

# TCC-MSA1003-Revio-ReadSeg - Bio Sample 1

SUCCESSFUL

Copy

Delete

▼ Analysis Overview
Status
Thumbnails
Display All
► Read Segmentation
► Data

## Display All

### Status



**Analysis ID** 109657  
**From Multi-Job** [109653](#)  
**Status** SUCCESSFUL: 4 tasks finished  
**Created By** etseng  
**Date Created** 2023-12-11, 03:00:36 PM  
**Date Updated** 2023-12-12, 08:55:04 AM  
**Application** Read Segmentation  
**SMRT Link Version** 13.0.0.207600

### Inputs

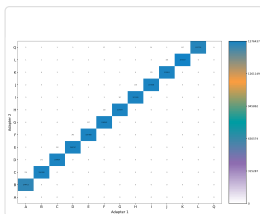
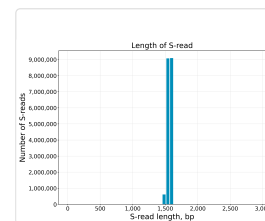
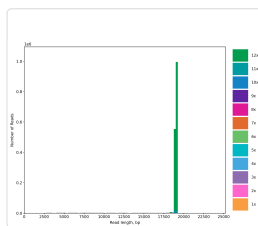
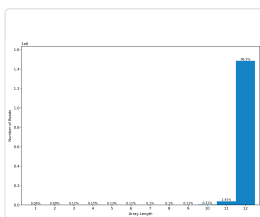
Data Type ↑↓	Name ↑↓
BarcodeSet	<a href="#">Barcode Sets: MAS-Seq Adapter v2 (MAS12)</a>
ConsensusReadSet	<a href="#">HiFi Reads: 20231020-4_84028_13.0VnV-MAS-Seq_PE8_24hPE...</a>

### Path

/pbi/flash/smrtlink/smrtlink-alpha/smrtlink/userdata/jobs\_root/0000/0000109/0000109657

## ► Analysis Parameters

### Thumbnails

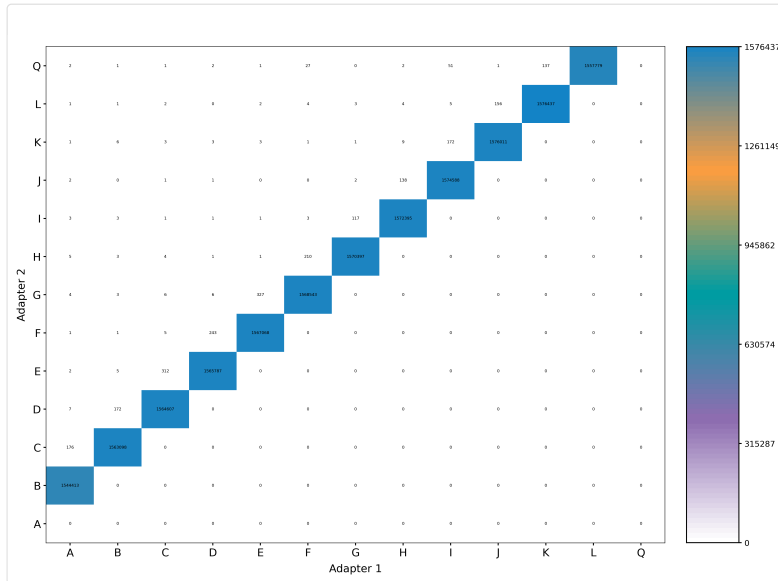
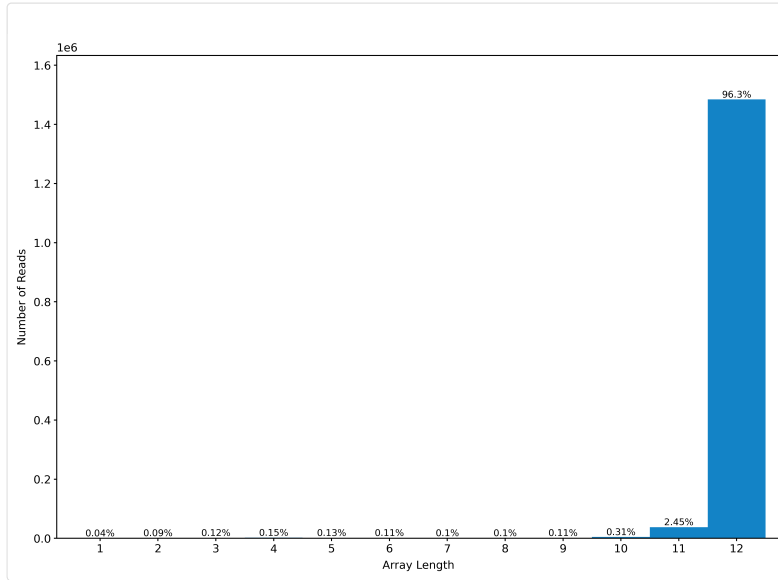


## Read Segmentation

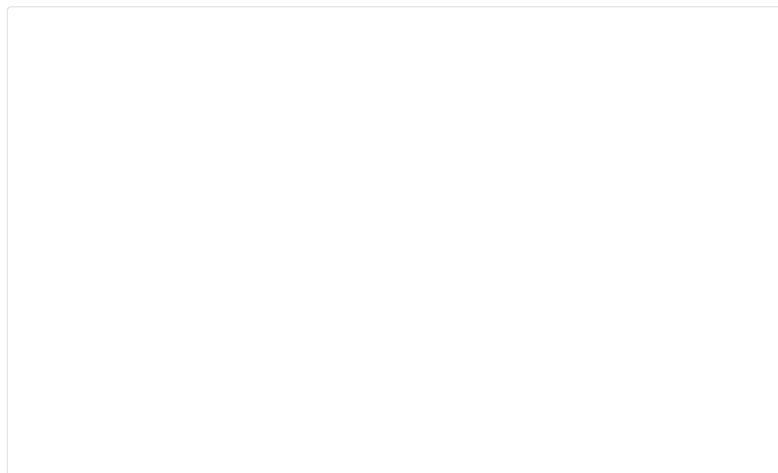
Value	Analysis Metric
1,579,179	Reads
18,801,123	Segmented reads (S-reads)
1,566	Mean length of S-reads

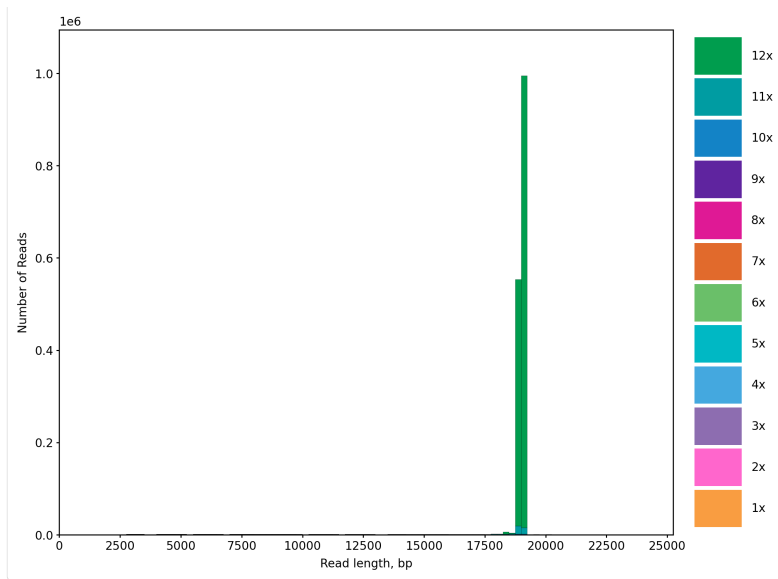
Value	Analysis Metric
96.35 %	Percent of reads with full arrays
11.91	Mean array size (concatenation factor)

## Segmentation Statistics

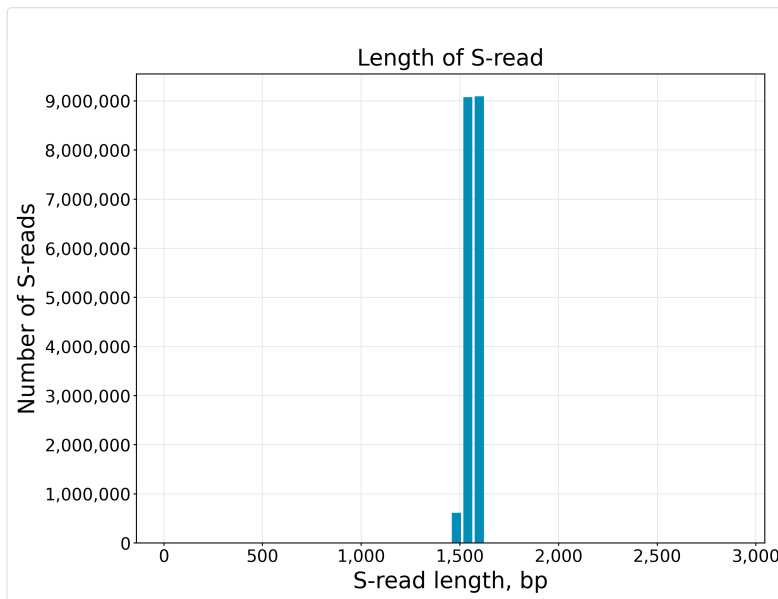


## Length of Reads





### S-read Length Distribution



### File Downloads

**Example:**analysis-Bio Sample 1-109657