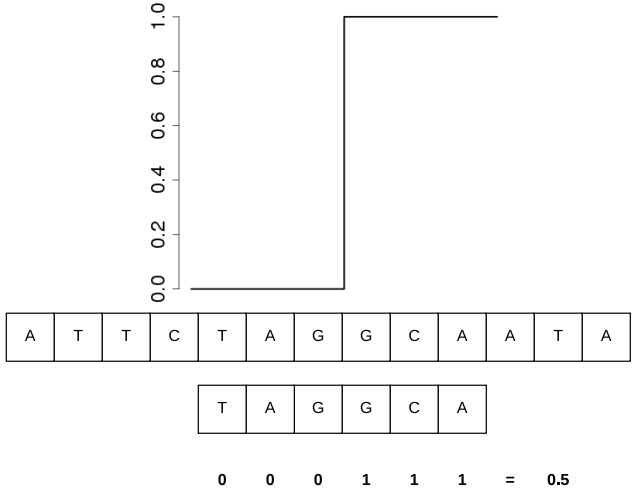
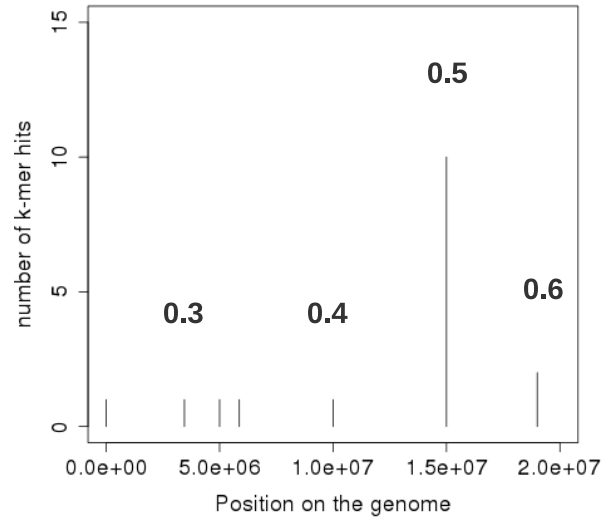
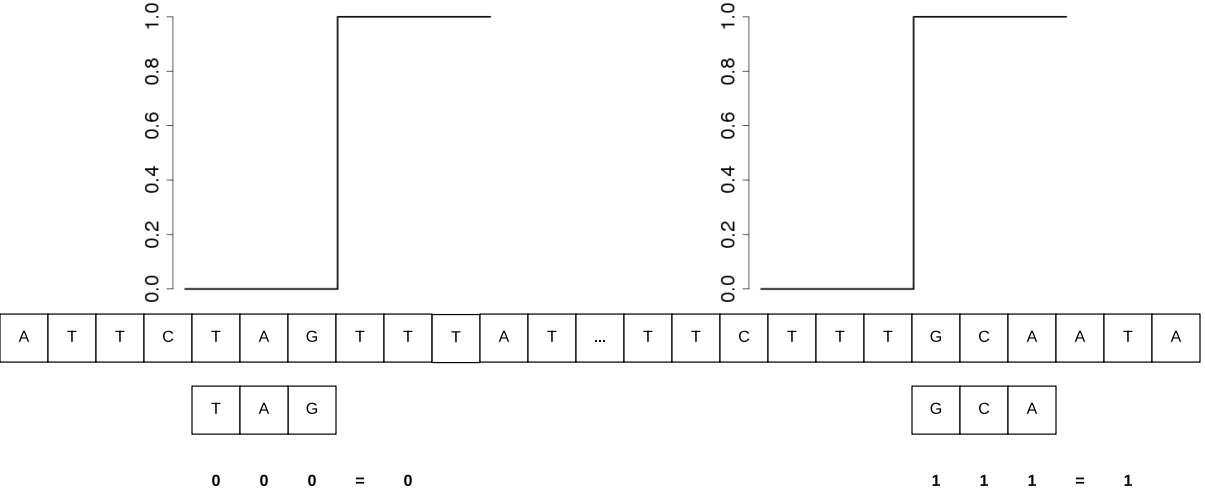

0.0
0.2
0.4
0.6
0.8
1.0
A T T C T A G G C A A T A
T A G G C A
0 0 0 1 1 1 = 0.5


number of k-mer hits
0
5
10
15
0.5
0.3
0.4
0.6
0.0e+00
5.0e+06
1.0e+07
1.5e+07
2.0e+07
Position on the genome


0.0
0.2
0.4
0.6
0.8
1.0
A T T C T A G T T T A T ... T T C T T T G C A A T A
T A G
0 0 0 = 0
G C A
1 1 1 = 1