

10x-ALSmice-SingleNucleiExomeEnrichment-SNAM-justReadSeg - SNAM-3

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-3

Analysis ID

13172

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, 02:57:06 PM

Application

Read Segmentation

SMRT Link Version

25.4.0.278749

Inputs

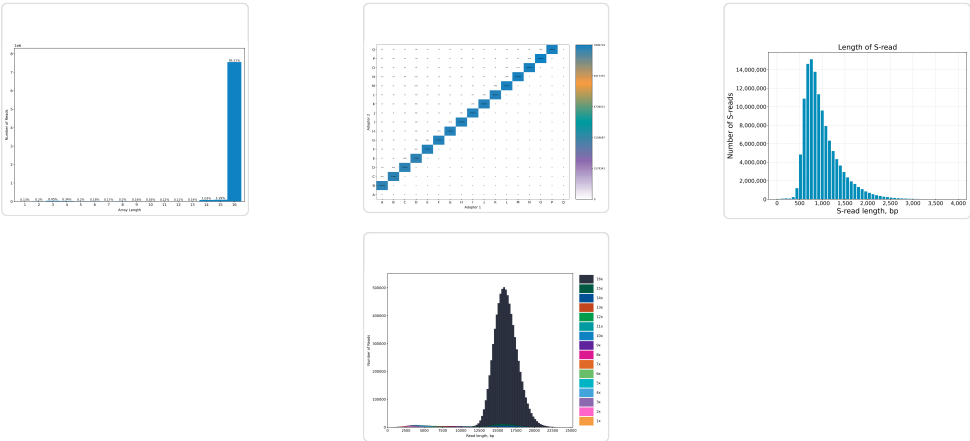
Data Type ⬆	Name ⬆
BarcodeSet	Barcodes: MAS-Seq Adapter v1 (MAS16)
ConsensusReadSet	HiFi reads: SNAM-3-Cell3 (all samples)

Path

/pbi/flash/smartlink/smartlink-appslabht/smartlink/userdata/jobs_root/0000/0000013/0000013172

➤ Analysis Parameters

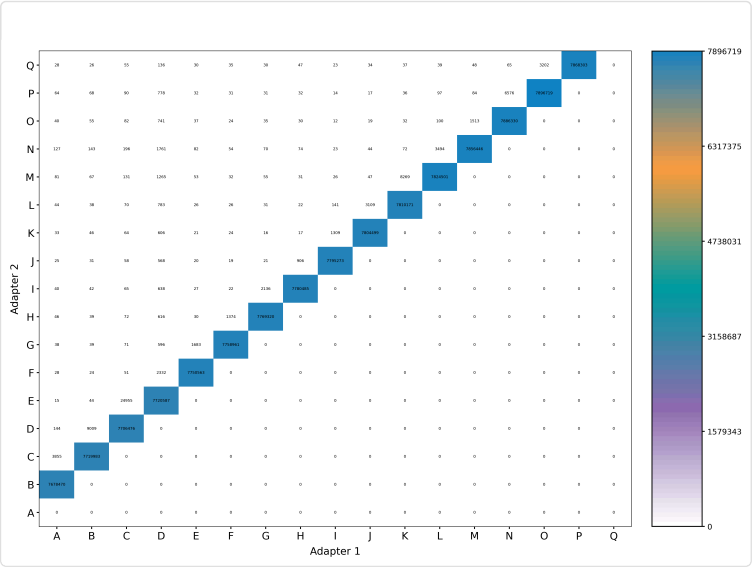
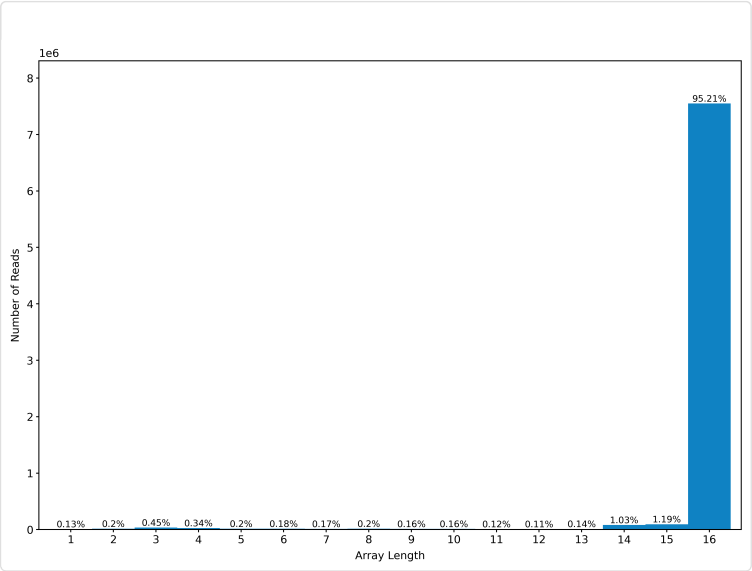
Thumbnails



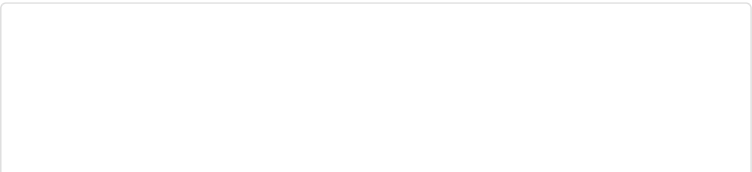
Read Segmentation

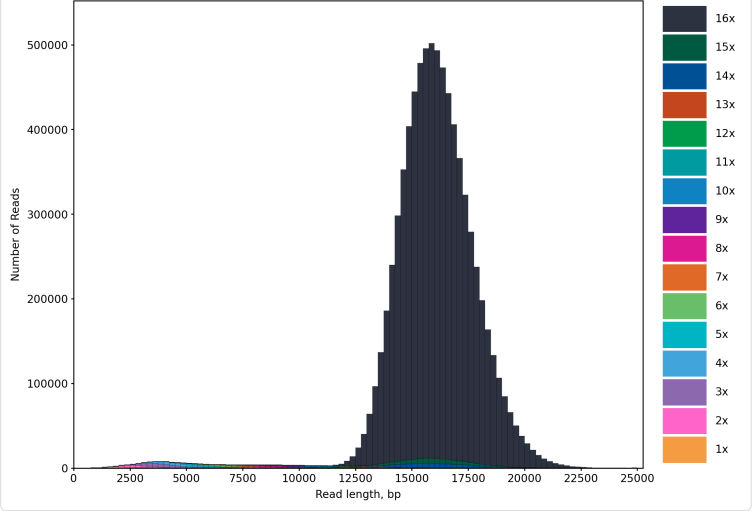
Value	Analysis Metric
7,933,609	Reads
124,627,087	Segmented reads (S-reads)
992	Mean length of S-reads
95.19 %	Percent of reads with full arrays
15.71	Mean array size (concatenation factor)

Segmentation Statistics

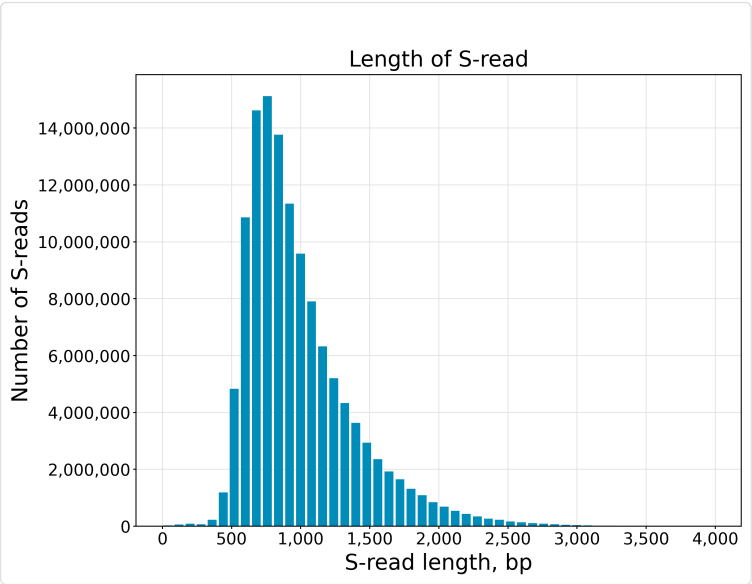


Length of Reads





S-read Length Distribution



File Downloads

Edit Output File Name Prefix

Example:analysis-SNAM-3-13172

File ↕	Type ↕	Size	Path
Analysis Log	log	15 KB	 /pbi/flash/smrtlink/smrtlink-...
Non-passing read:	bam	2 GB	 /pbi/dept/secondary/siv/smr...
Segmented Reads	bam	84 GB	 /pbi/dept/secondary/siv/smr...
SMRT Link Log	log	12 KB	 /pbi/flash/smrtlink/smrtlink-...