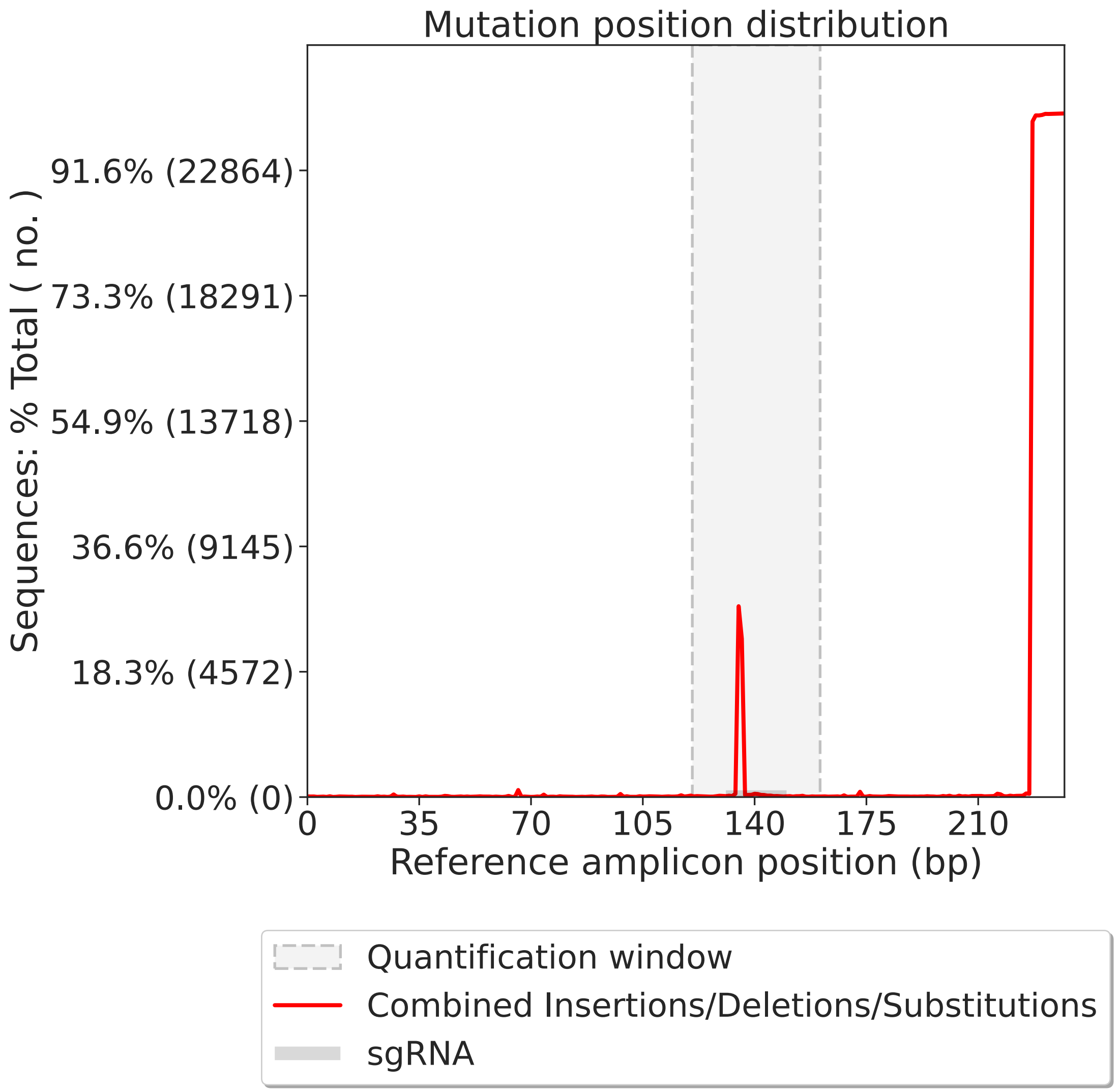
Mutation position distribution
Sequences: % Total (no.)
91.6% (22864)
73.3% (18291)
54.9% (13718)
36.6% (9145)
18.3% (4572)
0.0% (0)
0
35
70
105
140
175
210
Reference amplicon position (bp)
Quantification window
Combined Insertions/Deletions/Substitutions
sgRNA