

Mutation position distribution: CASP2

Sequences: % Total (% CASP2, no.)

26.3% (85.3% , 4642)

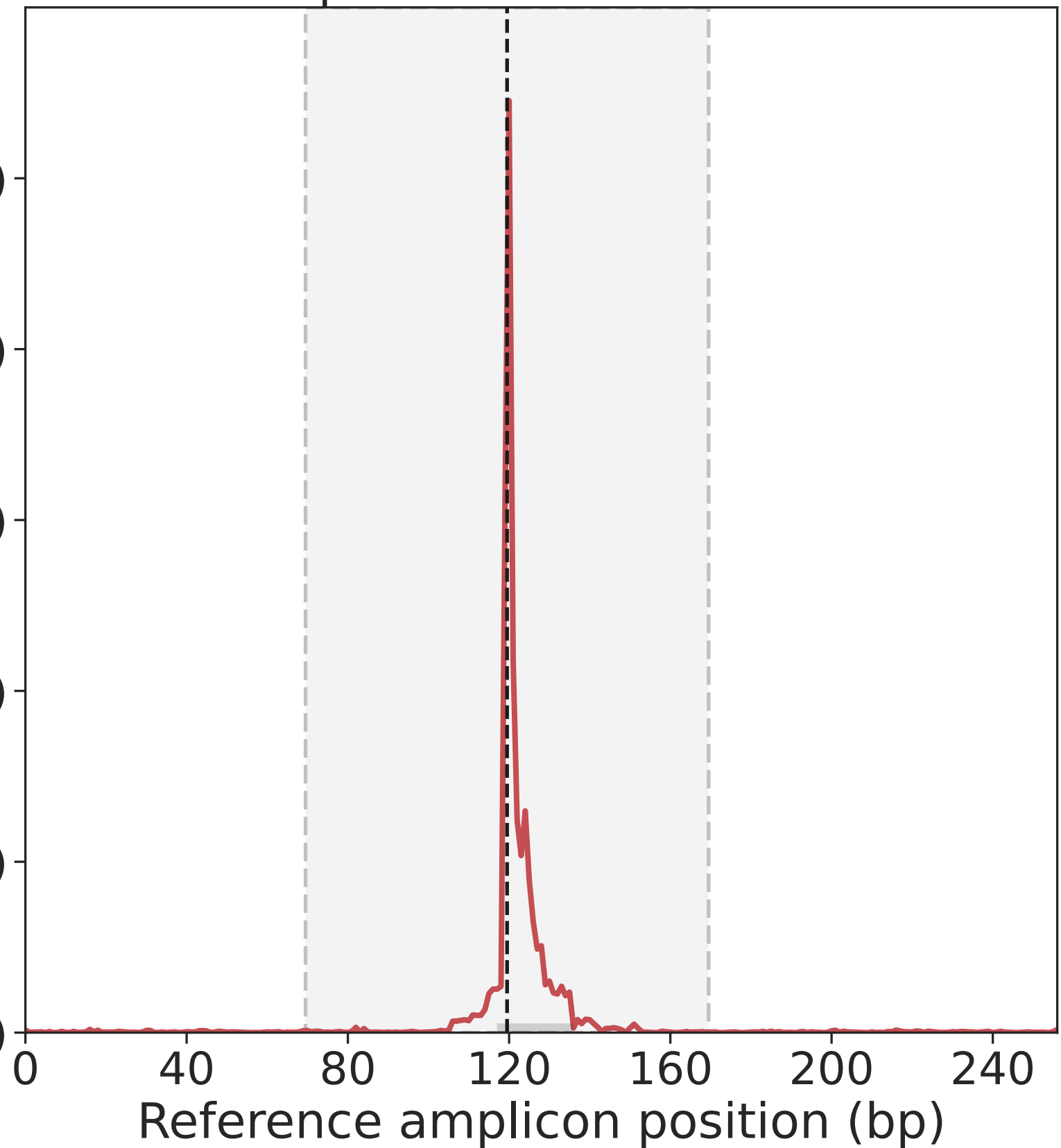
21.1% (68.3% , 3714)

15.8% (51.2% , 2785)

10.5% (34.1% , 1857)

5.3% (17.1% , 928)

0.0% (0.0% , 0)



Quantification window



Combined Insertions/Deletions/Substitutions: CASP2



Predicted cleavage position



sgRNA