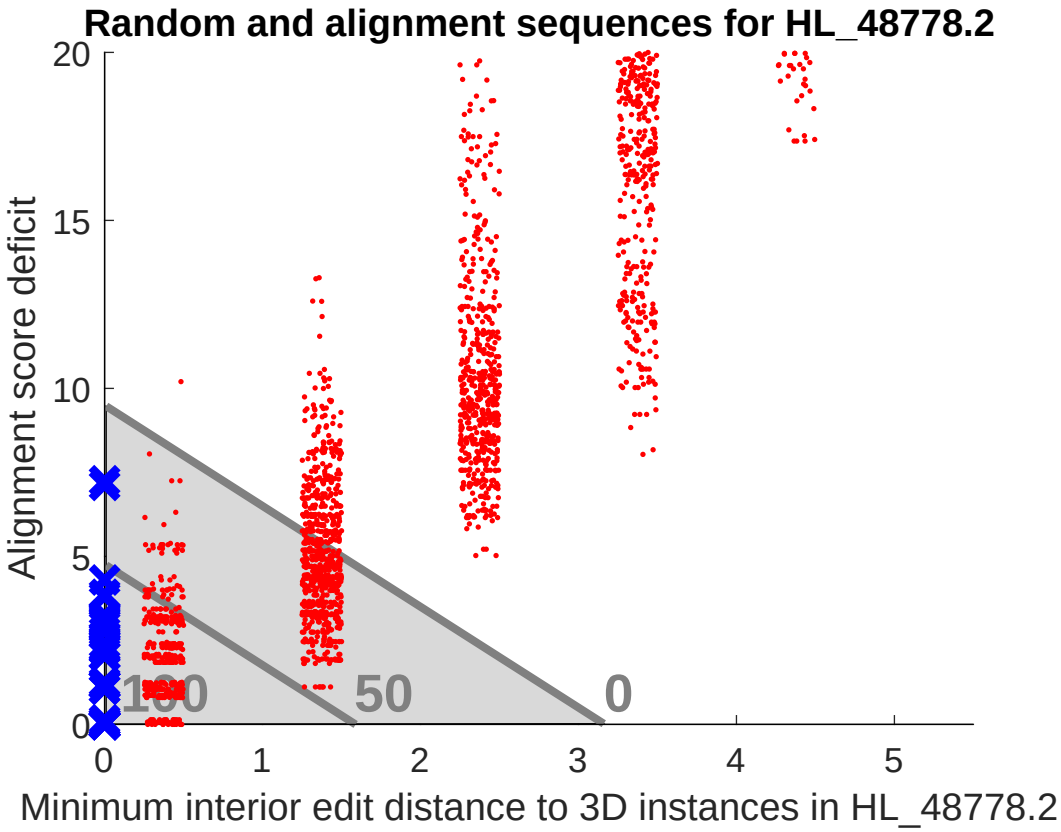
Random and alignment sequences for HL_48778.2
Alignment score deficit
Minimum interior edit distance to 3D instances in HL_48778.2
100
50
0
0
5
10
15
20
0
1
2
3
4
5