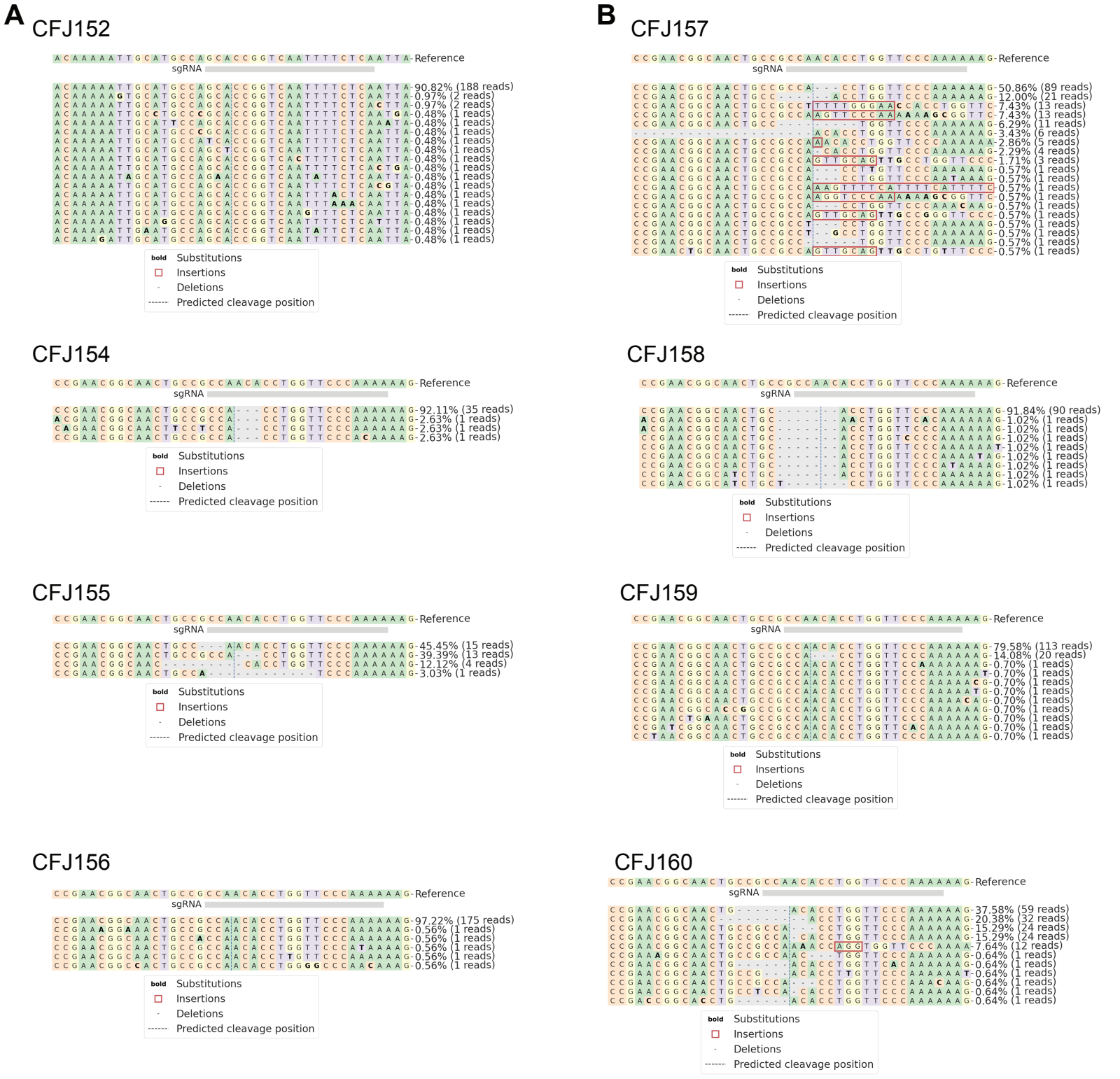
A
CFJ152
Reference
sgRNA
90.82% (188 reads)
0.97% (2 reads)
0.97% (2 reads)
0.48% (1 reads)
0.48% (1 reads)
0.48% (1 reads)
0.48% (1 reads)
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0.48% (1 reads)
0.48% (1 reads)
0.48% (1 reads)
0.48% (1 reads)
0.48% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
CFJ154
Reference
sgRNA
92.11% (35 reads)
2.63% (1 reads)
2.63% (1 reads)
2.63% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
CFJ155
Reference
sgRNA
45.45% (15 reads)
39.39% (13 reads)
12.12% (4 reads)
3.03% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
CFJ156
Reference
sgRNA
97.22% (175 reads)
0.56% (1 reads)
0.56% (1 reads)
0.56% (1 reads)
0.56% (1 reads)
0.56% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
B
CFJ157
Reference
sgRNA
50.86% (89 reads)
12.00% (21 reads)
7.43% (13 reads)
7.43% (13 reads)
6.29% (11 reads)
3.43% (6 reads)
2.86% (5 reads)
2.29% (4 reads)
1.71% (3 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
0.57% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
CFJ158
Reference
sgRNA
91.84% (90 reads)
1.02% (1 reads)
1.02% (1 reads)
1.02% (1 reads)
1.02% (1 reads)
1.02% (1 reads)
1.02% (1 reads)
1.02% (1 reads)
1.02% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
CFJ159
Reference
sgRNA
79.58% (113 reads)
14.08% (20 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
0.70% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position
CFJ160
Reference
sgRNA
37.58% (59 reads)
20.38% (32 reads)
15.29% (24 reads)
15.29% (24 reads)
7.64% (12 reads)
0.64% (1 reads)
0.64% (1 reads)
0.64% (1 reads)
0.64% (1 reads)
0.64% (1 reads)
0.64% (1 reads)
bold Substitutions
Insertions
Deletions
Predicted cleavage position