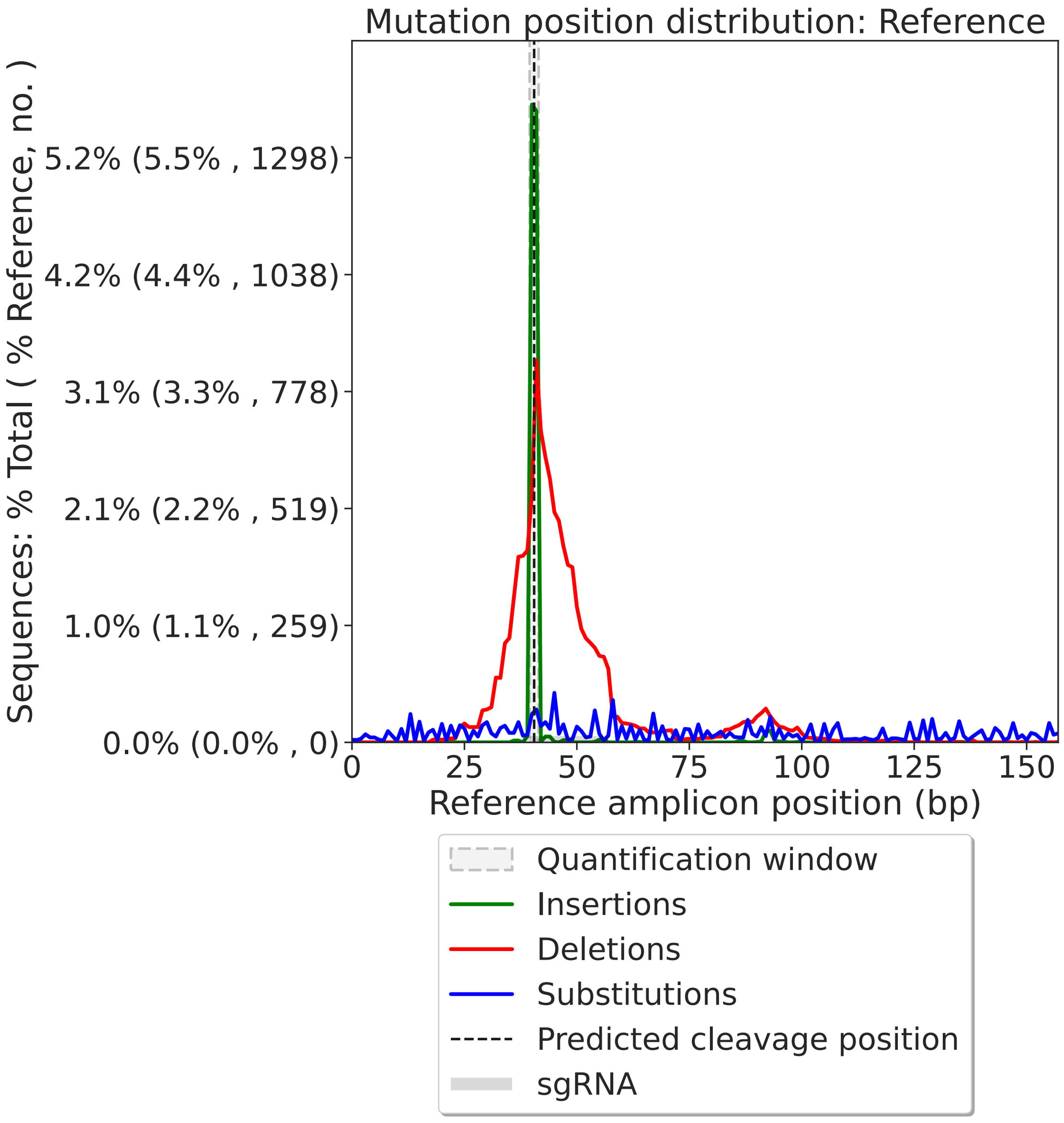
Mutation position distribution: Reference
Sequences: % Total (% Reference, no.)
5.2% (5.5% , 1298)
4.2% (4.4% , 1038)
3.1% (3.3% , 778)
2.1% (2.2% , 519)
1.0% (1.1% , 259)
0.0% (0.0% , 0)
0
25
50
75
100
125
150
Reference amplicon position (bp)
Quantification window
Insertions
Deletions
Substitutions
Predicted cleavage position
sgRNA