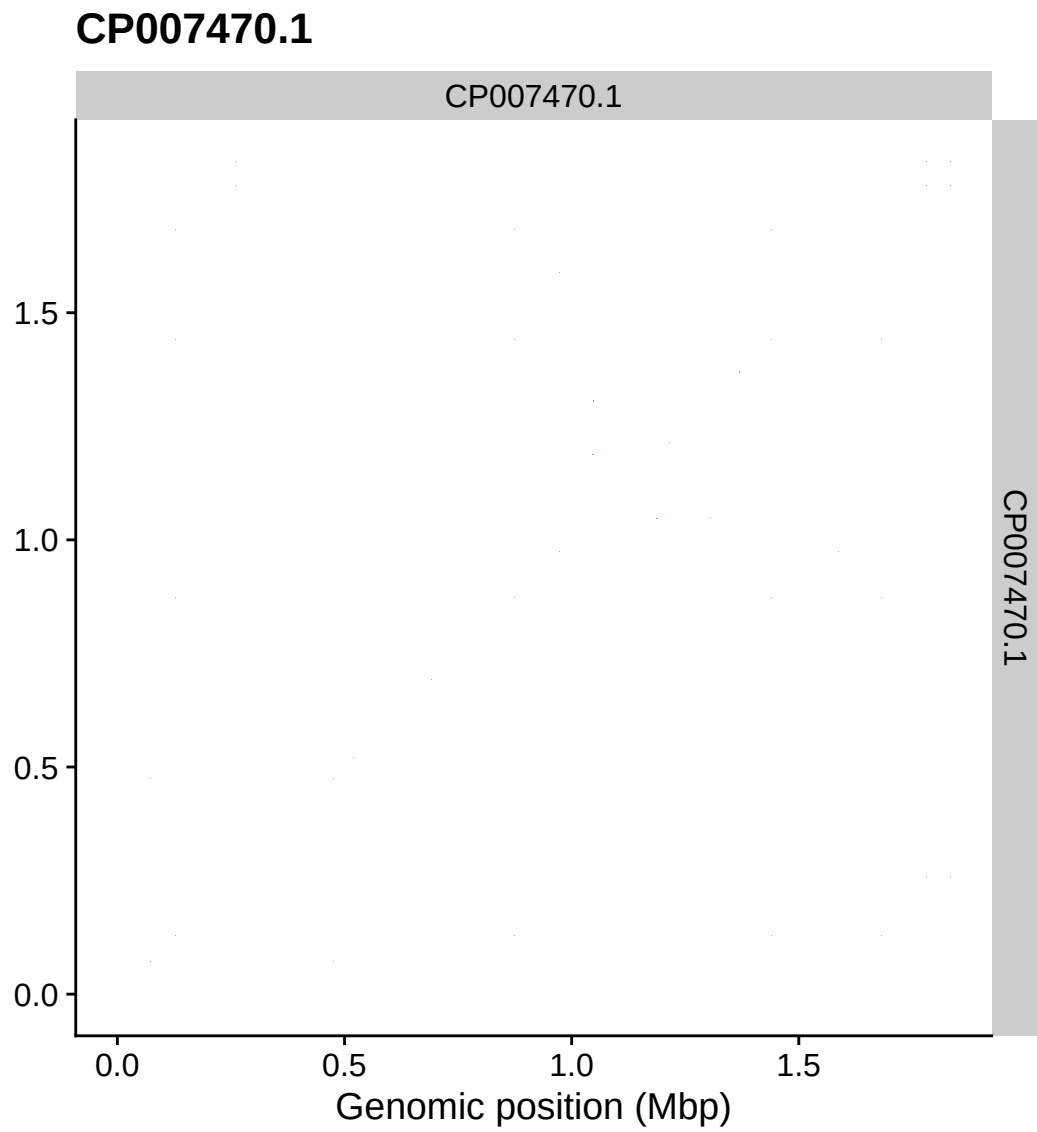
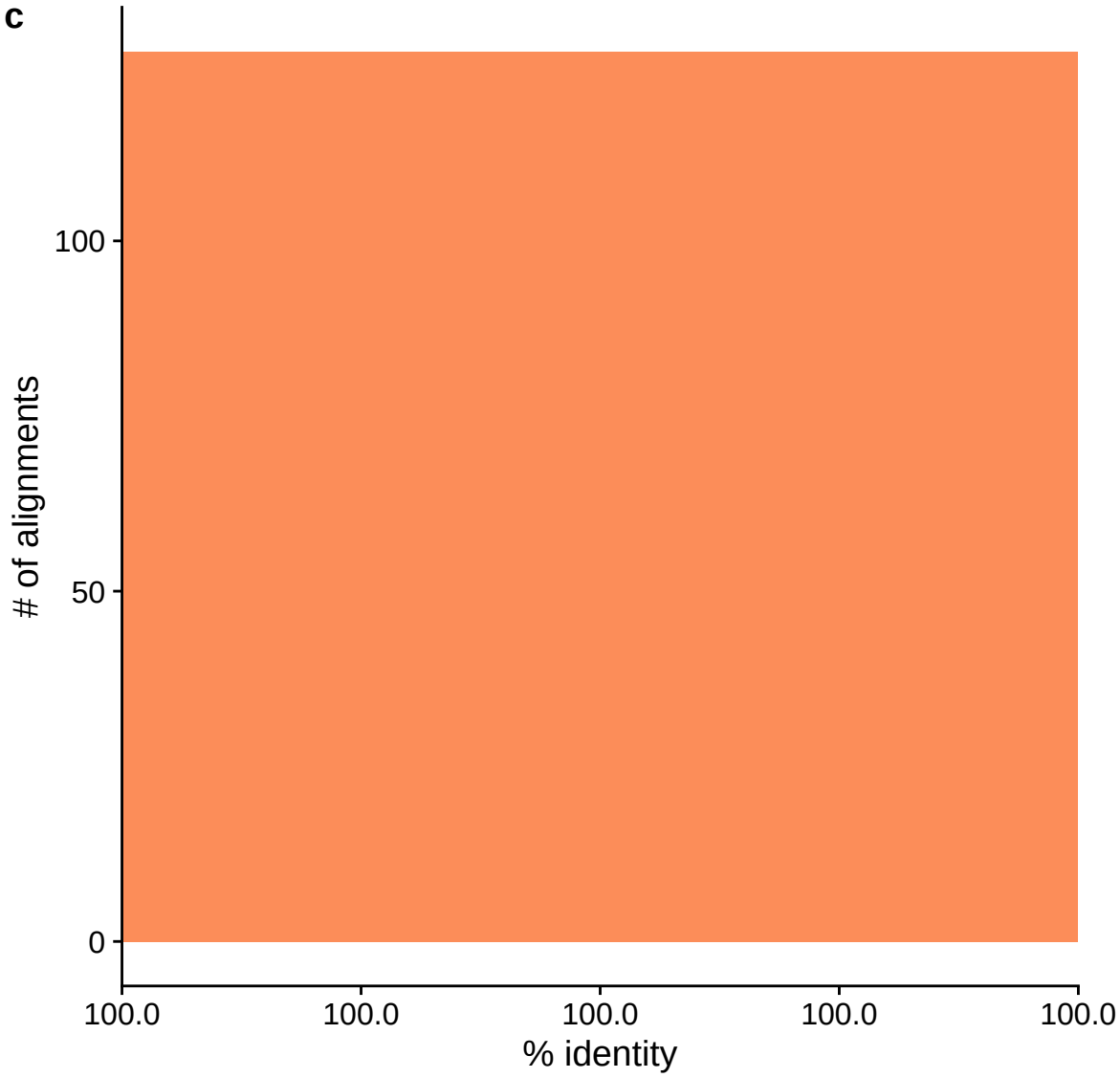
a


CP007470.1
CP007470.1
CP007470.1
1.5
1.0
0.5
0.0
0.0
0.5
1.0
1.5
Genomic position (Mbp)

b

c


100
50
0
# of alignments
100.0
100.0
100.0
100.0
100.0
% identity