

Tree scale: 0.1

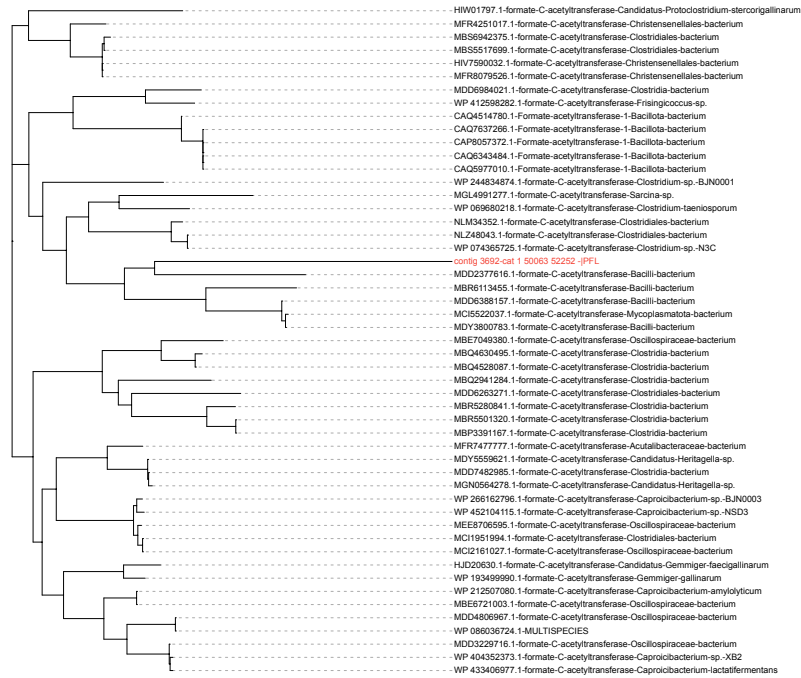


Fig S4. Maximum-likelihood phylogenetic tree of PFL proteins encoded by phage genomes and their closest reference homologs.