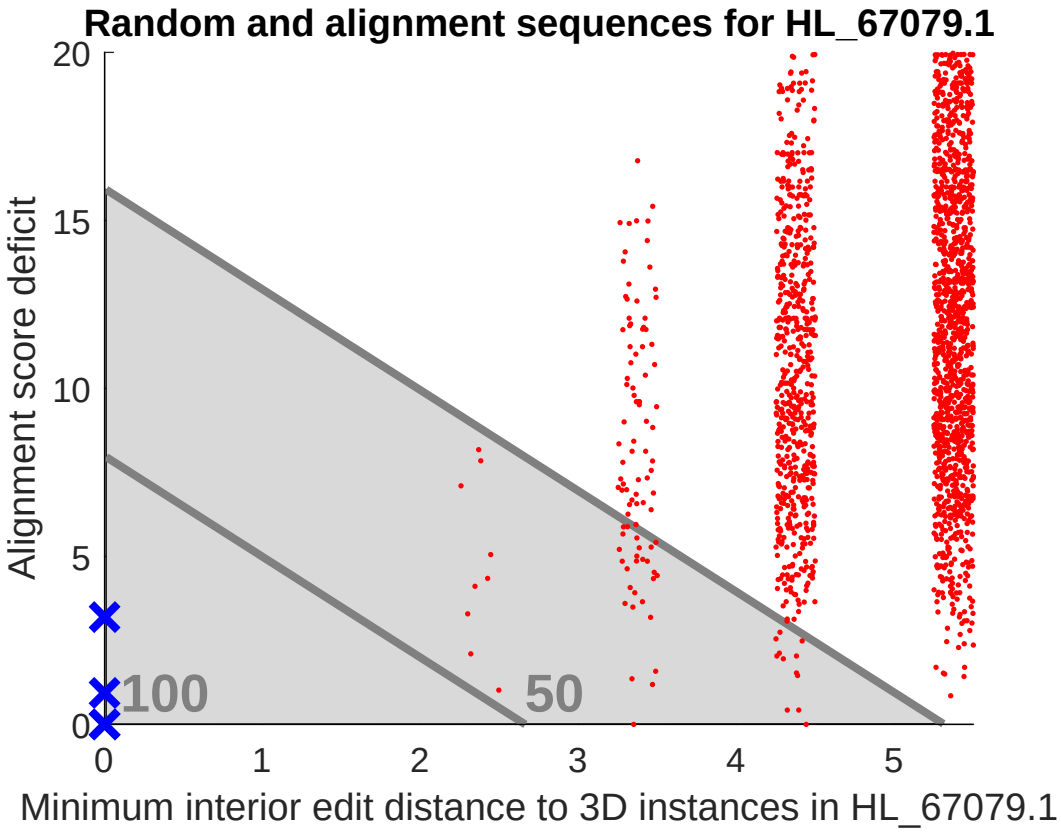
Random and alignment sequences for HL_67079.1
Alignment score deficit
Minimum interior edit distance to 3D instances in HL_67079.1
100
50
0
5
10
15
20
0
1
2
3
4
5