



Available alignment modes for Bismark (v0.11.1)



	library type	(option '--gzip')	(option '--bam')	Bowtie 1		Bowtie 2	
		temporary files	output file	FastQ	FastA	FastQ	FastA
Single-End (default)	directional	--gzip	SAM	yes	yes	yes	yes
			BAM	yes	yes	yes	yes
		uncompressed	SAM	yes	yes	yes	yes
			BAM	yes	yes	yes	yes
	non-directional	--gzip	SAM	yes	yes	yes	yes
			BAM	yes	yes	yes	yes
		uncompressed	SAM	yes	yes	yes	yes
			BAM	yes	yes	yes	yes
Paired-End (default)	directional	--gzip	SAM	yes	-	yes	-
			BAM	yes	-	yes	-
		uncompressed	SAM	yes	yes	yes	yes
			BAM	yes	yes	yes	yes
	non-directional	--gzip	SAM	yes	-	yes	-
			BAM	yes	-	yes	-
		uncompressed	SAM	yes	yes	yes	yes
			BAM	yes	yes	yes	yes
Single-End	PBAT	uncompressed	SAM	yes	-	yes	-
			BAM	yes	-	yes	-
Paired-End	PBAT	uncompressed	SAM	yes	-	yes	-
			BAM	yes	-	yes	-