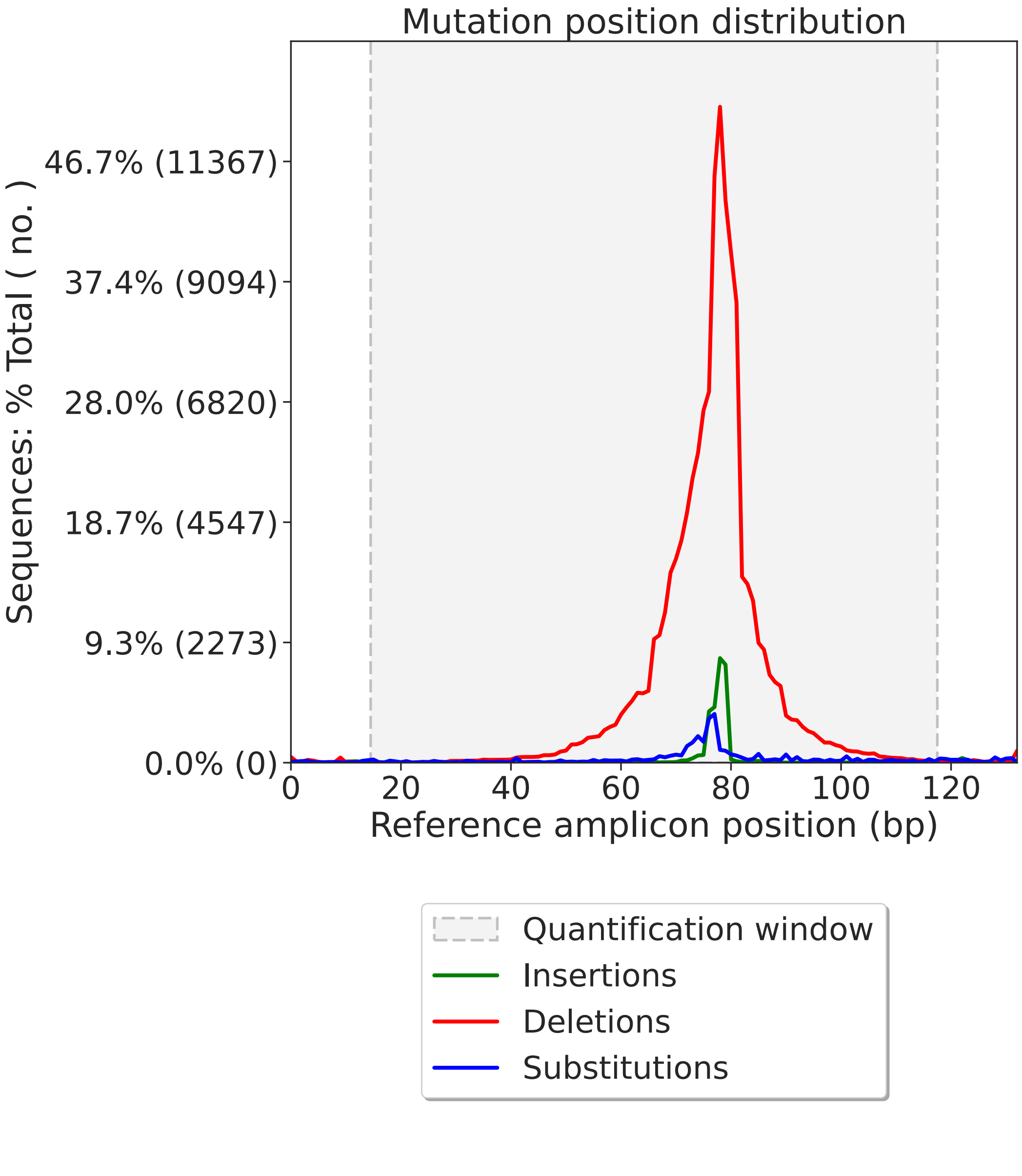
Mutation position distribution
Sequences: % Total (no.)
46.7% (11367)
37.4% (9094)
28.0% (6820)
18.7% (4547)
9.3% (2273)
0.0% (0)
0
20
40
60
80
100
120
Reference amplicon position (bp)
Quantification window
Insertions
Deletions
Substitutions