

Mutation position distribution: WT

Sequences: % Total ( % WT, no. )

10.1% (19.4% , 2479)

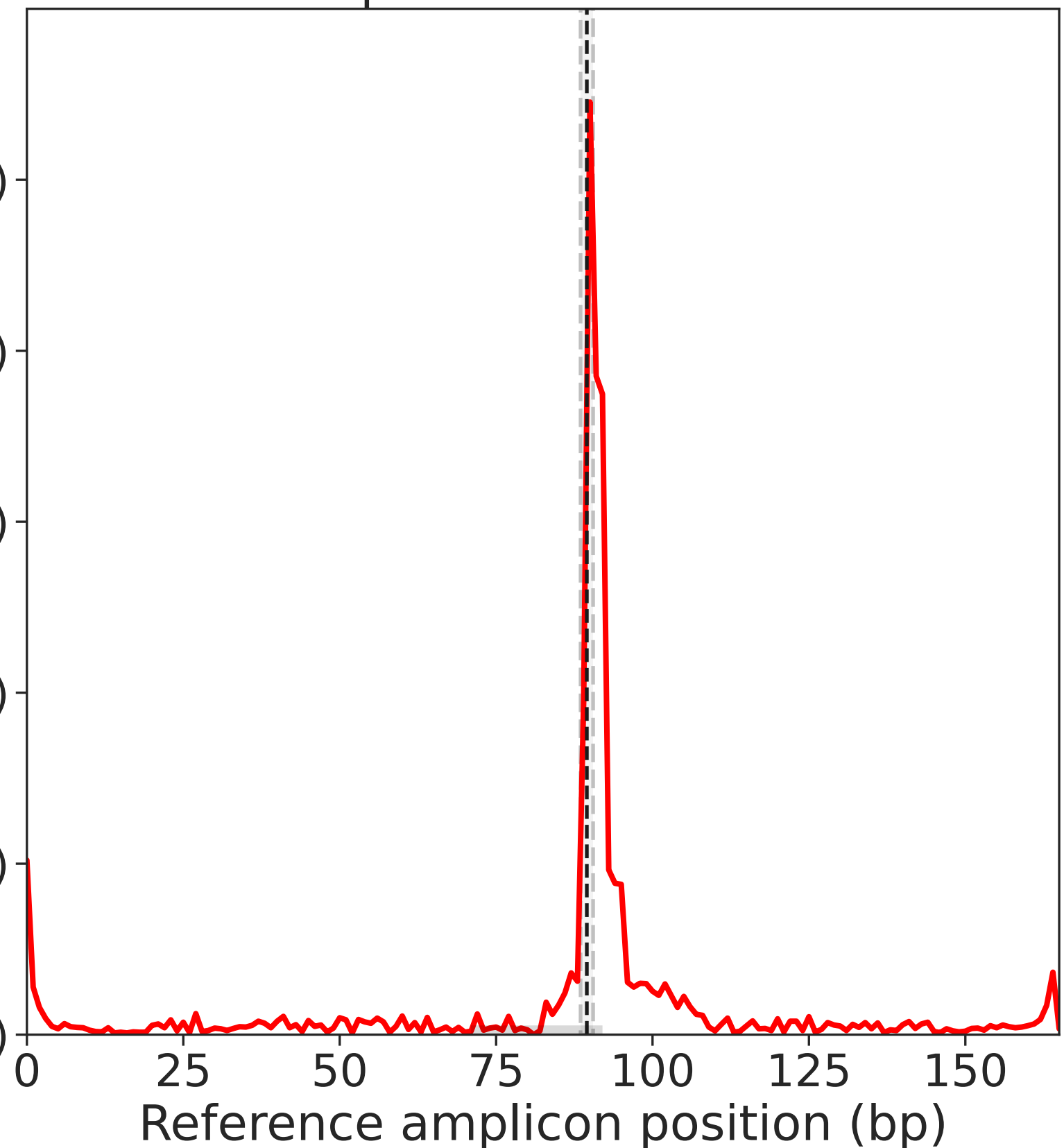
8.1% (15.6% , 1983)

6.0% (11.7% , 1487)

4.0% (7.8% , 991)

2.0% (3.9% , 495)

0.0% (0.0% , 0)



Quantification window



Combined Insertions/Deletions/Substitutions: WT



Predicted cleavage position



sgRNA