

status



normal



paired



tumor

format



BAM



CRAM

peak\_vmem\_MB

CNV

ASCAT

CNVKIT\_BATCH

Preprocessing

APPLYBQSR

BASERECALIBRATOR

GATHERBQSRREPORTS

GATK4\_MARKDUPLICATES

INDEX\_CRAM

MERGE\_CRAM

MOSDEPTH

SAMTOOLS\_BAMTOCRAM

SAMTOOLS\_STATS

SAMTOOLS\_STATS\_CRAM

SNP/Indels

FREEBAYES

GETPILEUPSUMMARIES\_NORMAL

GETPILEUPSUMMARIES\_TUMOR

HAPLOTYPECALLER

MSISENSORPRO\_MSI\_SOMATIC

MUTECT2

STRELKA\_SINGLE

STRELKA\_SOMATIC

SV

MANTA\_GERMLINE

MANTA\_SOMATIC

MERGE\_MANTA\_DIPLOID

TIDBIT\_SV