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Running Status

Selecting Tools for Basic Analysis of DDBJ ANNOTATION PIPELINE

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Reference Genome Mapping

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Mapping tool can be selected up to two.
 When blat is included, only Single-end query is executable.

Run	Tool	Help	Version	Base calls		Analysis					Output format			Visualization	Comment
				Base space	Color space	Paired end	SNP	Indel	Depth	Coverage	error rate	.gff	.bed		
<input type="checkbox"/>	BLAT		34						✓	✓				✓	Single analysis only
<input type="checkbox"/>	Mag		0.7.1			✓	✓	✓	✓	✓		✓	✓		
<input type="checkbox"/>	bwa		0.4.9			✓		✓	✓	✓				✓	
<input type="checkbox"/>	SSAHA2		2.3.0.1			✓	✓	✓							SNP is single analysis only
<input type="checkbox"/>	SOAP		2.1.8			✓	✓	✓	✓	✓					
<input type="checkbox"/>	Bowtie (SAMtools)		0.12.0 (0.1.7)	✓	✓	✓	✓				✓			✓	
<input type="checkbox"/>	TopHat		1.0.11 (BETA)	✓		✓								✓	

De Novo Assembly