History for:

🗏 Illumina High-Throughput Sequencing Import log

Illumina High-Throughput Sequencing Import 2.0

29 Jun 2023 12:26:49

CLC Genomics Server 23.0.4 Version:

Modified by:

Select files /disk1/CLC-Upload/markop/RawFASTQ/AM24_1_1.fastq.gz, /disk1/CLC-

Upload/markop/RawFASTQ/AM24_1_2.fastq.gz, /disk1/CLC-Upload/ markop/RawFASTQ/AM24_2_1.fastq.gz, /disk1/CLC-Upload/markop/ RawFASTQ/AM24_2_2.fastq.gz, /disk1/CLC-Upload/markop/RawFASTQ/

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NM48_INF_3_2.fastq.qz

Discard read names No Discard quality scores No Paired reads Yes

Read orientation Forward Reverse

Minimum distance Maximum distance 1,000 Remove failed reads Yes

Quality scores NCBI/Sanger or Illumina Pipeline 1.8 and later

Join reads from different lanes No Custom read structure No Comments:	MiSeq de-multiplexing	No
	Join reads from different lanes	No
Comments:	Custom read structure	No
	Comments:	