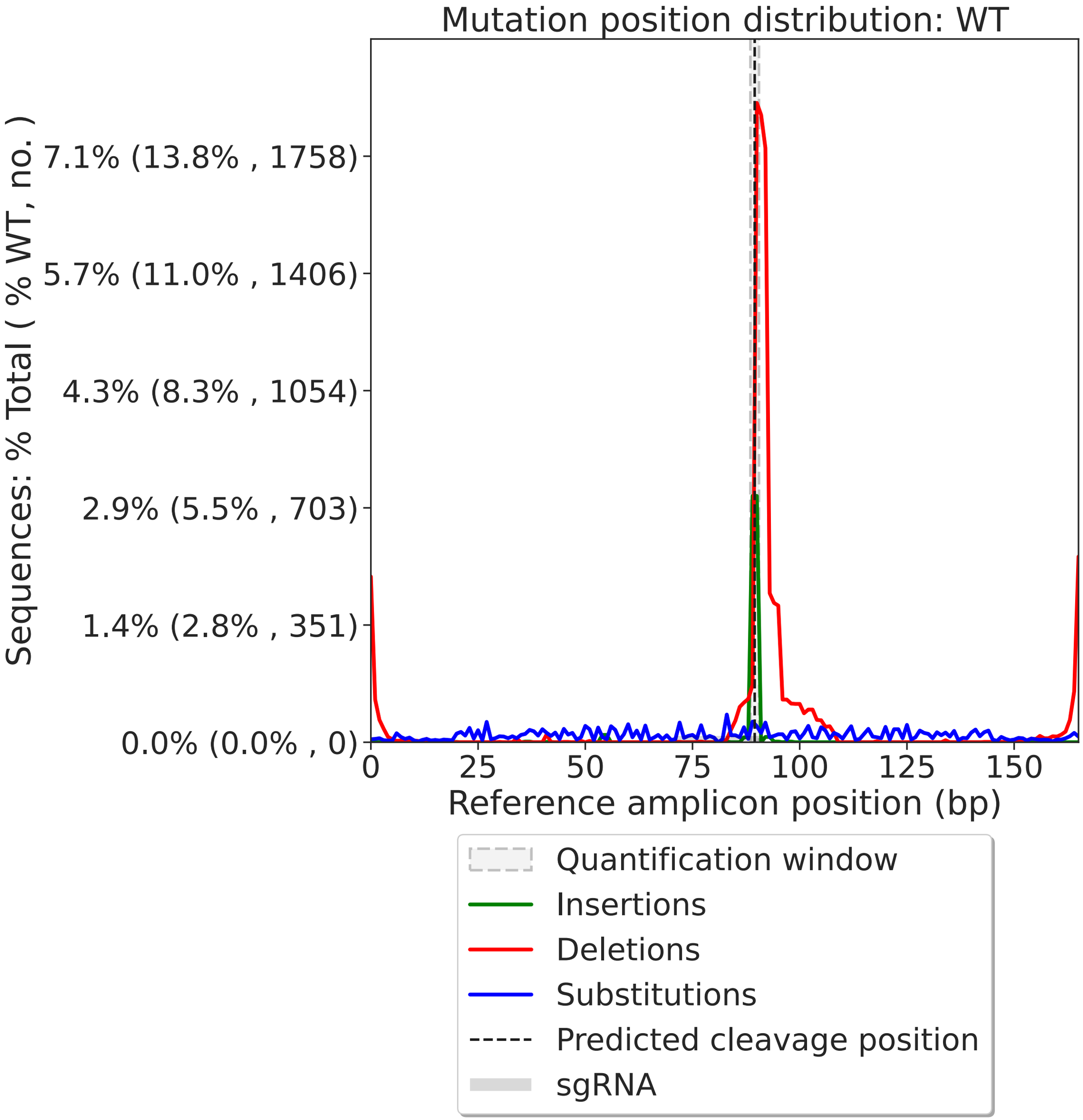
Mutation position distribution: WT
Sequences: % Total ( % WT, no. )
7.1% (13.8% , 1758)
5.7% (11.0% , 1406)
4.3% (8.3% , 1054)
2.9% (5.5% , 703)
1.4% (2.8% , 351)
0.0% (0.0% , 0)
0
25
50
75
100
125
150
Reference amplicon position (bp)
Quantification window
Insertions
Deletions
Substitutions
Predicted cleavage position
sgRNA