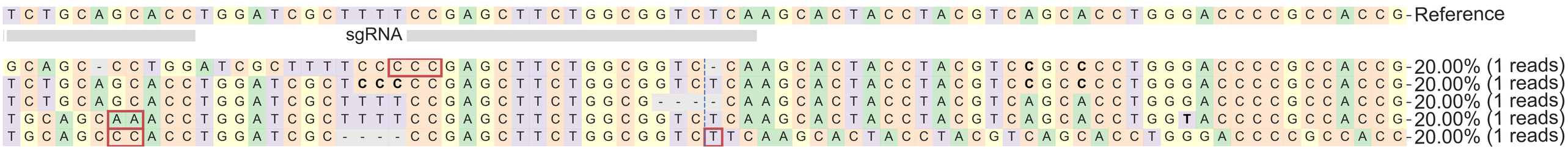
TCTGCAGCACCTGGATCGCTTTTCCGAGCTTCTGGCGGTCTCAAGCACTACCTACGTCAGCACCTGGGACCCCGCCACCG-Reference
sgRNA
GCAGC-CCTGGATCGCTTTTCCCCCCGAGCTTCTGGCGGTC-CAAGCACTACCTACGTCCGCCCCTGGGACCCCGCCACCG-20.00% (1 reads)
TCTGCAGCACCTGGATCGCTCCCCCGAGCTTCTGGCGGTCTCAAGCACTACCTACGTCCGCCCCTGGGACCCCGCCACCG-20.00% (1 reads)
TCTGCAGCACCTGGATCGCTTTTCCGAGCTTCTGGCG---CAAGCACTACCTACGTCAGCACCTGGGACCCCGCCACCG-20.00% (1 reads)
TGCAGCAAACCTGGATCGCTTTTCCGAGCTTCTGGCGGTCTCAAGCACTACCTACGTCAGCACCTGGTACCCCGCCACCG-20.00% (1 reads)
TGCAGCCCACCTGGATCGC----CCGAGCTTCTGGCGGTCTTCAAGCACTACCTACGTCAGCACCTGGGACCCCGCCACC-20.00% (1 reads)
bold Substitutions
Insertions
- Deletions
------ Predicted cleavage position