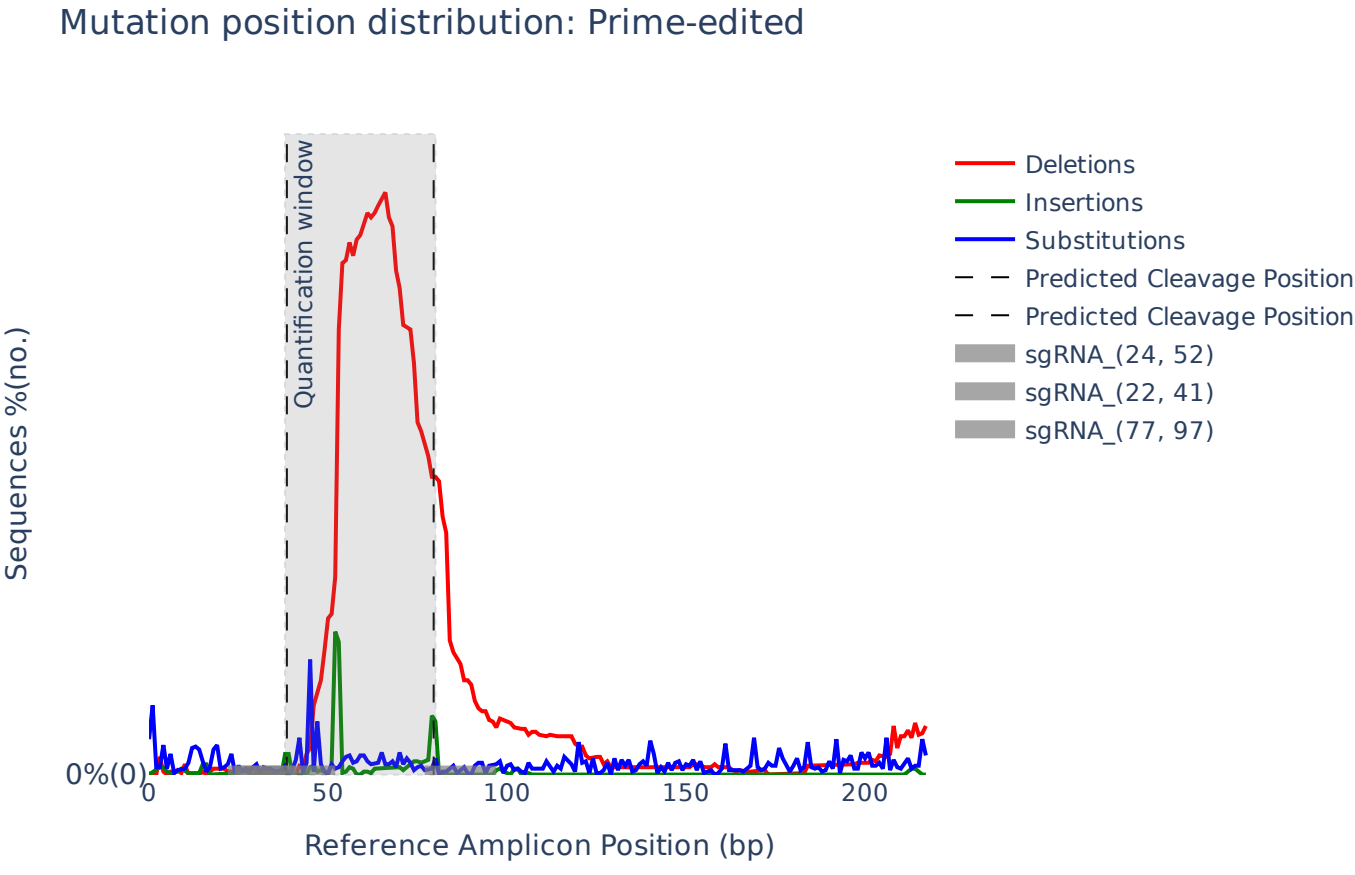
Mutation position distribution: Prime-edited
Quantification window
Deletions
Insertions
Substitutions
Predicted Cleavage Position
Predicted Cleavage Position
sgRNA_(24, 52)
sgRNA_(22, 41)
sgRNA_(77, 97)
Sequences %(no.)
0%(0)
0
50
100
150
200
Reference Amplicon Position (bp)