

10x-ALSmice-SingleNucleiExomeEnrichment-SNAM-justReadSeg - SNAM-5

SUCCESSFUL

Send Log Files

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▼ Analysis Overview

Status

Thumbnails

Display All

➤ Read Segmentation

➤ Data

Display All

Status

Analysis

10x-ALSmice-SingleNucleiExomeEnrichment-SNAM-justReadSeg - SNAM-5

Analysis ID

13174

From Multi-Job

[13169](#)

Status

SUCCESSFUL: 4 tasks finished

Created By

etseng

Date Created

2025-11-18, 02:00:16 PM

Date Updated

2025-11-18, 03:01:57 PM

Application

Read Segmentation

SMRT Link Version

25.4.0.278749

Inputs

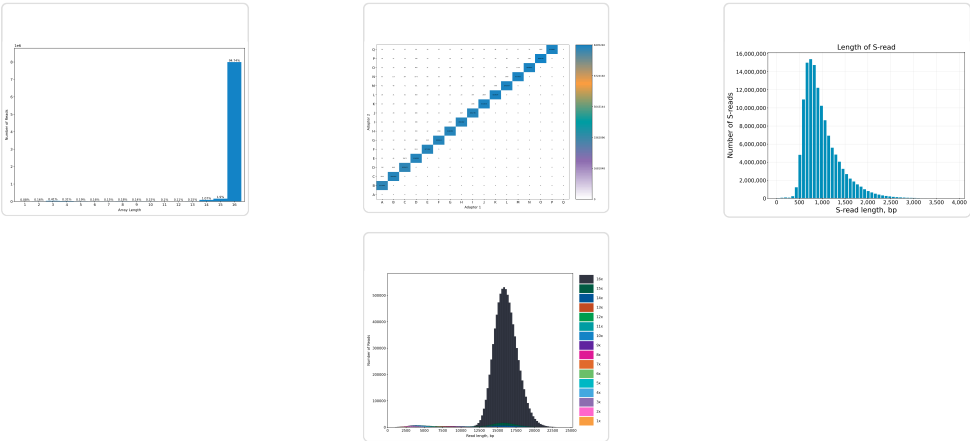
Data Type	Name
BarcodeSet	Barcodes: MAS-Seq Adapter v1 (MAS16)
ConsensusReadSet	HiFi reads: SNAM-5-Cell5 (all samples)

Path

/pbi/flash/smartlink/smartlink-appslabht/smartlink/userdata/jobs_root/0000/0000013/0000013174

➤ Analysis Parameters

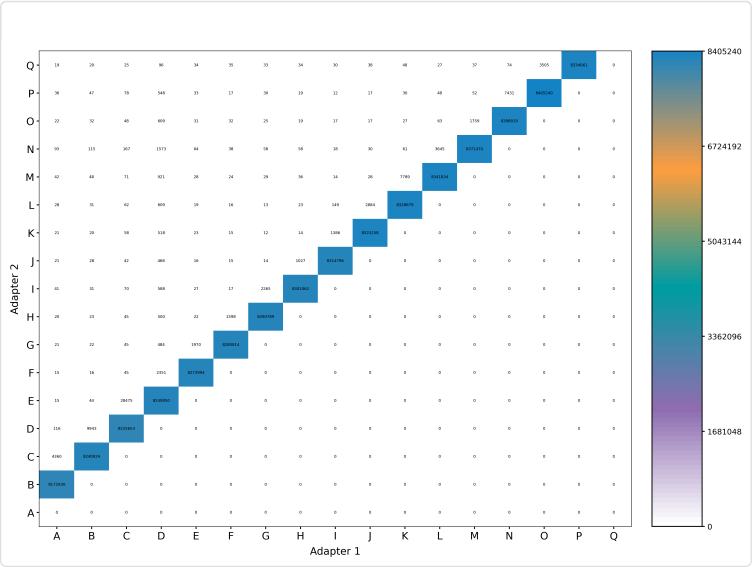
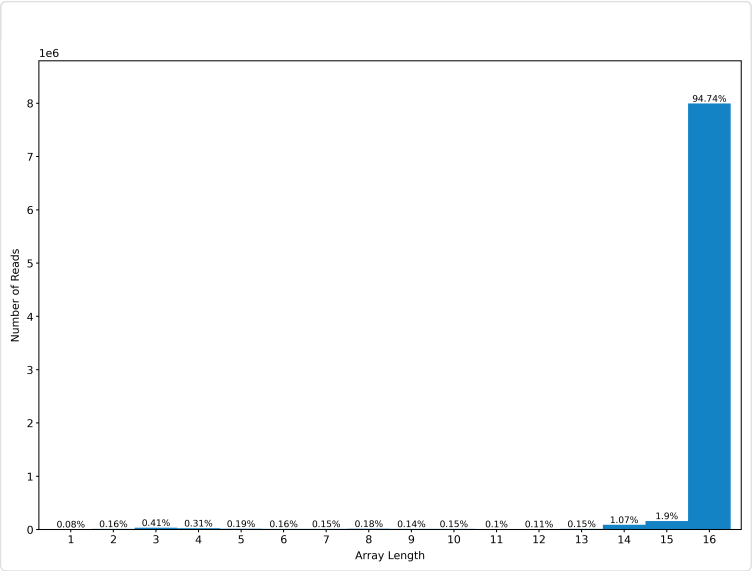
Thumbnails



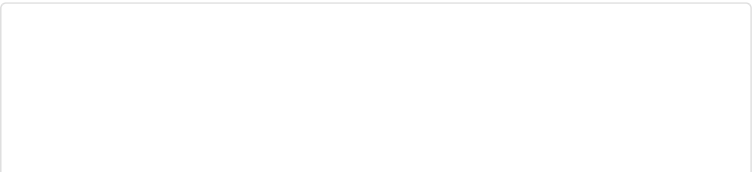
Read Segmentation

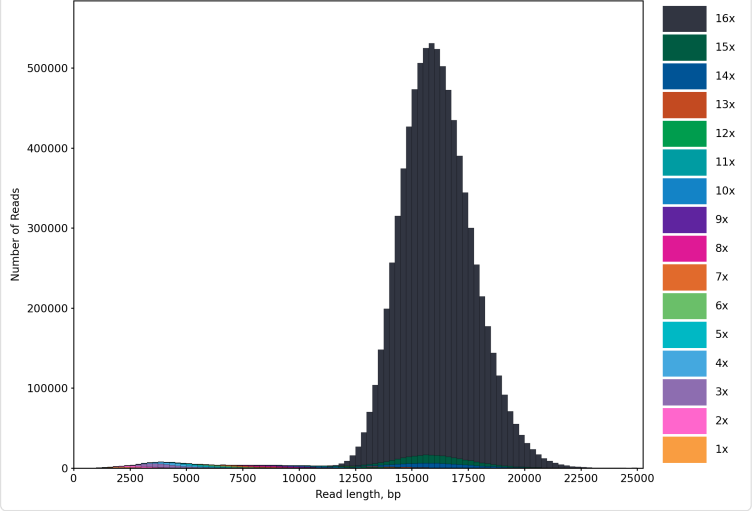
Value	Analysis Metric
8,442,542	Reads
132,845,136	Segmented reads (S-reads)
992	Mean length of S-reads
94.72 %	Percent of reads with full arrays
15.74	Mean array size (concatenation factor)

Segmentation Statistics

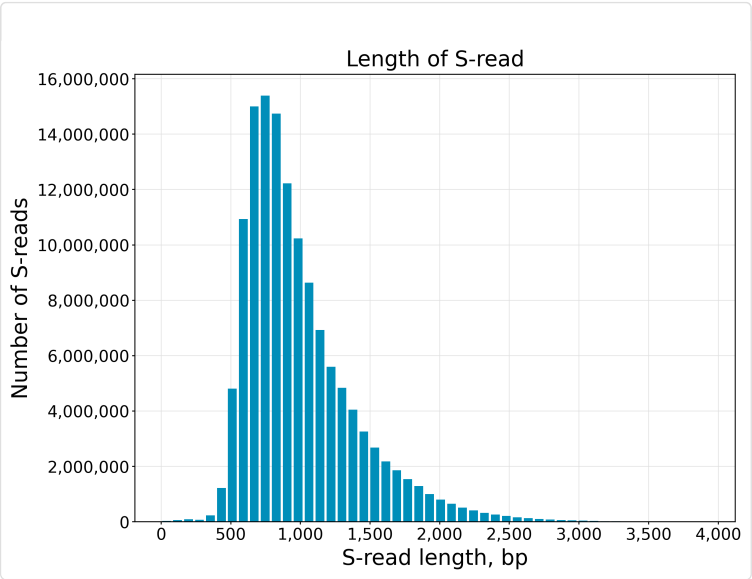


Length of Reads





S-read Length Distribution



File Downloads

Edit Output File Name Prefix

Example:analysis-SNAM-5-13174

File ⬆️⬆️	Type ⬆️⬆️	Size	Path
Analysis Log	log	15 KB	 /pbi/flash/smartlink/smartlink-...
Non-passing read:	bam	3 GB	 /pbi/dept/secondary/siv/smr...
Segmented Reads	bam	90 GB	 /pbi/dept/secondary/siv/smr...
SMRT Link Log	log	12 KB	 /pbi/flash/smartlink/smartlink-...