G G G C T C A T G A T C A T C G G C C G G A G C G G C G G C G T G G C G A C C T - Reference
sgRNA

G G G C T C A T G A T C A T C G G C C G G A G C G G C G G C G T G G C G A C C T - 62.04% (51323 reads)
G G G C T C A T G A T C A T C G G C C G - A G C G G C G G C G T G G C G A C C T - 23.60% (19520 reads)
G G G C T C A T G A T C A T C G G - - - - A G C G G C G G C G T G G C G A C C T - 1.79% (1483 reads)
G G G C T C A T G A T C A T C G G C - - G A G C G G C G G C G T G G C G A C C T - 1.50% (1240 reads)
G G G C T C A T G A T C A - - - - - - - G A G C G G C G G C G T G G C G A C C T - 1.25% (1038 reads)
G G G C T C A T G A T C A T C G - - - - A G C G G C G G C G T G G C G A C C T - 0.84% (694 reads)
G G G C T C A T G A T C - - - - - - - - G A G C G G C G G C G T G G C G A C C T - 0.49% (402 reads)
G G G C T C A T G A T C A T C G G - - - G A G C G G C G G C G T G G C G A C C T - 0.44% (360 reads)
G G G C T C A T G A T C A T - - - - - - G A G C G G C G G C G T G G C G A C C T - 0.26% (211 reads)

bold Substitutions

☐ Insertions

- Deletions

------ Predicted cleavage position