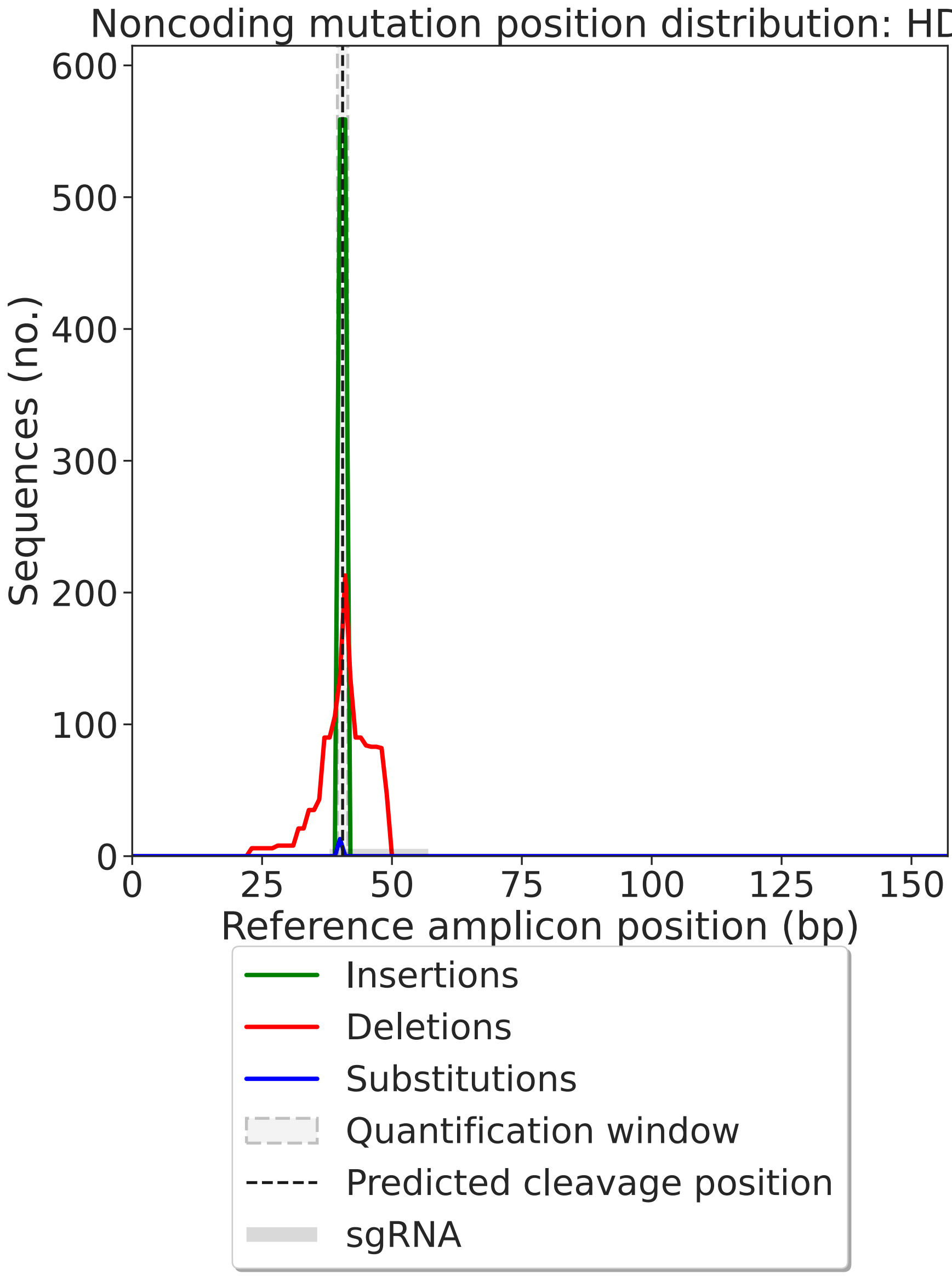
Noncoding mutation position distribution: HDR
Sequences (no.)
600
500
400
300
200
100
0
0
25
50
75
100
125
150
Reference amplicon position (bp)
Insertions
Deletions
Substitutions
Quantification window
Predicted cleavage position
sgRNA