

Spike recovery is evaluated on the alignments of spiked read sets or *de novo* assembled contig sets against viral database

Short reads from HTS

Original_readsR1_539895

Original_readsR2_539897

Trimming (keep read positions 10-110)

HS_trimmingR1_539358 (yielded Trimmed_readsR1_539360)

HS_trimmingR2_539359 (yielded Trimmed_readsR2_539361)

