

Mutation position distribution: P23H

Sequences: % Total (% P23H, no.)

20.6% (44.8% , 5069)

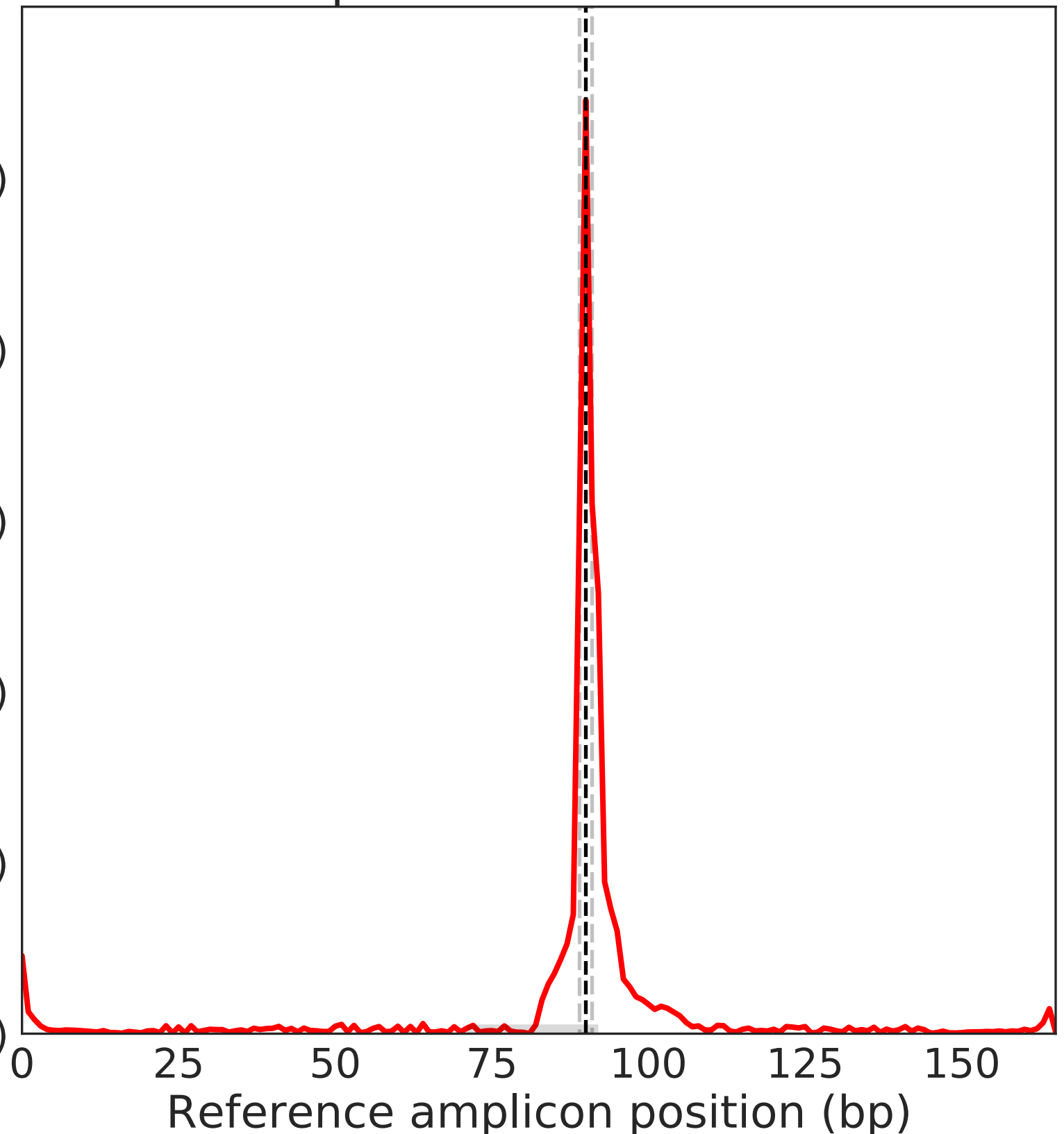
16.5% (35.9% , 4055)

12.4% (26.9% , 3041)

8.2% (17.9% , 2027)

4.1% (9.0% , 1013)

0.0% (0.0% , 0)



- Combined Insertions/Deletions/Substitutions: P23H
- - - Predicted cleavage position
- sgRNA
- Quantification window