

10x-ALSmice-SingleNucleiExomeEnrichment-SNAM-justReadSeg - SNAM-6

SUCCESSFUL

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▼ Analysis Overview

Status

Thumbnails

Display All

➤ Read Segmentation

➤ Data

Display All

Status

Analysis	10x-ALSmice-SingleNucleiExomeEnrichment-SNAM-justReadSeg - SNAM-6
Analysis ID	13175
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, 02:57:16 PM
Application	Read Segmentation
SMRT Link Version	25.4.0.278749

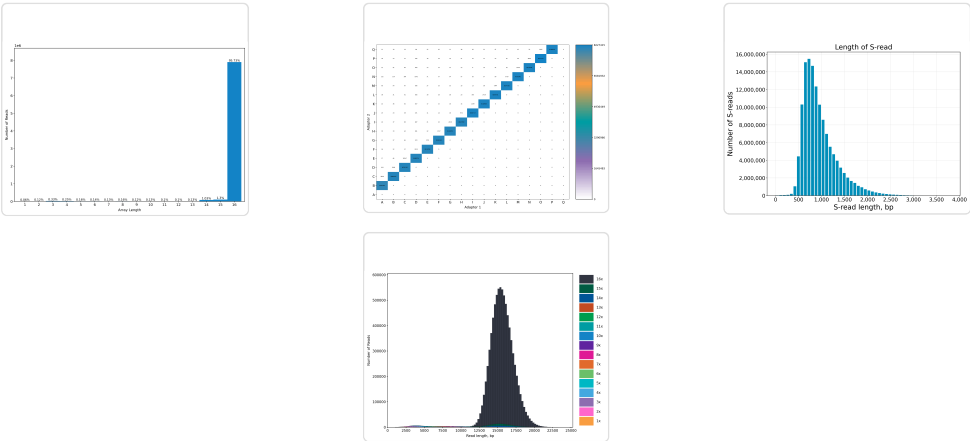
Inputs

Data Type	Name
BarcodeSet	Barcodes: MAS-Seq Adapter v1 (MAS16)
ConsensusReadSet	HiFi reads: SNAM-6-Cell6 (all samples)

Path /pbi/flash/smrtlink/smrtlink-appslabht/smrtlink/userdata/jobs_root/0000/0000013/0000013175

➤ Analysis Parameters

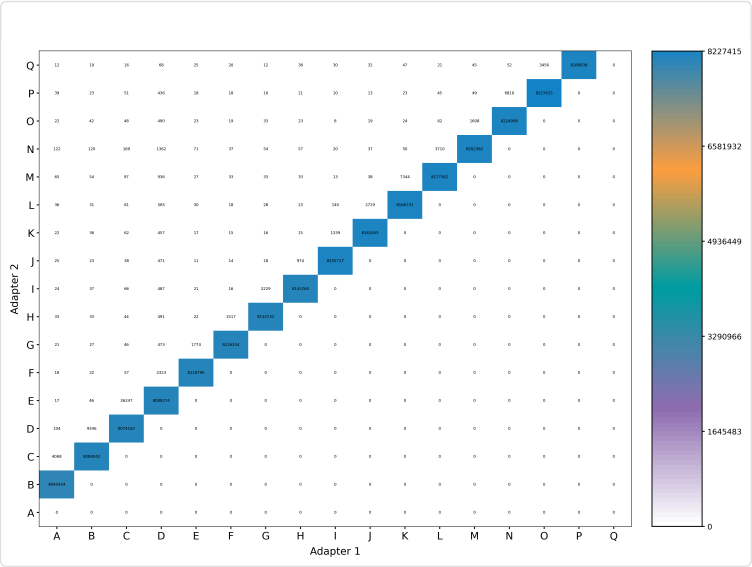
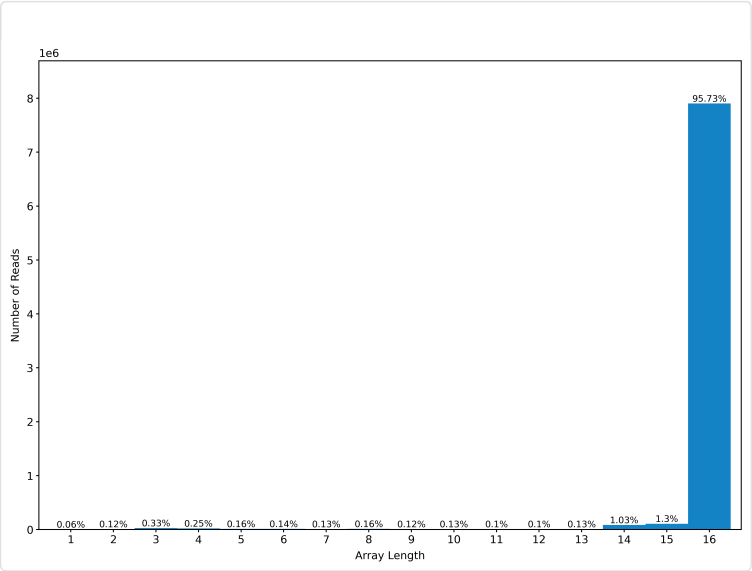
Thumbnails



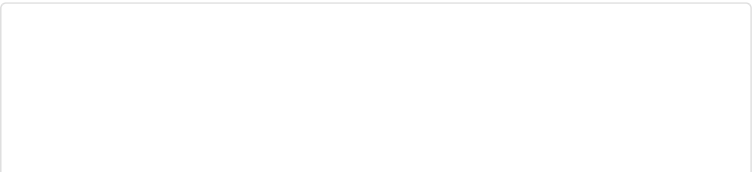
Read Segmentation

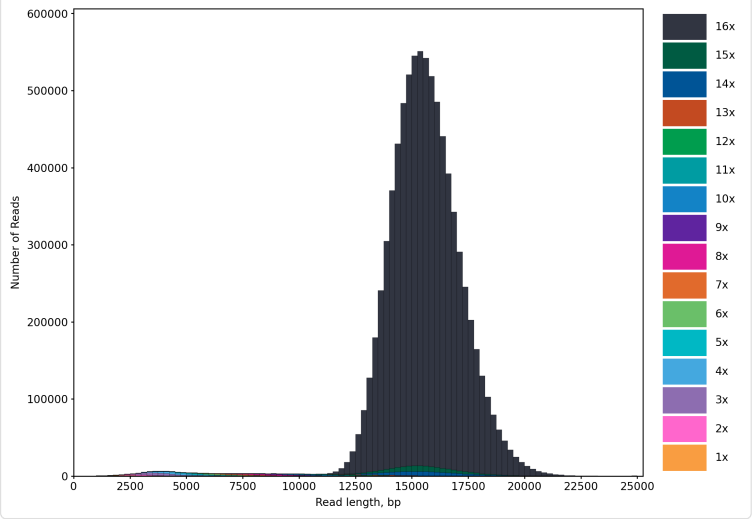
Value	Analysis Metric
8,258,305	Reads
130,317,641	Segmented reads (S-reads)
960	Mean length of S-reads
95.72 %	Percent of reads with full arrays
15.78	Mean array size (concatenation factor)

Segmentation Statistics

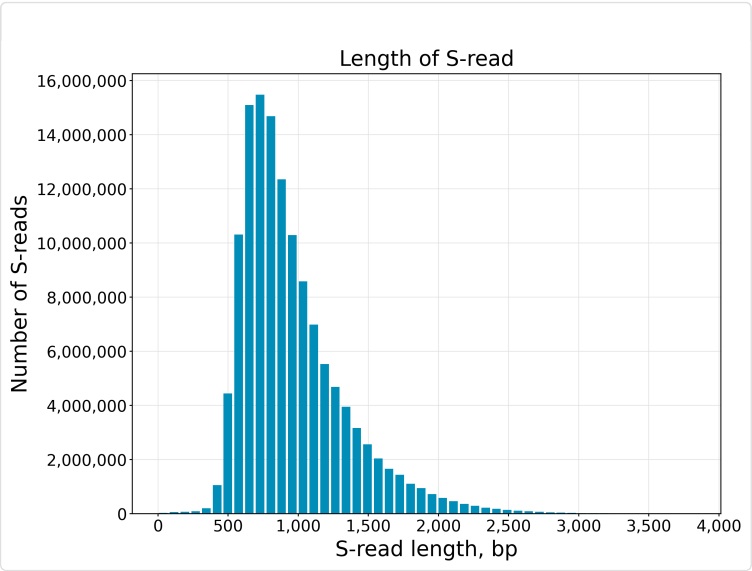


Length of Reads





S-read Length Distribution



File Downloads

Edit Output File Name Prefix

Example:analysis-SNAM-6-13175

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-...