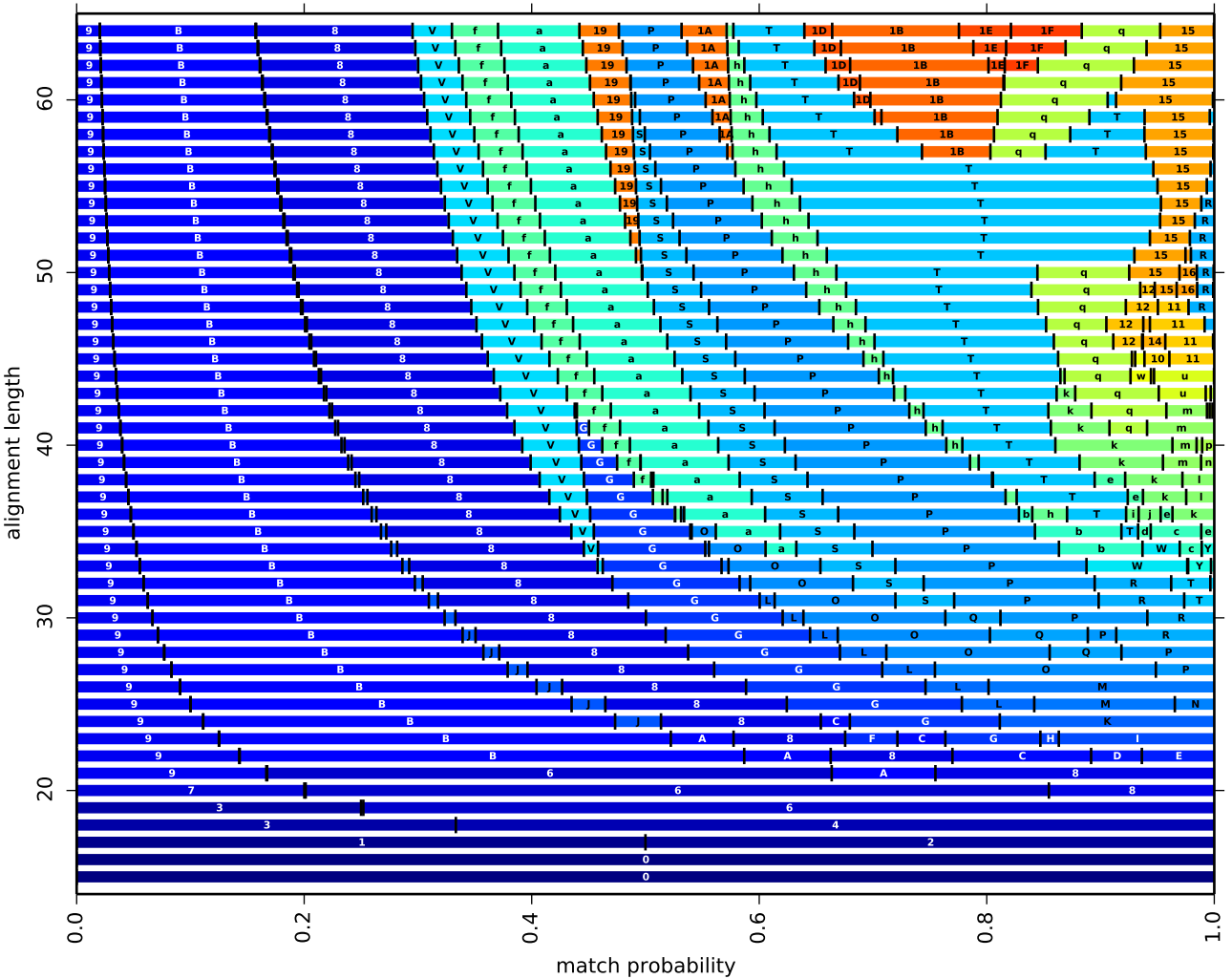
alignment length
60
50
40
30
20
0.0
0.2
0.4
0.6
0.8
1.0
match probability