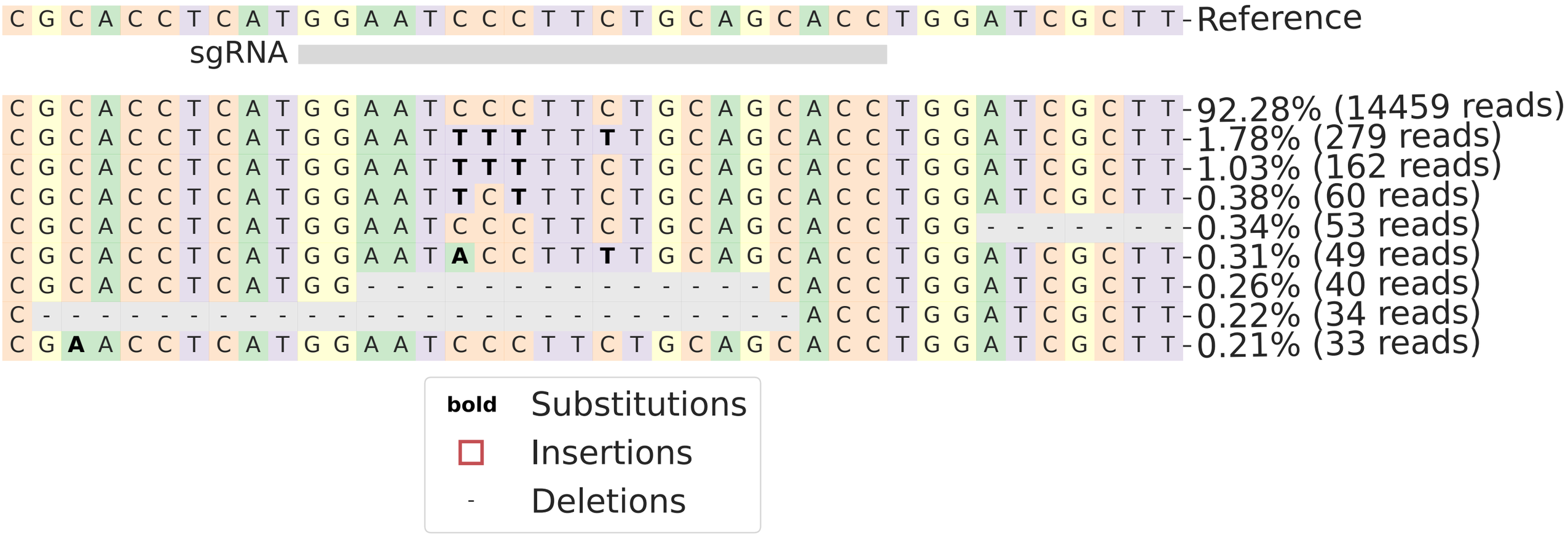
CGCACCTCATGGAATCCCTTCTGCAGCACCTGGATCGCTT-Reference
sgRNA
CGCACCTCATGGAATCCCTTCTGCAGCACCTGGATCGCTT-92.28% (14459 reads)
CGCACCTCATGGAATTTTTTTTGCAGCACCTGGATCGCTT-1.78% (279 reads)
CGCACCTCATGGAATTTTTTCTGCAGCACCTGGATCGCTT-1.03% (162 reads)
CGCACCTCATGGAATTCTTTCTGCAGCACCTGGATCGCTT-0.38% (60 reads)
CGCACCTCATGGAATCCCTTCTGCAGCACCTGG-------0.34% (53 reads)
CGCACCTCATGGAATACCTTTGCAGCACCTGGATCGCTT-0.31% (49 reads)
CGCACCTCATGG-----------CACCTGGATCGCTT-0.26% (40 reads)
C----------------ACCTGGATCGCTT-0.22% (34 reads)
CGAACCTCATGGAATCCCTTCTGCAGCACCTGGATCGCTT-0.21% (33 reads)
bold Substitutions
Insertions
- Deletions