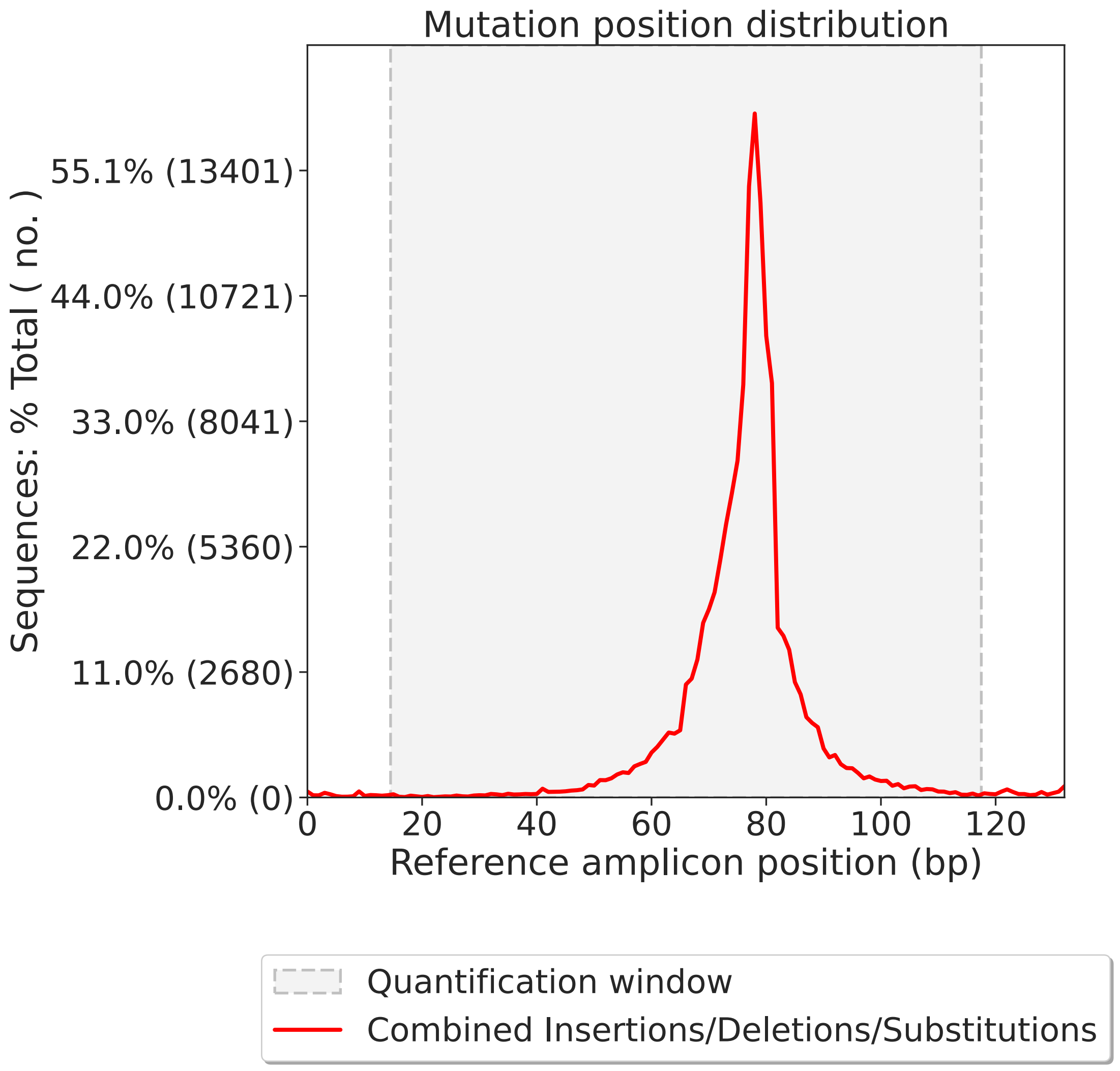
Mutation position distribution
Sequences: % Total (no.)
55.1% (13401)
44.0% (10721)
33.0% (8041)
22.0% (5360)
11.0% (2680)
0.0% (0)
0
20
40
60
80
100
120
Reference amplicon position (bp)
Quantification window
Combined Insertions/Deletions/Substitutions