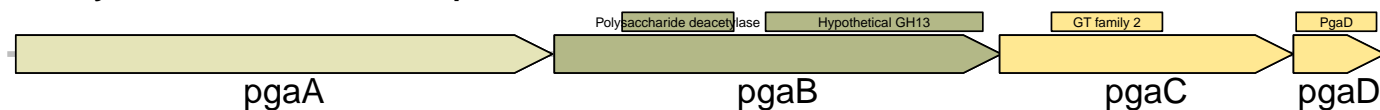
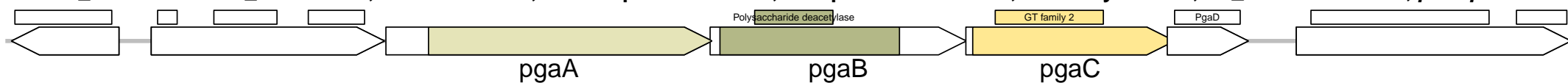


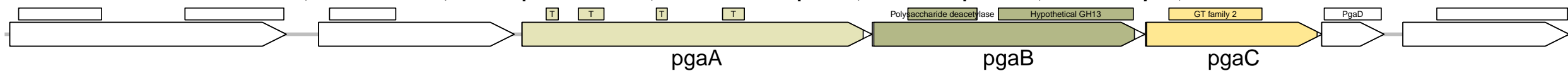
Query, Proteobacteria, Gammaproteobacteria, Enterobacteriales, Enterobacteriaceae, *Escherichia, coli*



OdNE_18-Q3-R46-58_BAT3C.415, Proteobacteria, Gammaproteobacteria, Betaproteobacteriales, Rhodocyclaceae, *Ca_Accumulibacter, phosphati*



Kalu_18-Q3-R12-55_BAT3C.208, Proteobacteria, Gammaproteobacteria, Ectothiorhodospirales, Thioalkalspiraceae, *Thioalkalispira, midas_s_282*



0 3000 6000 9000

Function of Matched Query Gene



Export



Modification



Synthase