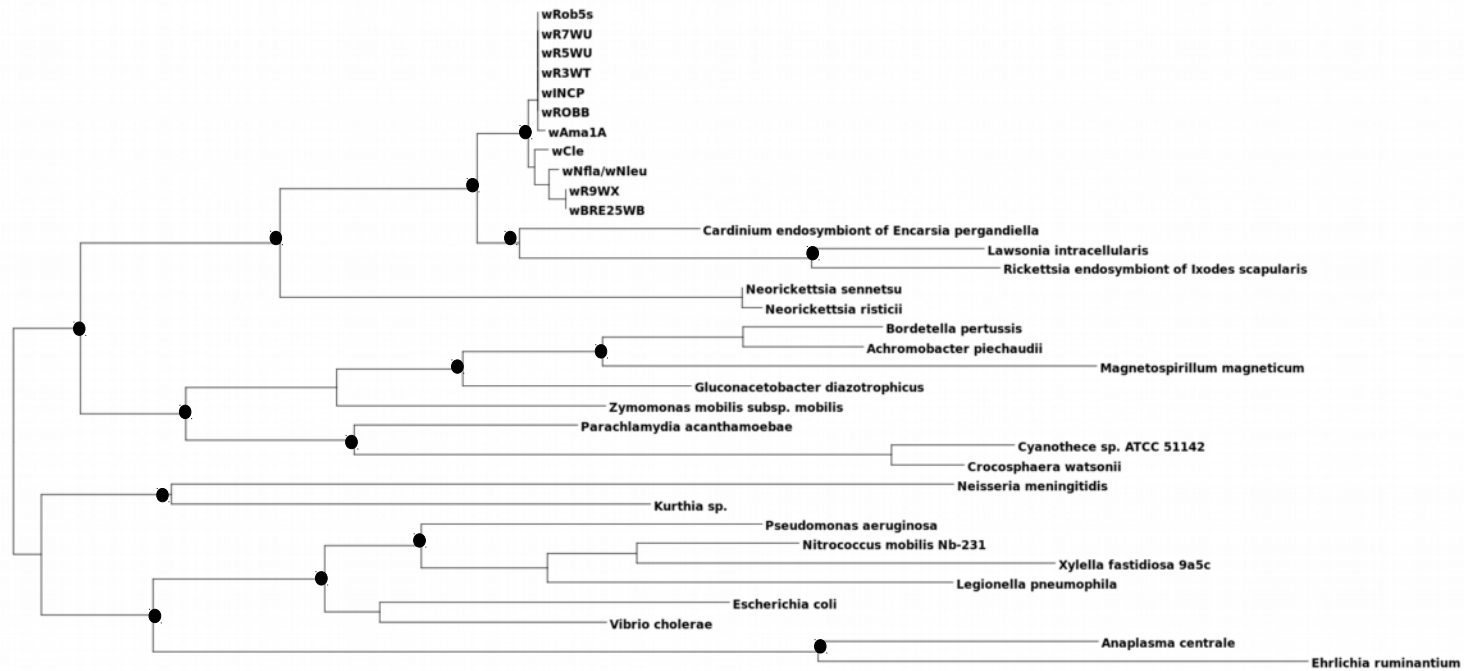
bioA



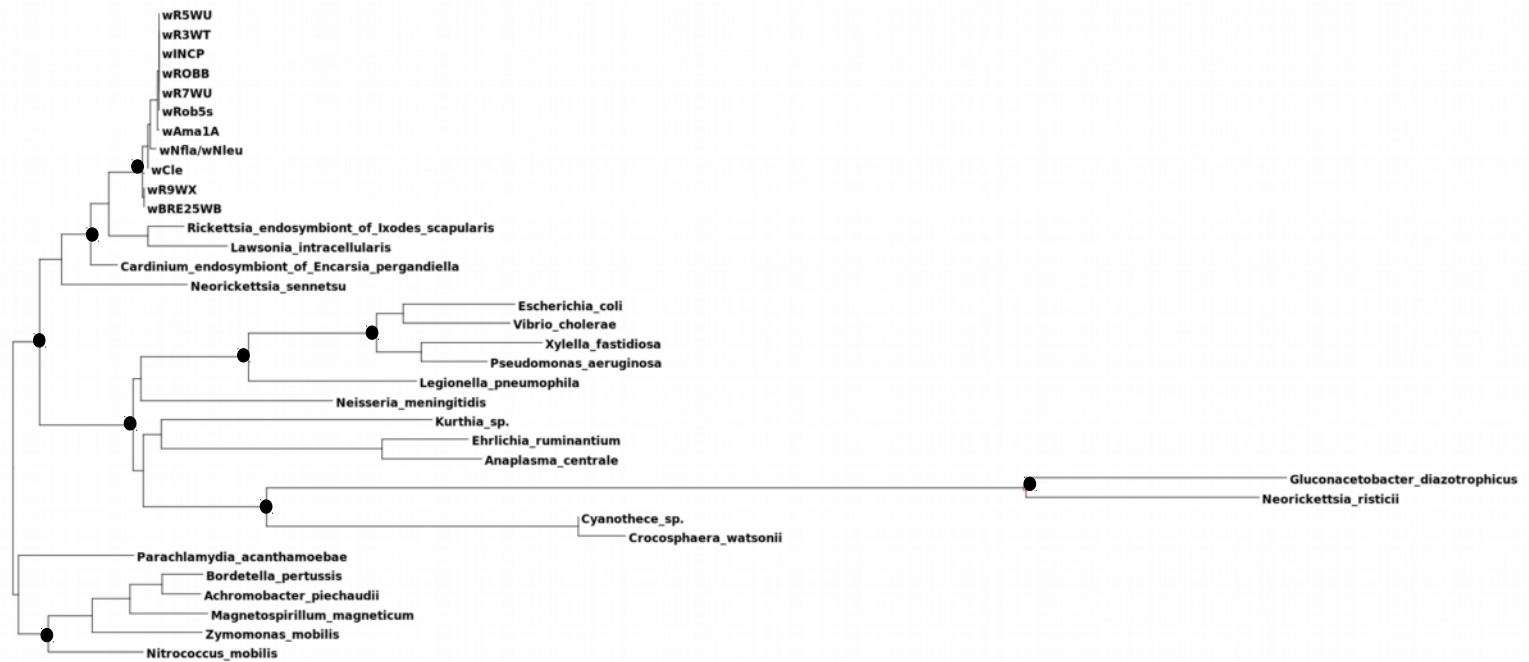
bioB



bioC

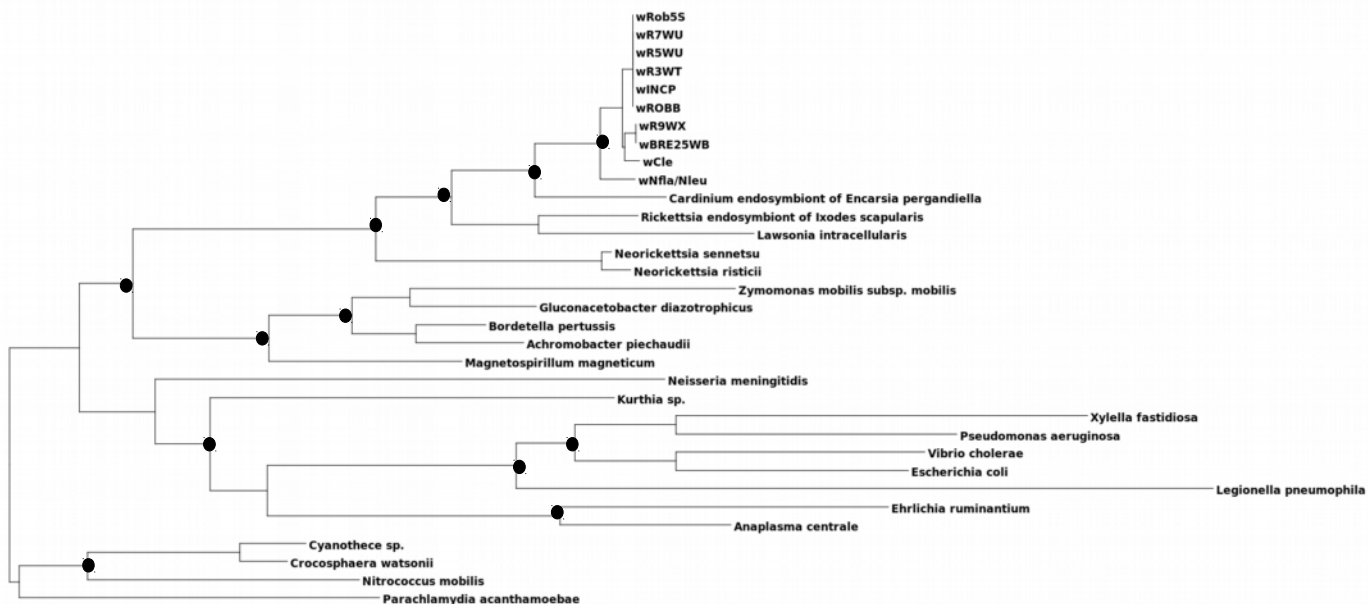


bioD



bioF

0.2



bioH

0.2

