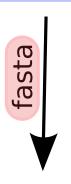Structural variant(s) (.vcf) query

Bedtools.getfasta

fasta

Reference RNA-seq

BLASTN_vdb

blastn_vdb -db SRR307006 -query
XM_006525543.2.fa -outfmt 15 -out test.bout15

sam

Samtools.view -bS .bam

bam

Bedtools.bamtobed

bed

Bedtools.coverage

Raw / differential histogram