

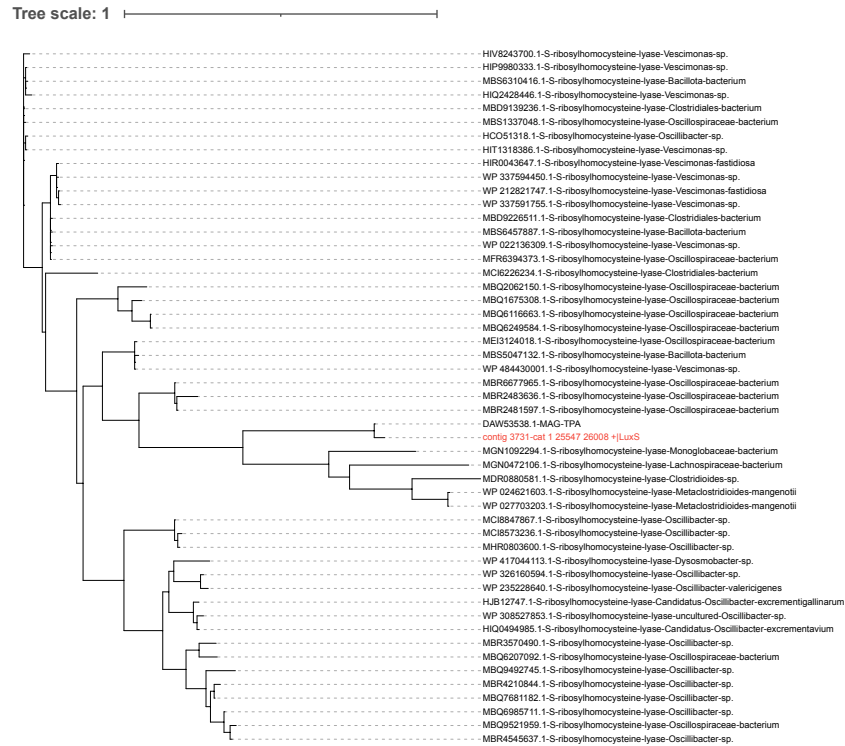


Fig S6. Maximum-likelihood phylogenetic tree of LuxS proteins encoded by phage genomes and their closest reference homologs.