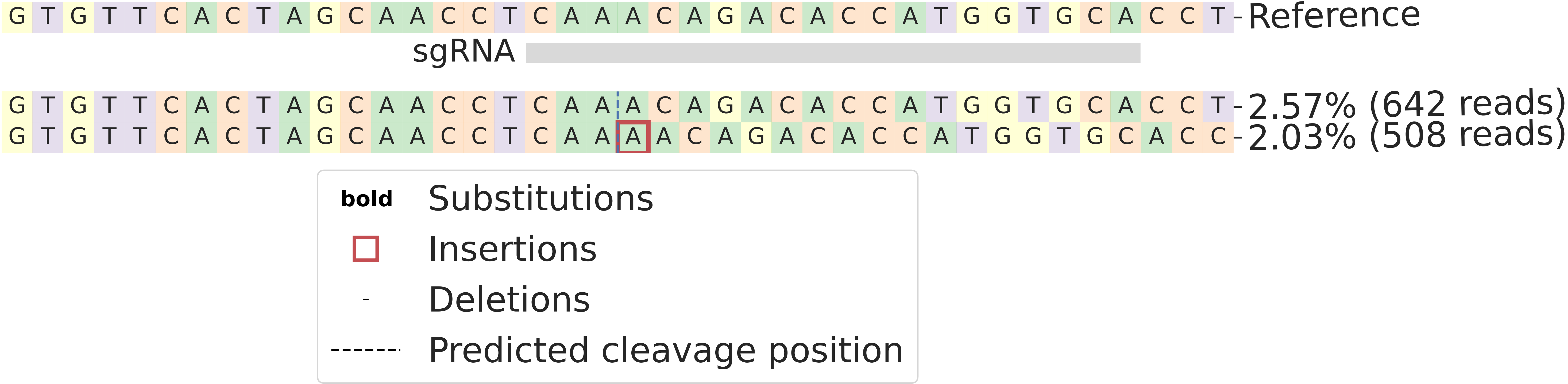

GTGTTCACTAGCAACCTCAAACAGACACCATGGTGCACCT - Reference

sgRNA

GTGTTCACTAGCAACCTCAAACAGACACCATGGTGCACCT - 2.57% (642 reads)

GTGTTCACTAGCAACCTCAAAACAGACACCATGGTGCACC - 2.03% (508 reads)

**bold** Substitutions

☐ Insertions

\- Deletions

\------ Predicted cleavage position