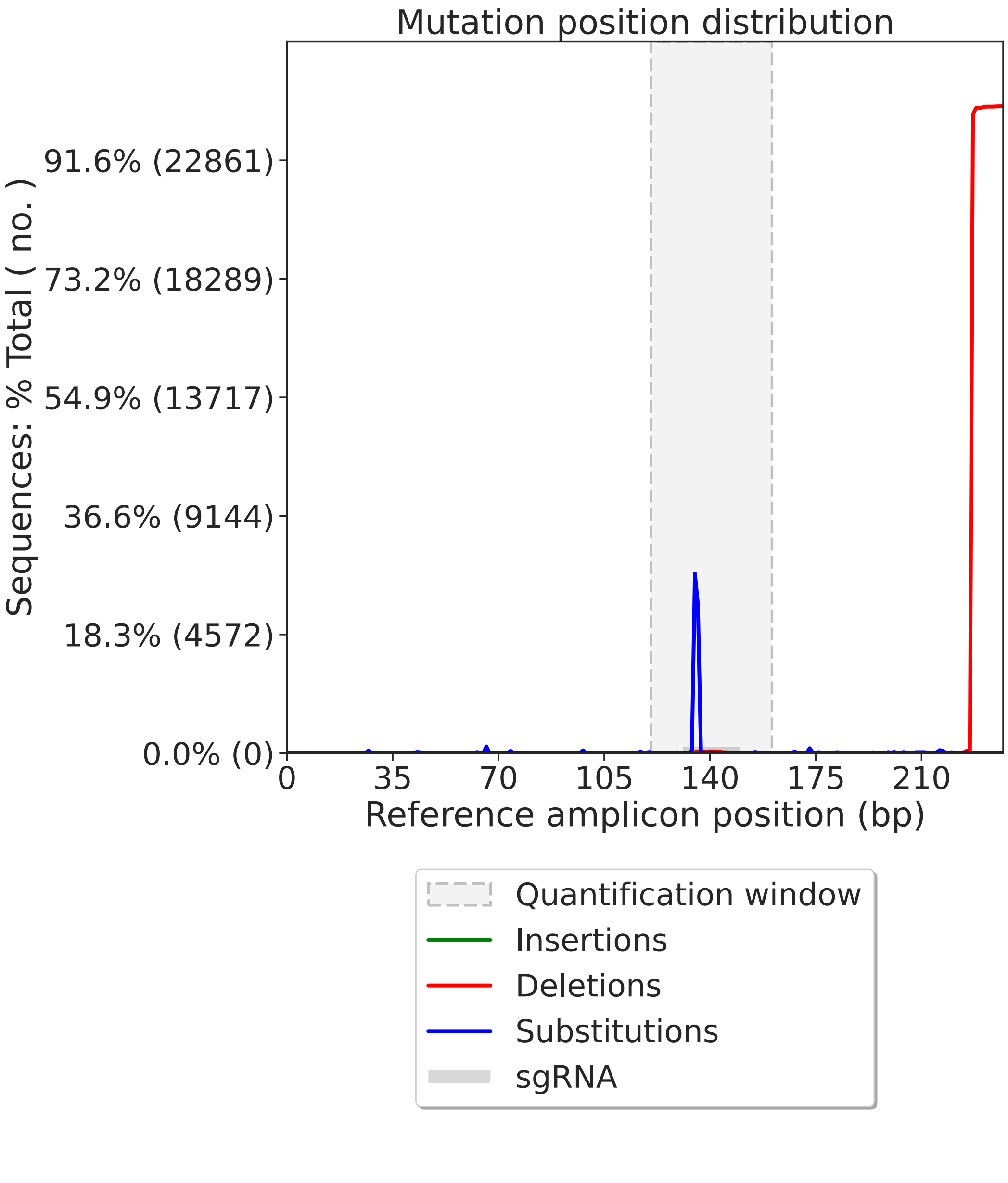
Mutation position distribution
Sequences: % Total (no.)
91.6% (22861)
73.2% (18289)
54.9% (13717)
36.6% (9144)
18.3% (4572)
0.0% (0)
0
35
70
105
140
175
210
Reference amplicon position (bp)
Quantification window
Insertions
Deletions
Substitutions
sgRNA