

Mutation position distribution: CASP3

Sequences: % Total (% CASP3, no.)

11.7% (93.9% , 10127)

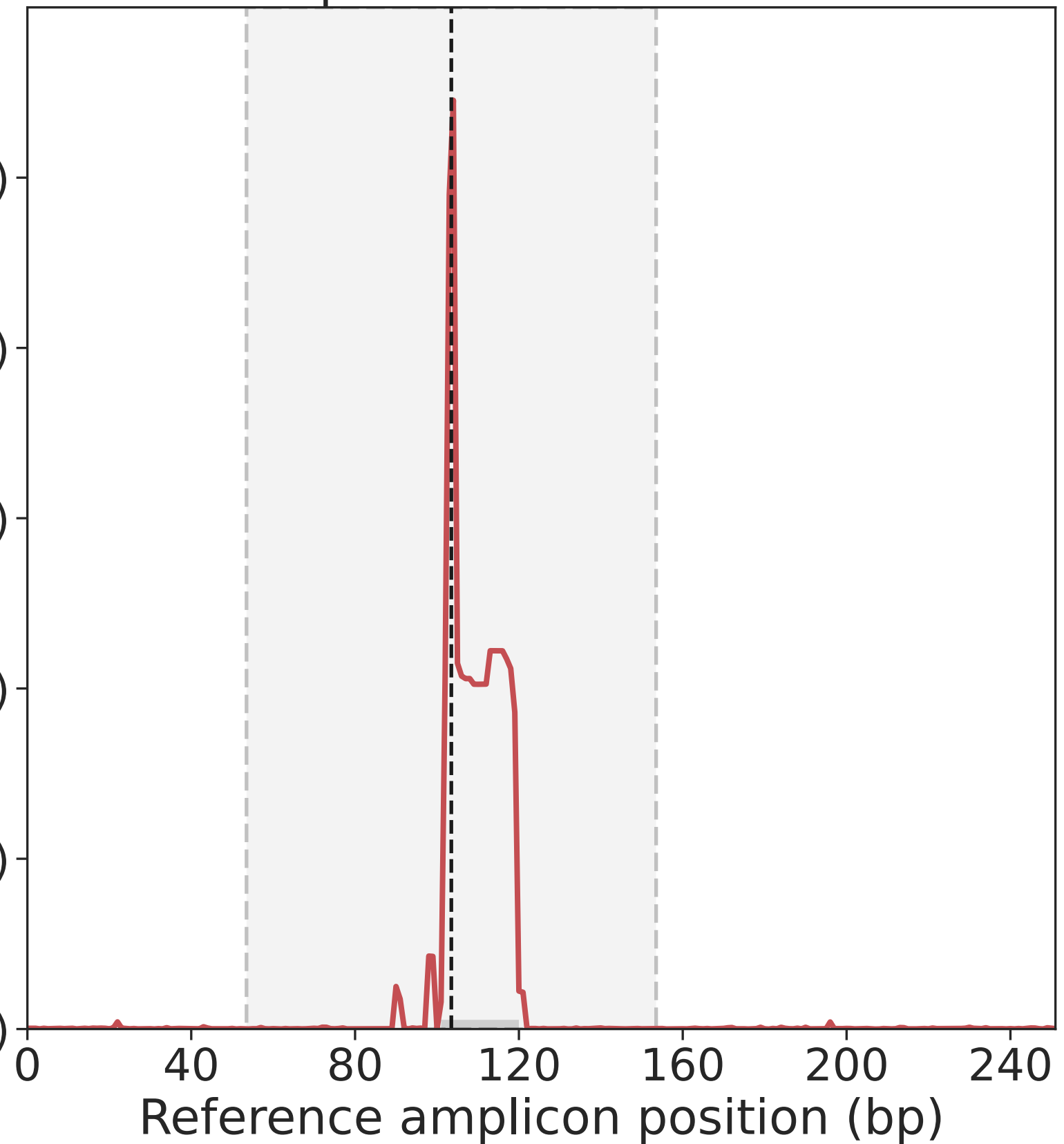
9.3% (75.1% , 8101)

7.0% (56.4% , 6076)

4.7% (37.6% , 4050)

2.3% (18.8% , 2025)

0.0% (0.0% , 0)



Quantification window



Combined Insertions/Deletions/Substitutions: CASP3



Predicted cleavage position



sgRNA