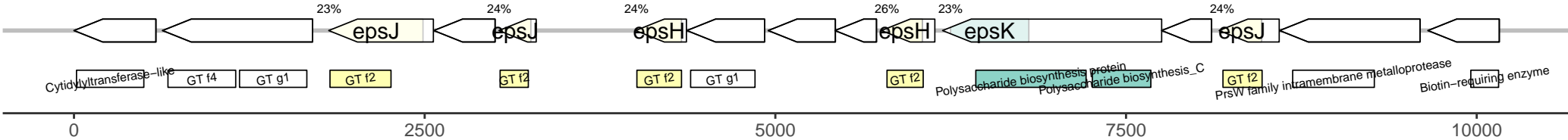
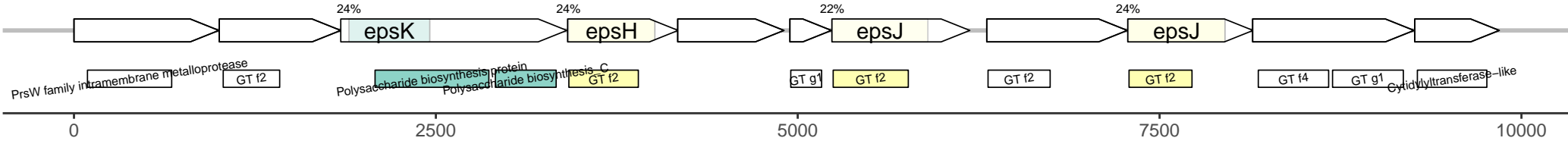


Percent_identity 20 25 30 35 40 Function Export GT NA

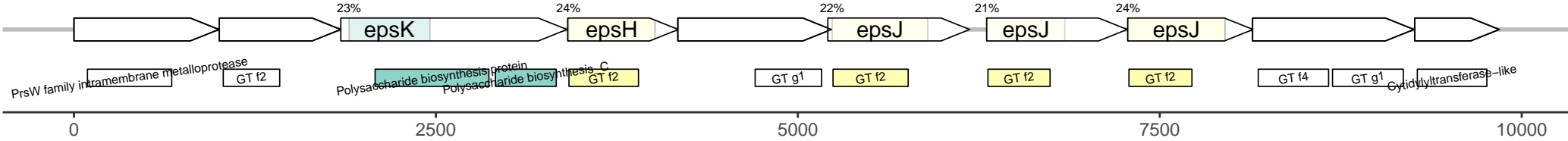
Aved_18-Q3-R54-62_MAXAC.030
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3



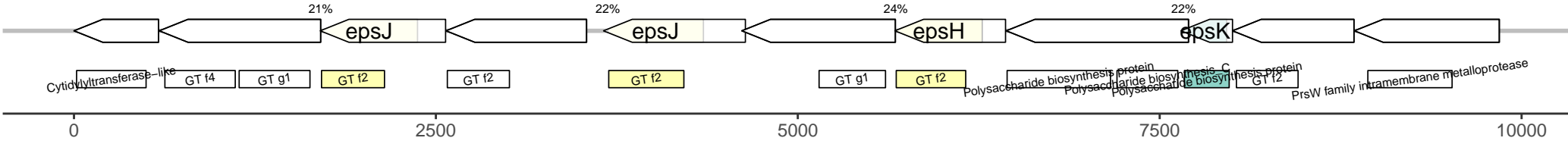
Damh_18-Q3-R51-60_BATAC.300
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3



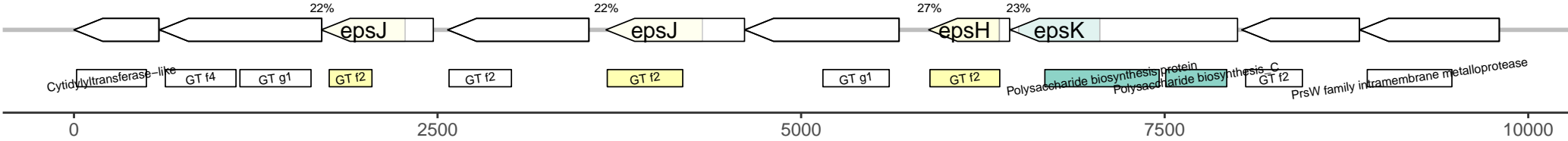
Ega_18-Q3-R5-49_BATAC.213
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3



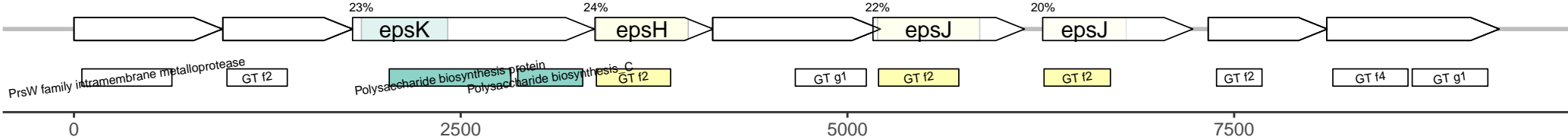
EsbW_18-Q3-R4-48_MAXAC.017
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3



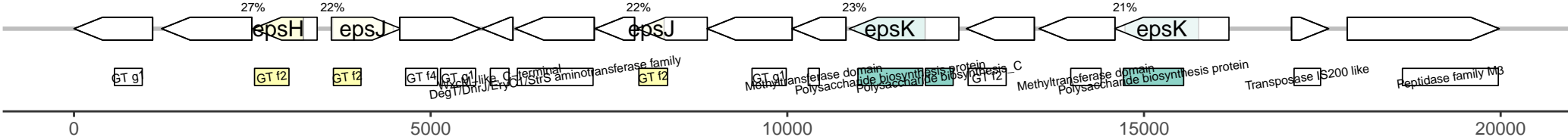
Hade_18-Q3-R52-61_MAXAC.028
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3



Mari_18-Q3-R65-66_MAXAC.025
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3



Mari_18-Q3-R65-66_MAXAC.226
Acidobacteria
Blastocatellia (Subgroup 4)
Blastocatellales
Blastocatellaceae
JGI:fa
NA



Skiv_18-Q3-R9-52_MAXAC.018
Bacteroidetes
Ignavibacteria
SJA-28
midas_f_31
midas_g_31
midas_s_3

