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

A G A T A G G T G G A C T A T G C A G T C A A T G G A C G A A C T G C C G C T A - 89.47% (48288 reads)
A G A T A G G T G G A C T A T G C A G T T C A A T G G A C G A A C T G C C G C T - 0.35% (190 reads)
A G G T A G G T G G A C T A T G C A G T C A A T G G A C G A A C T G C C G C T A - 0.31% (169 reads)
A G A T A G G T G G A C T A T G C A G T C A C T G G A C G A A C T G C C G C T A - 0.30% (161 reads)
A G A T A G G T G G A C T A T G C A G T C A A T G G G C G A A C T G C C G C T A - 0.27% (144 reads)
A G A T A G G T G G A C T A T G C A G T C A A T G G A C G A G C T G C C G C T A - 0.26% (139 reads)
A G A T A G G T G G G C T A T G C A G T C A A T G G A C G A A C T G C C G C T A - 0.23% (122 reads)
A G A T A G G T G G A C T A C G C A G T C A A T G G A C G A A C T G C C G C T A - 0.22% (120 reads)
A G A T A G G T G G A C T A T G C A G T C A A T G G A C G G A C T G C C G C T A - 0.22% (119 reads)
A G A T A G G T G G A C T A T G C G G T C A A T G G A C G A A C T G C C G C T A - 0.21% (113 reads)
A G A T A G G T G G A C T A T G C A G T C A G T G G A C G A A C T G C C G C T A - 0.20% (109 reads)

bold Substitutions

☐ Insertions

- Deletions

------ Predicted cleavage position