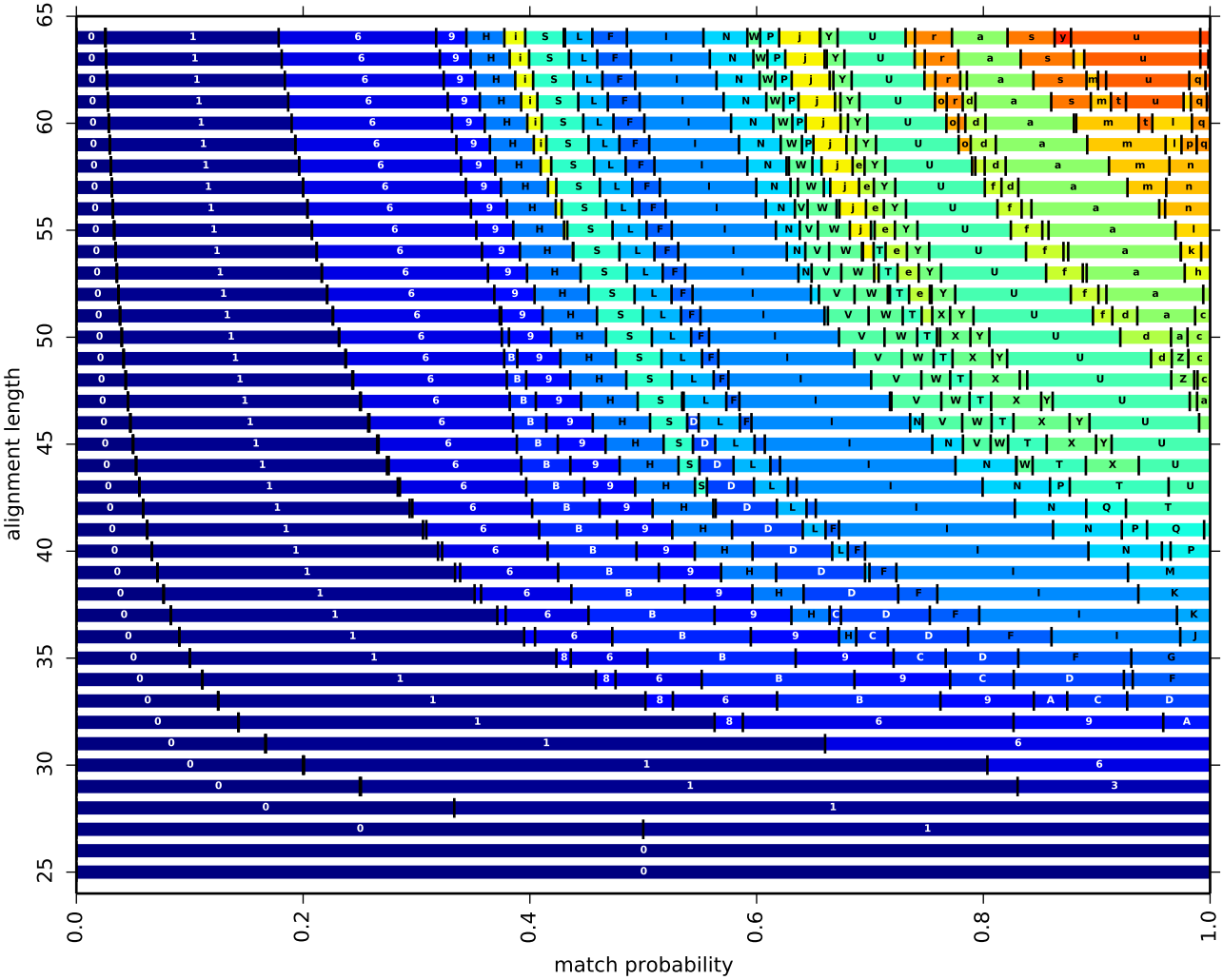
alignment length
match probability
65
60
55
50
45
40
35
30
25
0.0
0.2
0.4
0.6
0.8
1.0