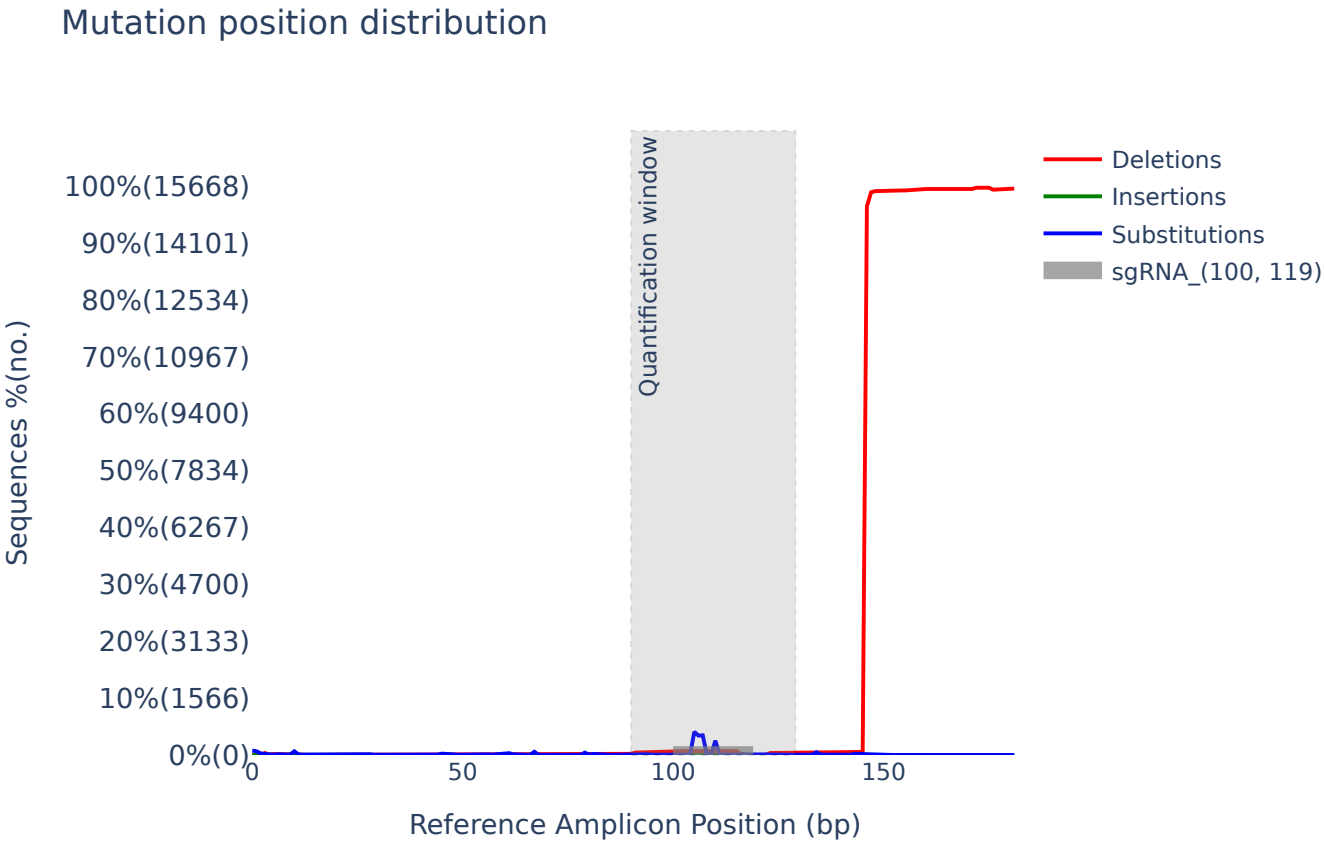
Mutation position distribution
100%(15668)
90%(14101)
80%(12534)
70%(10967)
60%(9400)
50%(7834)
40%(6267)
30%(4700)
20%(3133)
10%(1566)
0%(0)
Sequences %(no.)
Quantification window
Deletions
Insertions
Substitutions
sgRNA_(100, 119)
0
50
100
150
Reference Amplicon Position (bp)