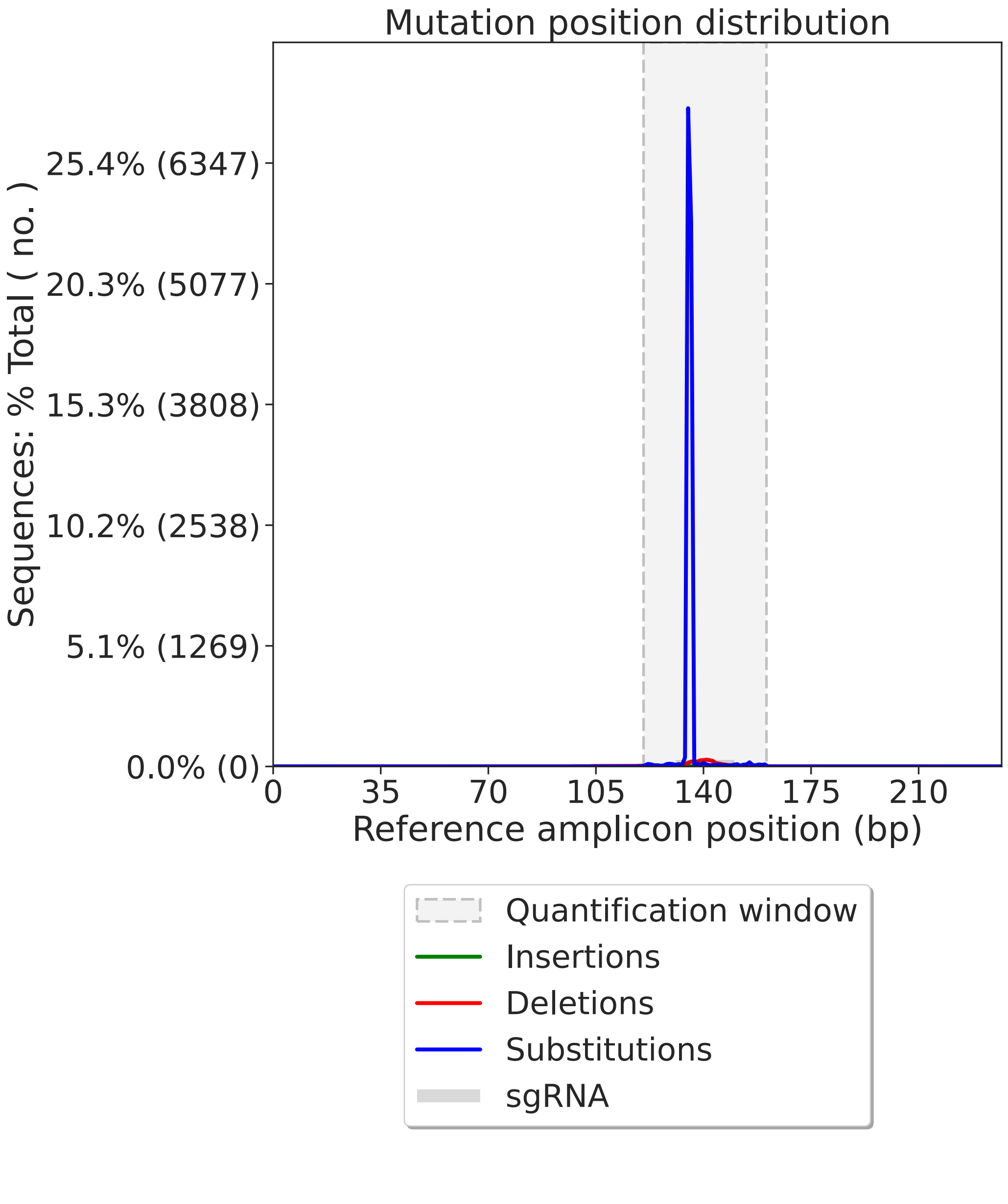
Mutation position distribution
Sequences: % Total (no.)
25.4% (6347)
20.3% (5077)
15.3% (3808)
10.2% (2538)
5.1% (1269)
0.0% (0)
0
35
70
105
140
175
210
Reference amplicon position (bp)
Quantification window
Insertions
Deletions
Substitutions
sgRNA