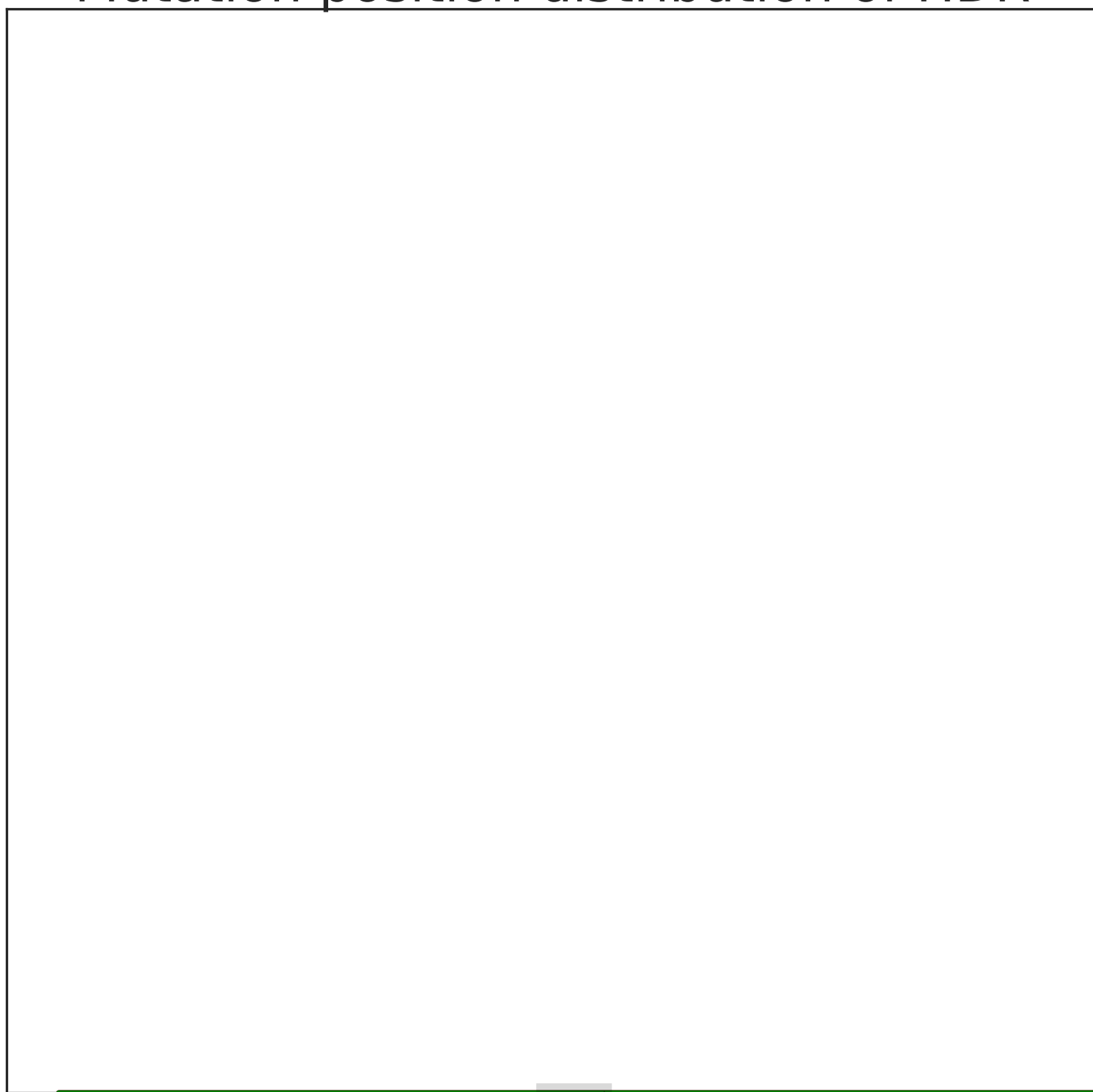


Mutation position distribution of HDR

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