

Mutation position distribution: AKT1S1

Sequences: % Total ( % AKT1S1, no. )

12.1% (30.5% , 4365)

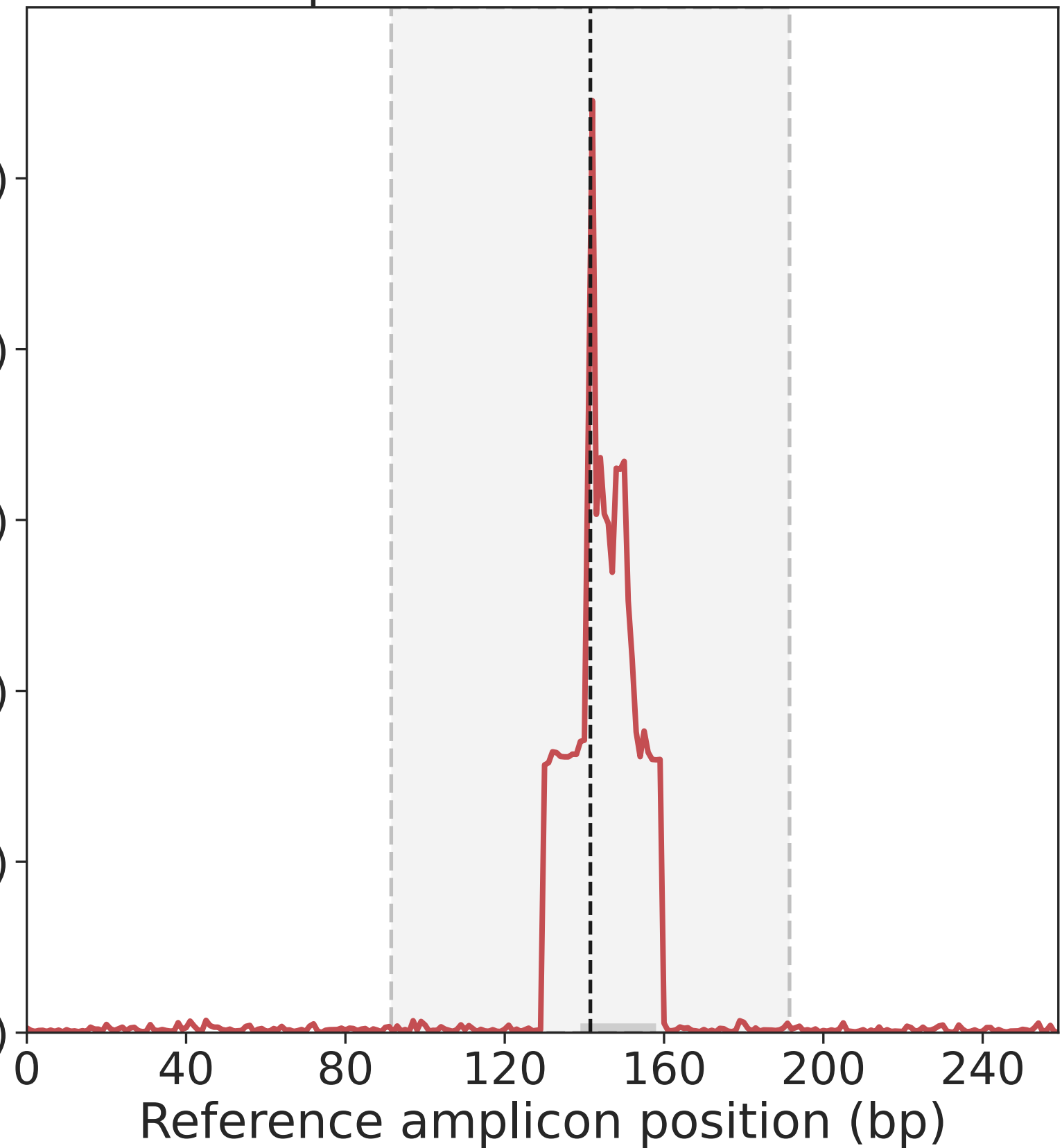
9.6% (24.4% , 3492)

7.2% (18.3% , 2619)

4.8% (12.2% , 1746)

2.4% (6.1% , 873)

0.0% (0.0% , 0)



- Quantification window
- Combined Insertions/Deletions/Substitutions: AKT1S1
- Predicted cleavage position
- sgRNA