Strong support multiple
variants if positioned called independently,
but reads all have identical sequence

Weak support for true insertion
due to alignment error
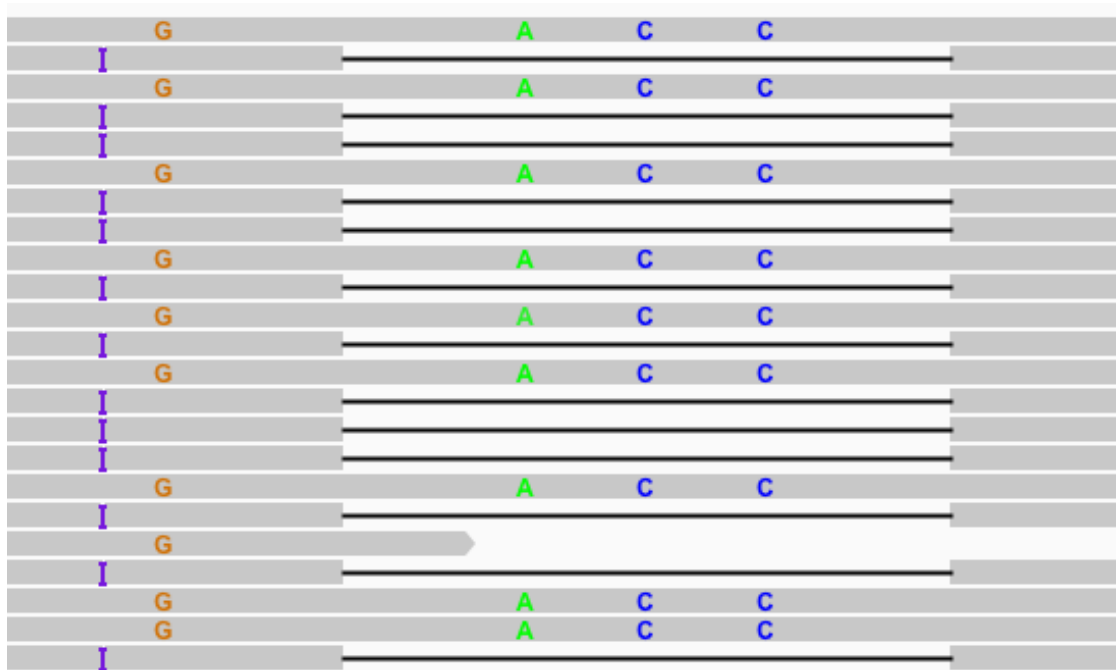
Haplotype caller: single complex substitution; TCGTGTT to TGGTACC
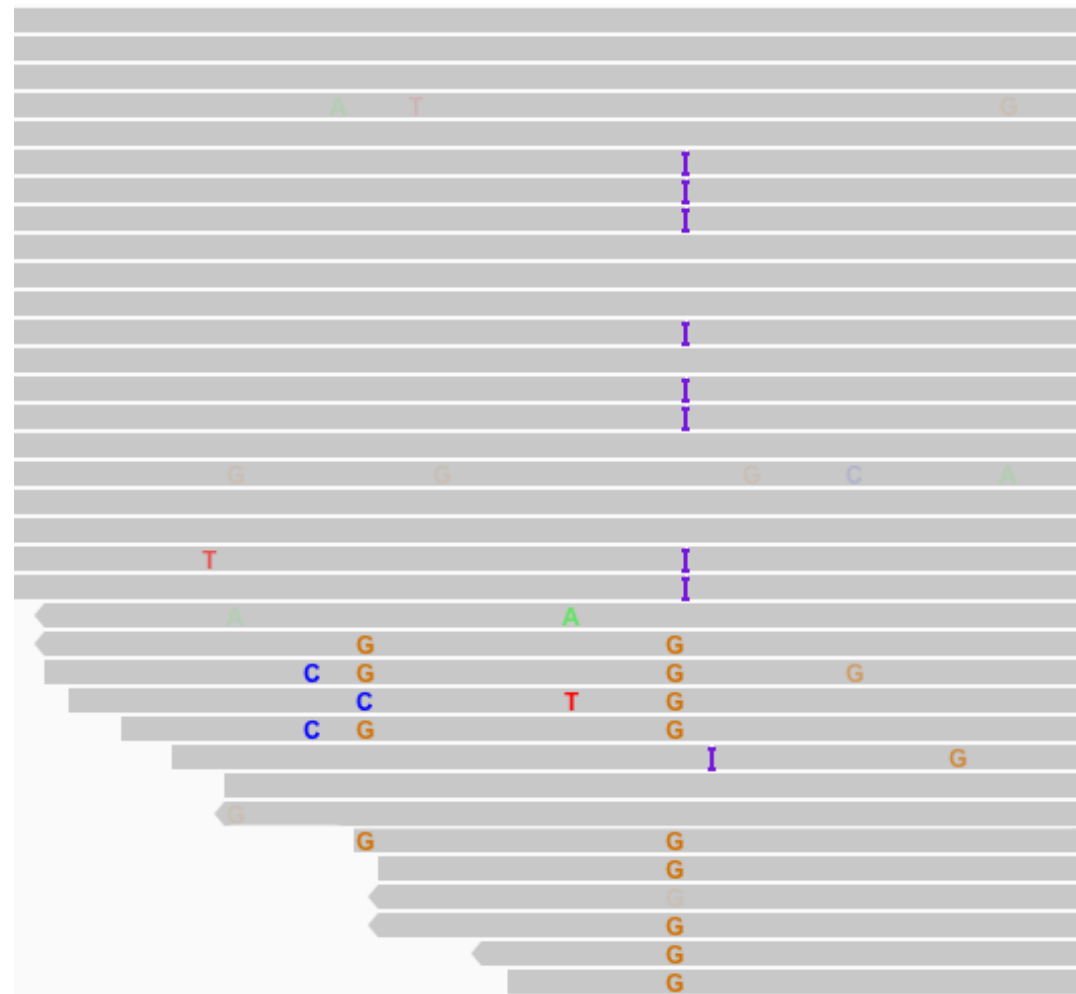
Haplotype caller can use misaligned
reads to call the insertion