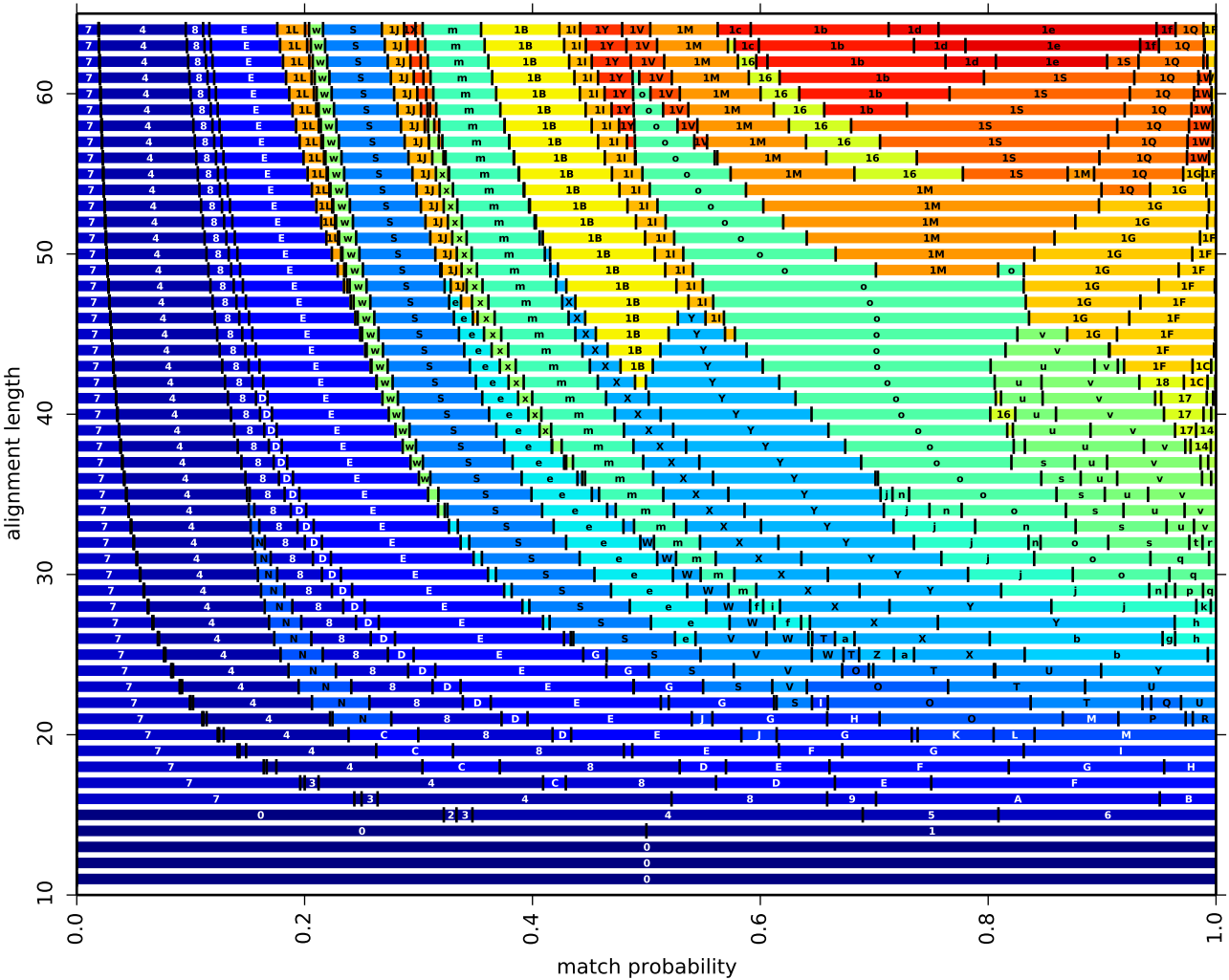
alignment length
match probability
0.0
0.2
0.4
0.6
0.8
1.0
10
20
30
40
50
60