GCAGTCATCTTAGTCATTACATGAGGTGTTCGTTGTAACT - Reference
PE Extension
PE spacer sgRNA
GCAGTCATCTTAGTCATTACATGAGGTGTTCGTTGCACCG - 0.44% (109 reads)
GCAGTCATCTTAGTCATTACATGAGGTGTTCGTTGCAACT - 0.39% (98 reads)
GCAGTCATCTTAGTCATTACATGAGGTGTTCGTTGCACCG - 0.26% (65 reads)
bold Substitutions
Insertions
- Deletions
Predicted cleavage position