

Mutation position distribution: EIF3I

Sequences: % Total (% EIF3I, no.)

3.0% (39.9% , 2135)

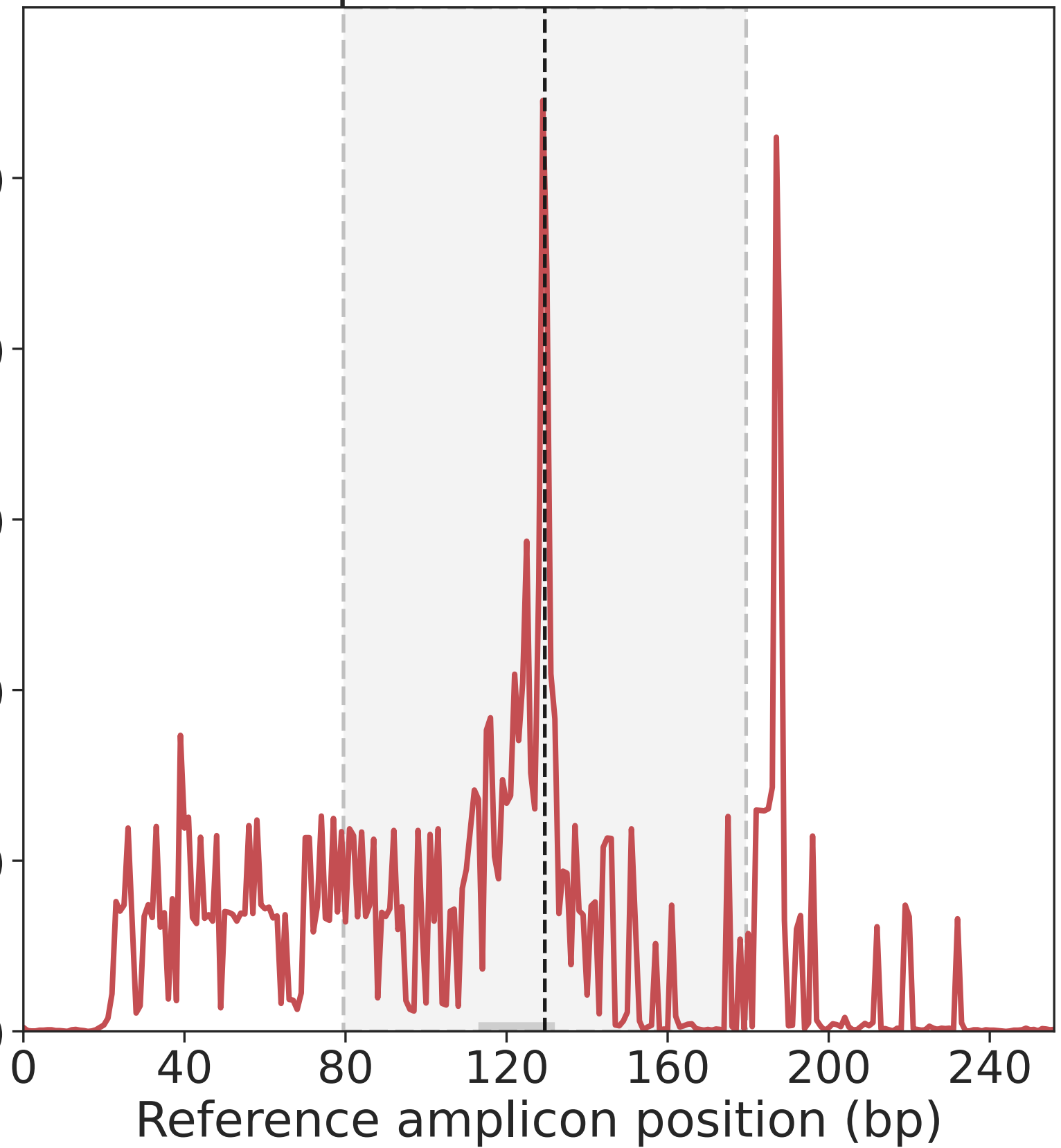
2.4% (31.9% , 1708)

1.8% (24.0% , 1281)

1.2% (16.0% , 854)

0.6% (8.0% , 427)

0.0% (0.0% , 0)



Quantification window



Combined Insertions/Deletions/Substitutions: EIF3I



Predicted cleavage position



sgRNA