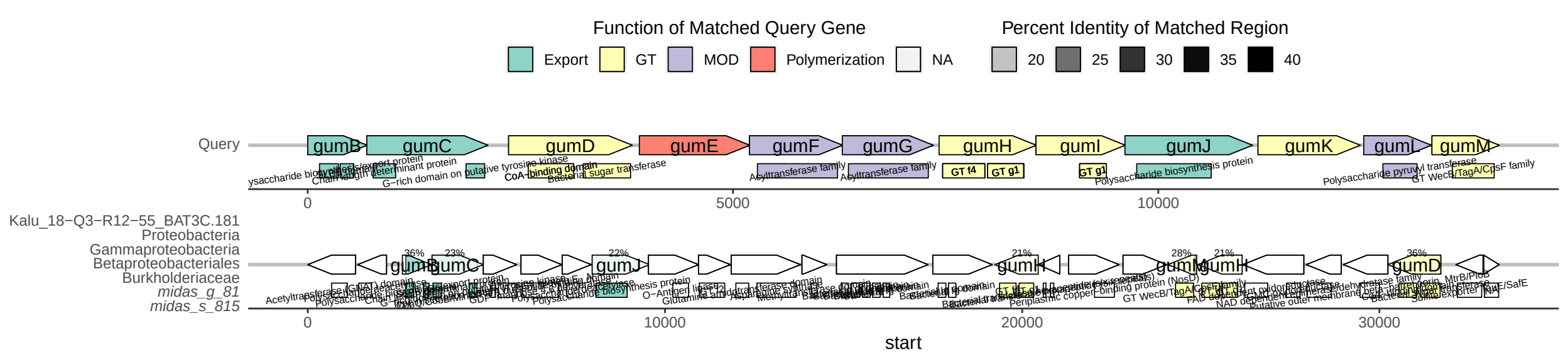
Function of Matched Query Gene
Export
GT
MOD
Polymerization
NA
Percent Identity of Matched Region
20
25
30
35
40
Query
gumB
gumC
gumD
gumE
gumF
gumG
gumH
gumI
gumJ
gumK
gumL
gumM
Acyltransferase family
Acyltransferase family
GT f4
GT g1
GT g1
Polysaccharide biosynthesis protein
Polysaccharide pyruvyl transferase
GT WecB/TagA/CpsF family
0
5000
10000
Kalu_18−Q3−R12−55_BAT3C.181
Proteobacteria
Gammaproteobacteria
Betaproteobacteriales
Burkholderiaceae
midas_g_81
midas_s_815
36%
23%
22%
21%
28%
21%
36%
gumB
gumC
gumJ
gumH
gumI
gumH
gumD
0
10000
20000
30000
start