



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

G G A A G G G C C T G A G T C C G A G C A G A A G A A G A A G G G C T C C C A T - 70.40% (17579 reads)  
 G G A A G G G C C T G A G T **T T** G A G C A G A A G A A G A A G G G C T C C C A T - 21.56% (5384 reads)  
 G G A A G G G C C T G A G T **T** C G A G C A G A A G A A G A A G G G C T C C C A T - 4.40% (1099 reads)  
 G G A A G G G C C T G A G T **G T** G A G C A G A A G A A G A A G G G C T C C C A T - 0.29% (73 reads)  
 G G A A G G G C C T G A G T **G** C G A G C A G A A G A A G A A G G G C T C C C A T - 0.26% (65 reads)

**bold** Substitutions  
 Insertions  
 - Deletions