

10x-ALSmice-SingleNucleiExomeEnrichment-SNAM-justReadSeg - SNAM-1

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-1

Analysis ID

13170

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, 02:59:16 PM

Application

Read Segmentation

SMRT Link Version

25.4.0.278749

Inputs

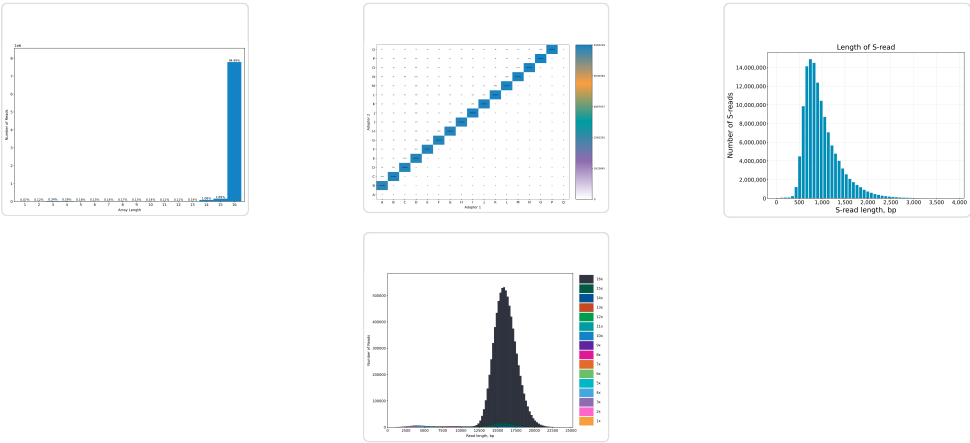
Data Type	Name
BarcodeSet	Barcodes: MAS-Seq Adapter v1 (MAS16)
ConsensusReadSet	HiFi reads: SNAM-1-Cell1 (all samples)

Path

/pbi/flash/smartlink/smartlink-appslabht/smartlink/userdata/jobs_root/0000/0000013/0000013170

Analysis Parameters

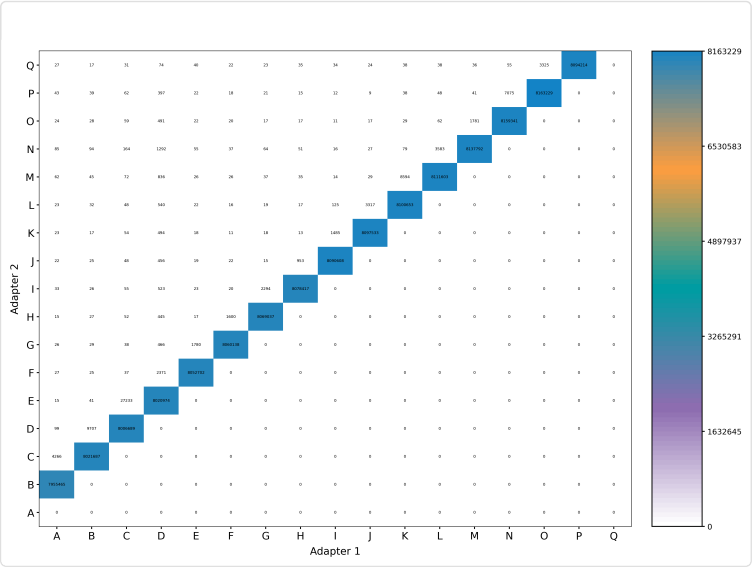
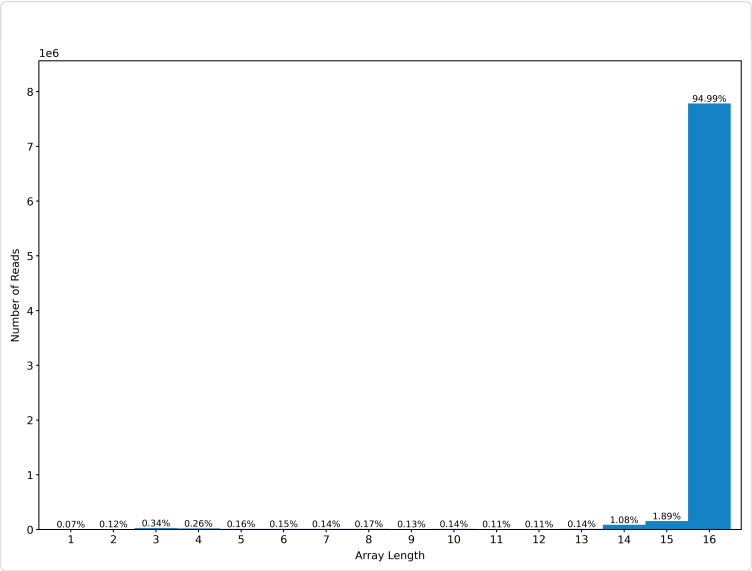
Thumbnails



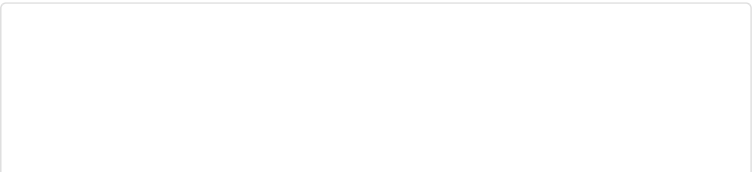
Read Segmentation

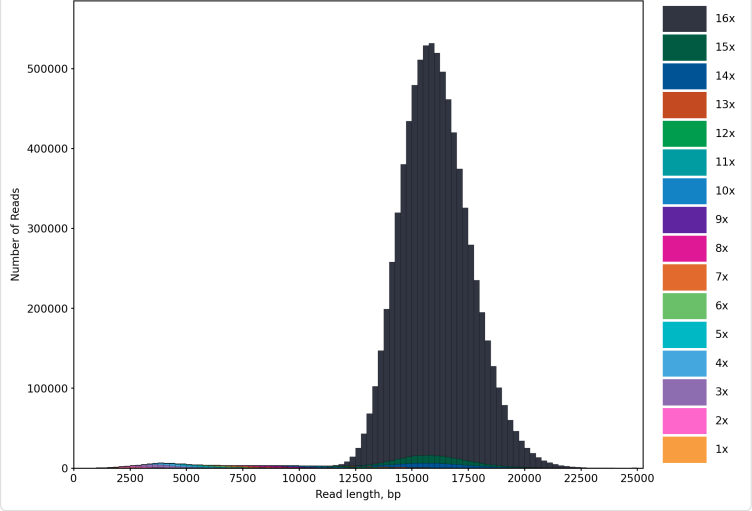
Value	Analysis Metric
8,197,233	Reads
129,220,082	Segmented reads (S-reads)
988	Mean length of S-reads
94.98 %	Percent of reads with full arrays
15.76	Mean array size (concatenation factor)

Segmentation Statistics

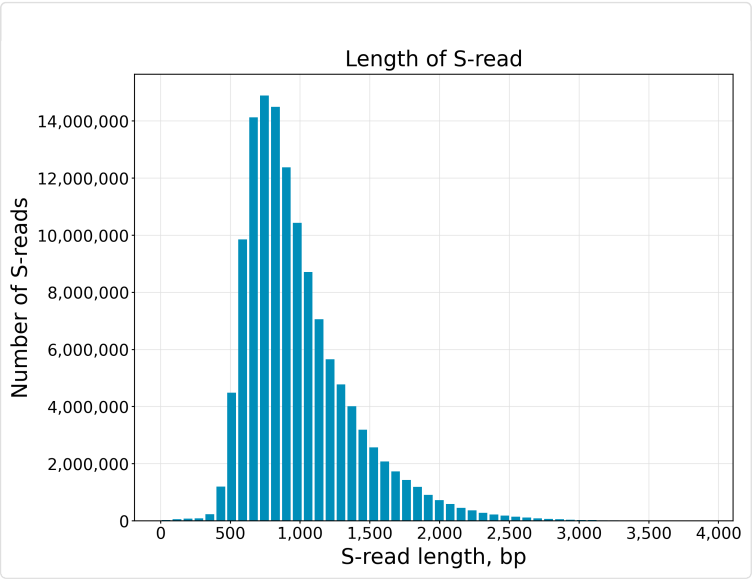


Length of Reads





S-read Length Distribution



File Downloads

Edit Output File Name Prefix

Example:analysis-SNAM-1-13170

File ⬆	Type ⬆	Size	Path
Analysis Log	log	15 KB	📁 /pbi/flash/smartlink/smartlink-c
Non-passing read:	bam	3 GB	📁 /pbi/dept/secondary/siv/smr
Segmented Reads	bam	87 GB	📁 /pbi/dept/secondary/siv/smr
SMRT Link Log	log	12 KB	📁 /pbi/flash/smartlink/smartlink-c