



Available alignment modes for Bismark (v0.21.0)



	library type	temporary files	output file	multi-core	Bowtie 2		HISAT2	
					FastQ	FastA	FastQ	FastA
Single-End (default)	directional	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
	PBAT	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
	non-directional	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
	SLAM-seq (experimental)	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
Paired-End (default)	directional	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
	PBAT	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
	non-directional	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes
	SLAM-seq (experimental)	uncompressed/ --gzip	BAM/SAM	1/yes	yes	yes	yes	yes