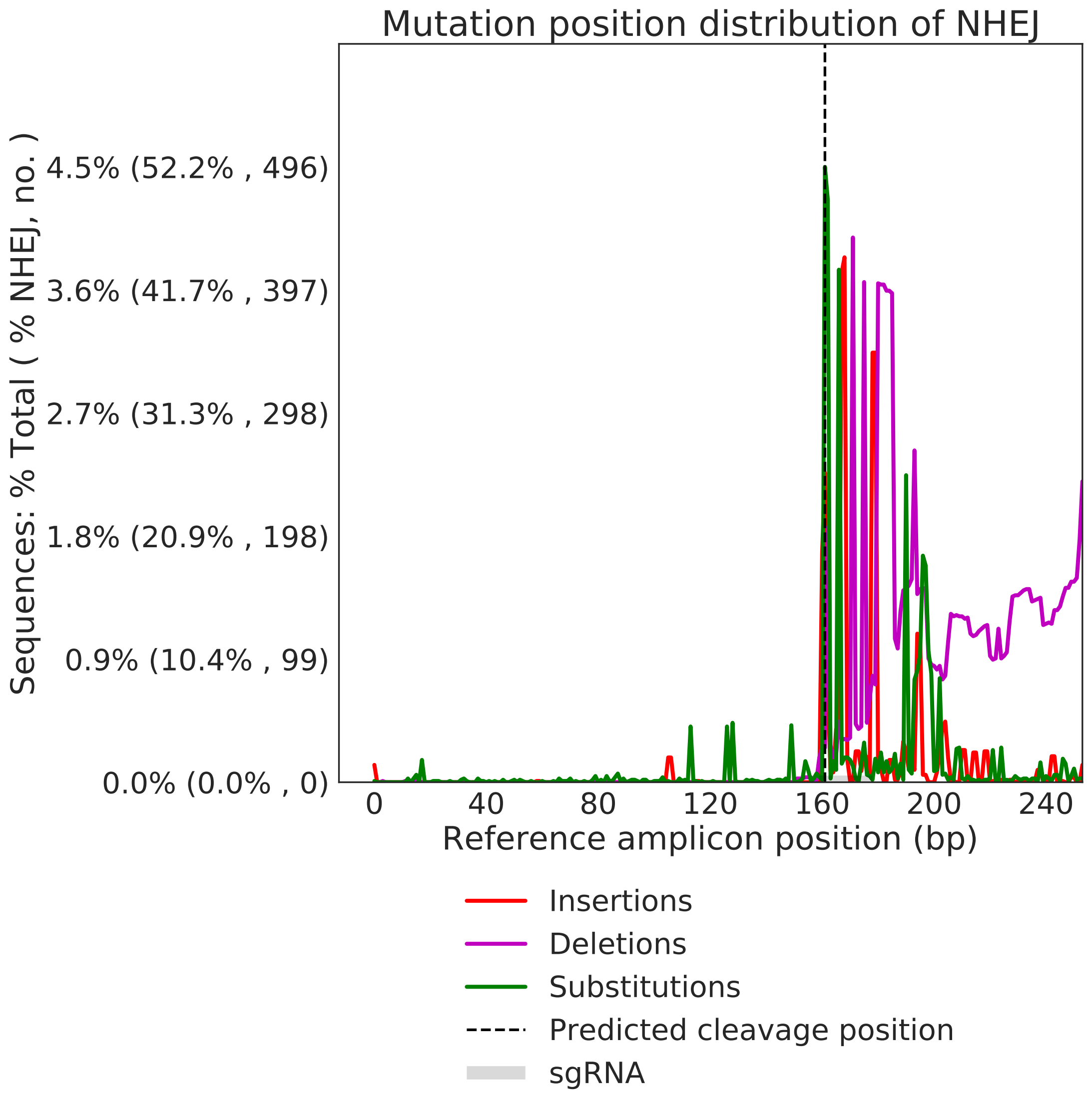
Mutation position distribution of NHEJ
Sequences: % Total (% NHEJ, no.)
4.5% (52.2% , 496)
3.6% (41.7% , 397)
2.7% (31.3% , 298)
1.8% (20.9% , 198)
0.9% (10.4% , 99)
0.0% (0.0% , 0)
0
40
80
120
160
200
240
Reference amplicon position (bp)
Insertions
Deletions
Substitutions
Predicted cleavage position
sgRNA