

Founder molecule ATGTACATCTACAGTCAGGT

Technical errors due to PCR and sequencing introduce 1 deletions and 1 substitution

HiFi read 1 ATGTACTTCTACAGTCAGGT

HiFi read 2 TGTACATCTACAGTCAGGT

isoseq3 tag --design 10B-6U-T: clips 16 bases from 5'

TAG result 1 ATGTACTTCTACAGTC

TAG result 2 TGTACATCTACAGTCA

isoseq3 dedup: alignment identifies 1 mismatch and tags are shifted by 1 base pair

Alignment 1 ATGTACTTCTACAGTC

Alignment 2 TGTACATCTACAGTCA

Tag shift →

Mismatches

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