

Percent_identity

20

25

30

35

40

Function

Export

Polymerization

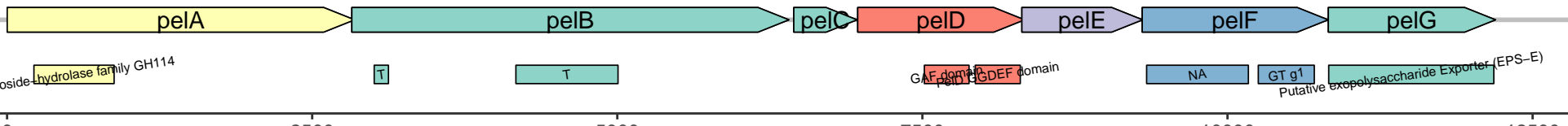
SY

Hydrolase

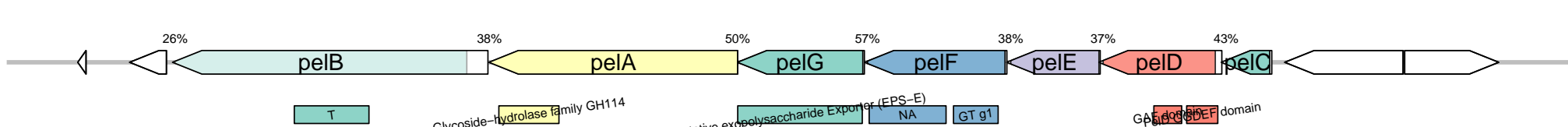
REG

NA

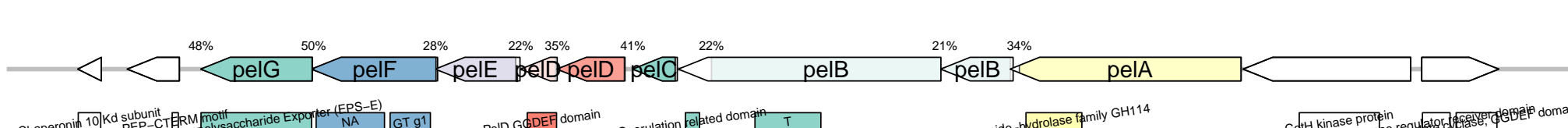
Query



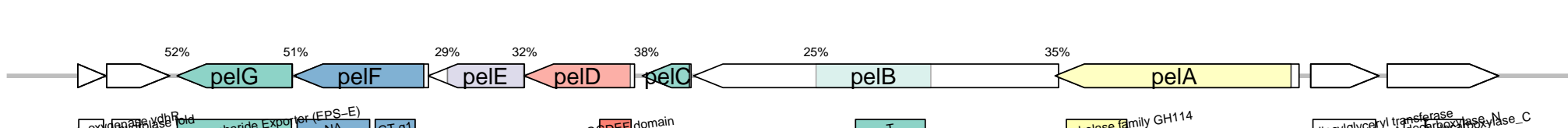
Aved_18-Q3-R54-62_BAT3C.540
Proteobacteria
Deltaproteobacteria
Myxococcales
P3OB-42
midas_g_6017
midas_s_601



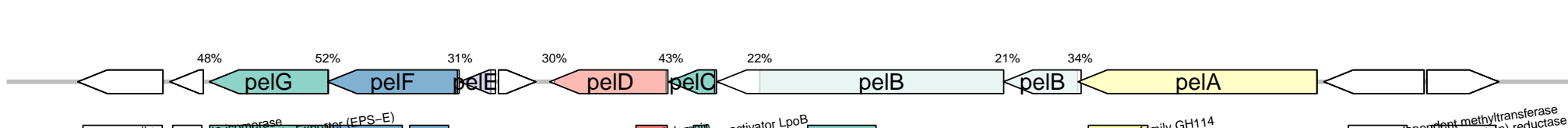
Bjer_18-Q3-R1-45_MAXAC.141
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Candidatus Symbiobacter
midas_s_30



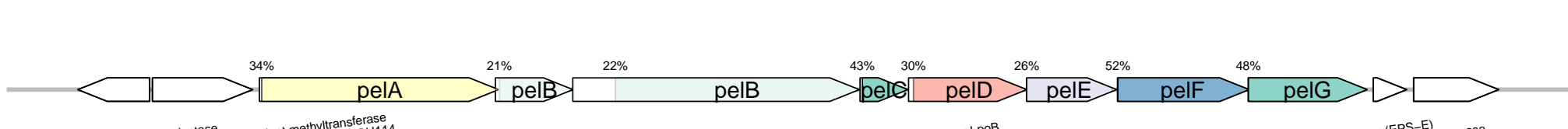
Hjor_18-Q3-R7-51_BATAC.292
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Lautropia
midas_s_13



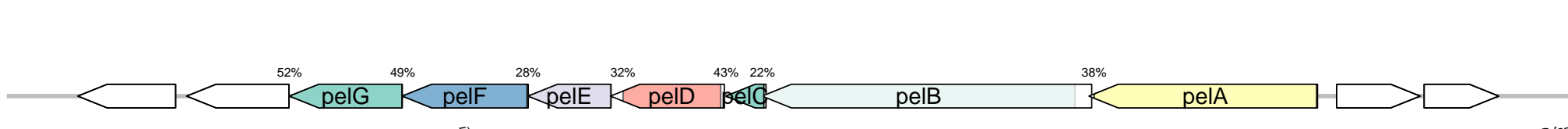
Bjer_18-Q3-R1-45_BATAC.569
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Limnohabibans
midas_s_283



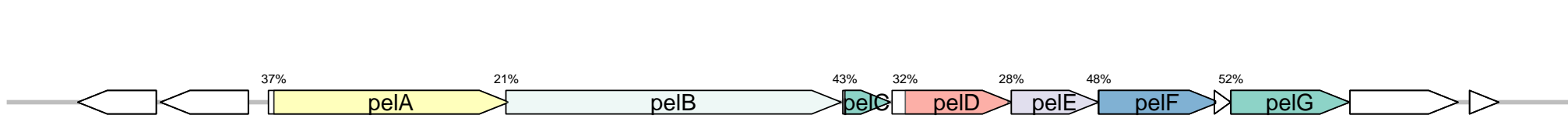
Lyne_18-Q3-R50-59_MAXAC.018
Proteobacteria
Gammaproteobacteria
Burkholderiaceae
Limnohabibans
midas_s_283



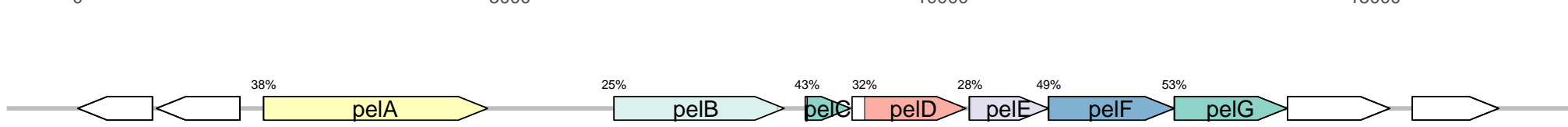
Ega_18-Q3-R5-49_BATAC.333
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Rhodoferritiferreducen
midas_s_283



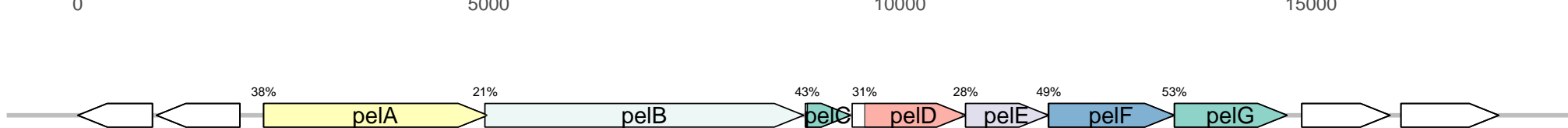
EsbE_18-Q3-R3-47_BATAC.414
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Rhodoferritiferreducen
midas_s_283



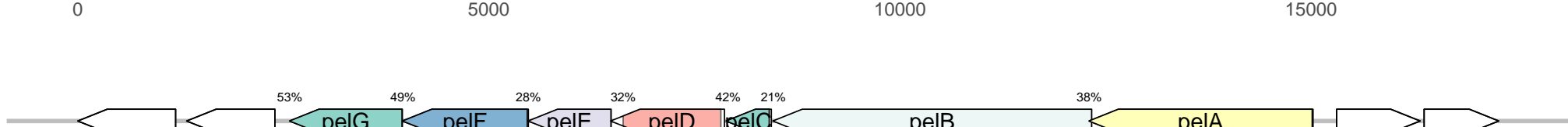
Viby_18-Q3-R106-67_MAXAC.046
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Rhodoferritiferreducen
midas_s_283



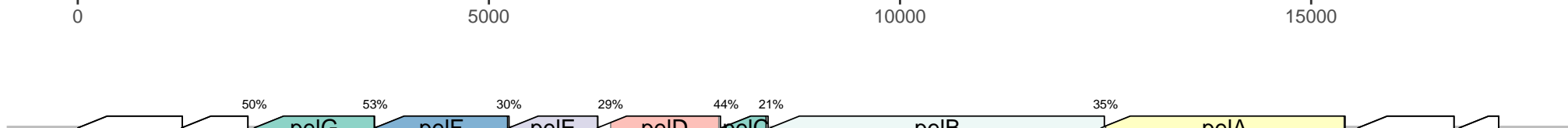
AalE_18-Q3-R2-46_BAT3C.116
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Rhodoferritiferreducen
midas_s_283



AalW_18-Q3-R10-53_MAXAC.061
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
Rhodoferritiferreducen
midas_s_283



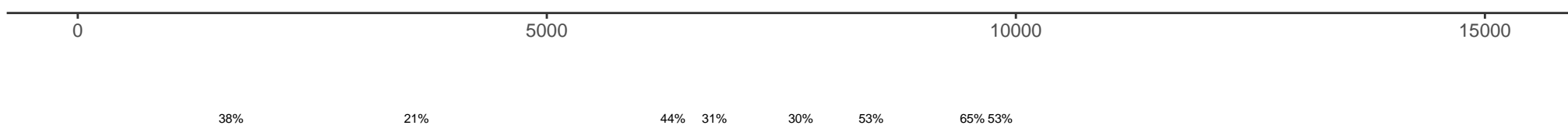
AalE_18-Q3-R2-46_BAT3C.211
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Ferrireducen
limneticus
midas_s_283



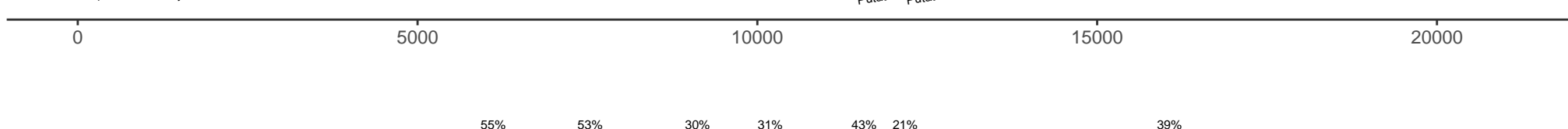
Ega_18-Q3-R5-49_MAXAC.071
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
Candidatus aalborgensis
midas_s_283



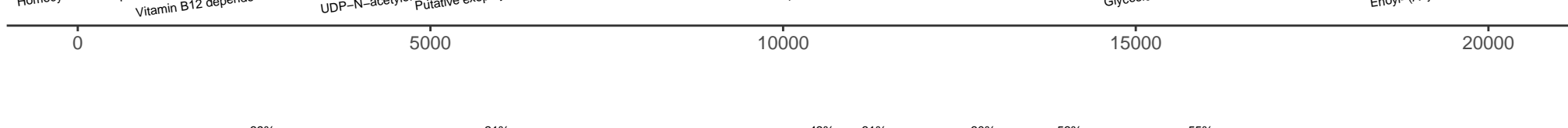
OdNE_18-Q3-R46-58_MAXAC.062_sub
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
Candidatus aalborgensis
midas_s_283



EsbE_18-Q3-R3-47_MAXAC.131
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
midas_s_42



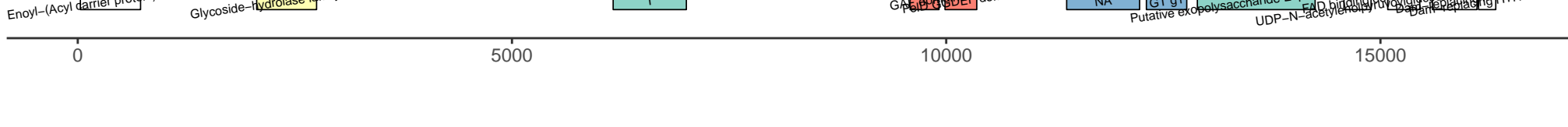
Hade_18-Q3-R52-61_MAXAC.304
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
midas_s_42



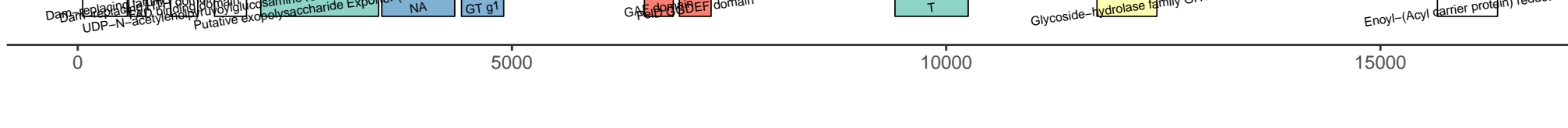
Hjor_18-Q3-R7-51_MAXAC.072
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
midas_s_42



Mari_18-Q3-R65-66_MAXAC.196
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
midas_s_42



Vibo_18-Q3-R45-57_BATAC.19
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Propionivibrio
midas_s_42



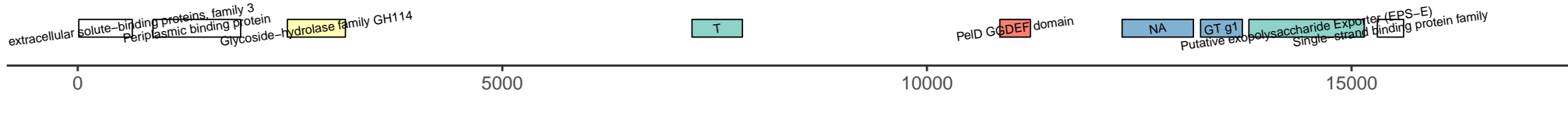
Aved_18-Q3-R54-62_MAXAC.403
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Sulfuritalea
midas_s_312



AalW_18-Q3-R10-53_MAXAC.406
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Rhodocyclaceae
Sulfuritalea
midas_s_312



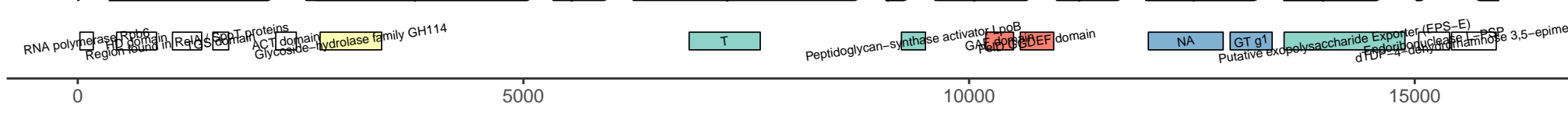
Kalu_18-Q3-R12-55_BATAC.288
Proteobacteria
Gammaproteobacteria
Oceanospirillales
Halellaceae
Halella
midas_s_97



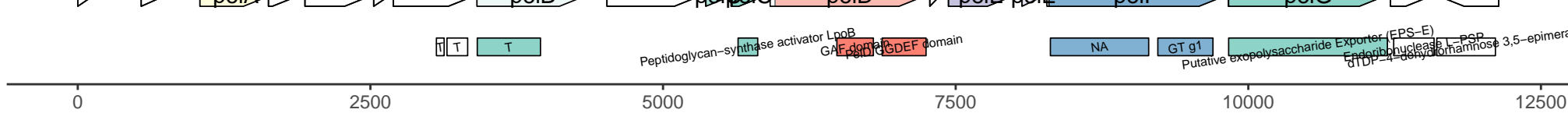
Aved_18-Q3-R54-62_BATAC.645_sub
Proteobacteria
Gammaproteobacteria
Pseudomonadales
Moraxellaceae
Agitococcus_lubricus_group
midas_s_317



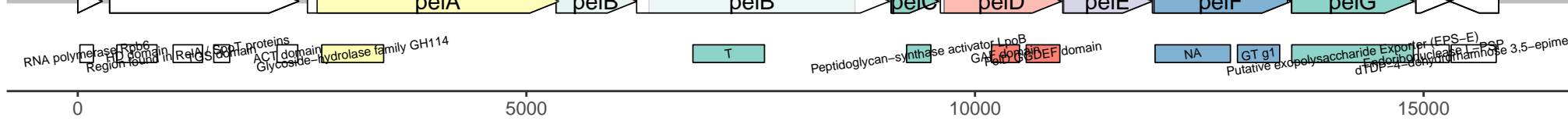
EsbE_18-Q3-R3-47_BATAC.309
Proteobacteria
Gammaproteobacteria
Pseudomonadales
Moraxellaceae
Agitococcus_lubricus_group
midas_s_317



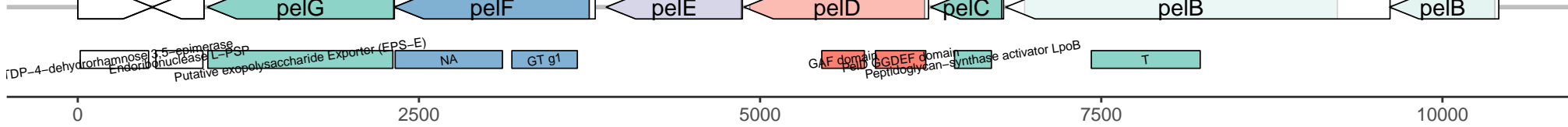
Hade_18-Q3-R52-61_BATAC.311
Proteobacteria
Gammaproteobacteria
Pseudomonadales
Moraxellaceae
Agitococcus_lubricus_group
midas_s_317



Lyne_18-Q3-R50-59_BAT3C.546
Proteobacteria
Gammaproteobacteria
Pseudomonadales
Moraxellaceae
Agitococcus_lubricus_group
midas_s_317



Vibo_18-Q3-R45-57_BAT3C.142
Proteobacteria
Gammaproteobacteria
Pseudomonadales
Moraxellaceae
Agitococcus_lubricus_group
midas_s_633



start