


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

sgRNA 

C G C A C C T C A T G G A A T C C C T T C T G C A G C A C C T G G A T C G C T T - 96.38% (23648 reads)
C G C A C C T C A T G G A A T C C C T T C T G C A G C A C C T G G A T C G **A** T T - 0.29% (70 reads)
C G C A C C T C A T G G A A T C C C T T C T G C A G C A C C T G **T** A T C G C T T - 0.24% (59 reads)

bold Substitutions
 Insertions
- Deletions