Mutation position distribution of HDR

Sequences: % Total (% HDR, no.)

0.0% (nan% , 0)

Reference amplicon position (bp)
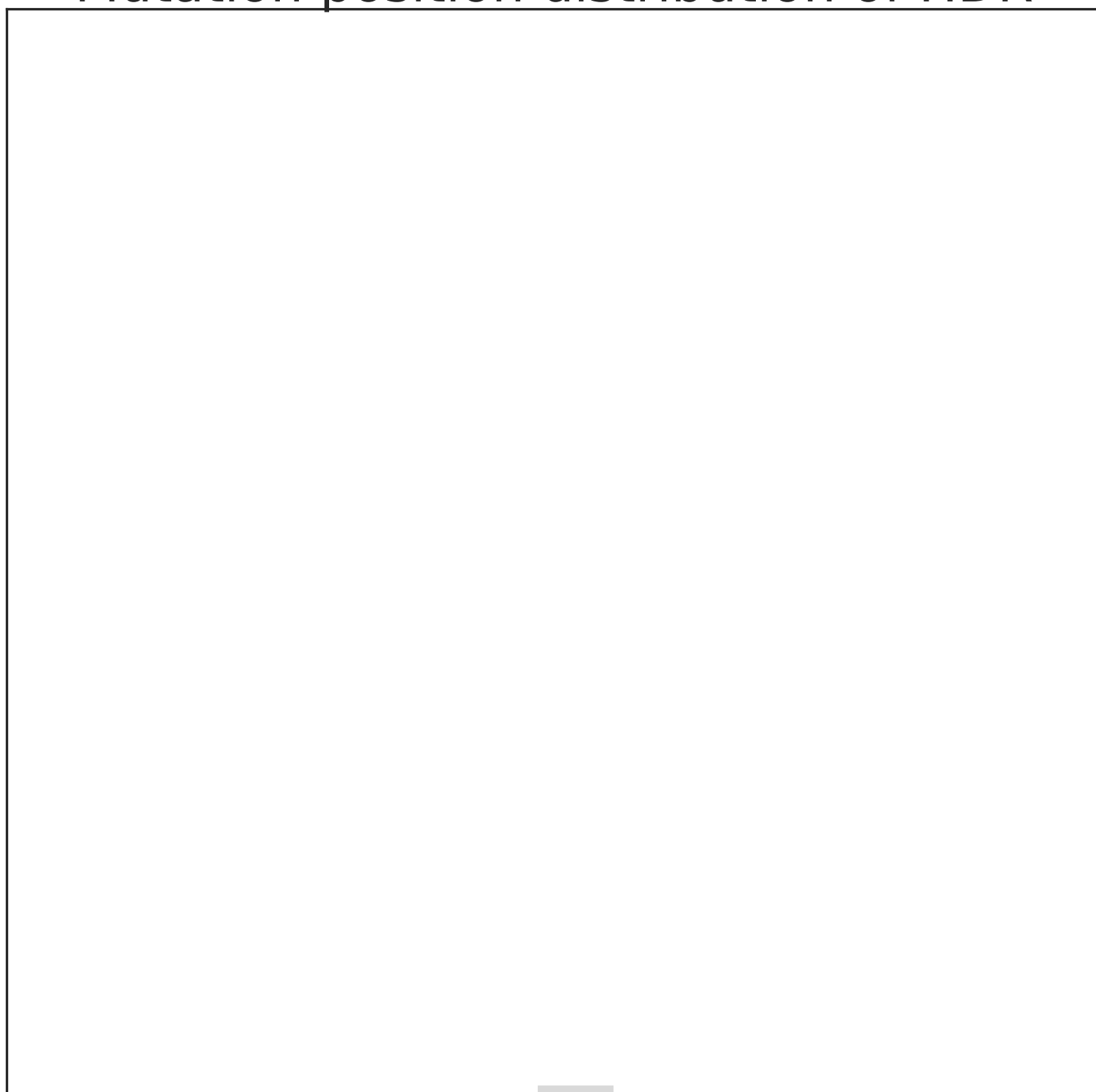
—— Insertions
—— Deletions
—— Substitutions
---- Predicted cleavage position
▬ sgRNA