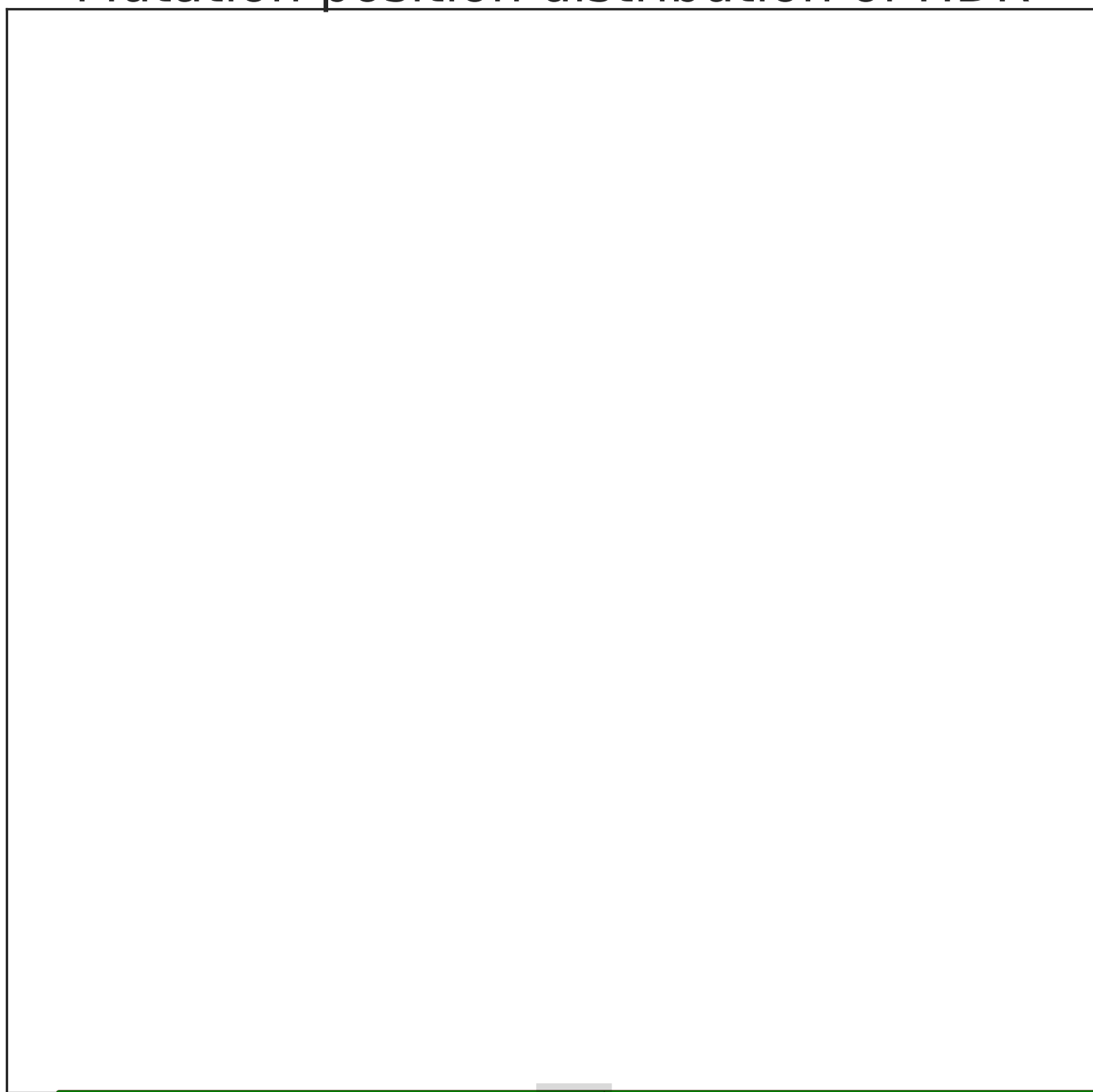


Mutation position distribution of HDR

Sequences: % Total (% HDR, no.)

0.0% (nan% , 0)



0 40 80 120 160 200 240

Reference amplicon position (bp)

- Insertions
- Deletions
- Substitutions
- - - - Predicted cleavage position
- █ sgRNA