



Tool Reference

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Tools By Category

Category	Tool
FASTQ/BAM Processing	<ul style="list-style-type: none">• applybqsr• bam2fq• bamsort• bqsr• fq2bam (BWA-MEM + GATK)• fq2bamfast (BWA-MEM + GATK)• fq2bam_meth

	<ul style="list-style-type: none"> • markdup • minimap2 (Beta)
Variant Calling	<ul style="list-style-type: none"> • deepsomatic • deepvariant • deepvariant_germline • germline (GATK Germline Pipeline) • haplotypcaller • mutectcaller • pacbio_germline (Beta) • postpon • prepon • somatic (Somatic Variant Caller)
RNA	<ul style="list-style-type: none"> • rna_fq2bam • starfusion
Quality Control	<ul style="list-style-type: none"> • bammetrics • collectmultiplemetrics
Variant Processing	<ul style="list-style-type: none"> • dbsnp
GVCF Processing	<ul style="list-style-type: none"> • genotypegvcf • indexgvcf

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