



SUCCESSFUL

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## Display All

### Status



**Analysis ID** 109620  
**Status** SUCCESSFUL: 4 tasks finished  
**Created By** etseng  
**Date Created** 2023-12-11, 01:38:09 PM  
**Date Updated** 2023-12-11, 02:09:38 PM  
**Application** Read Segmentation  
**SMRT Link Version** 13.0.0.207600

### Inputs

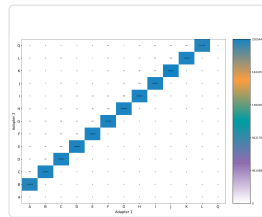
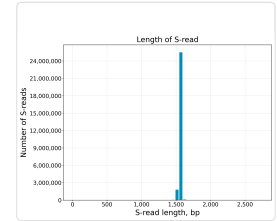
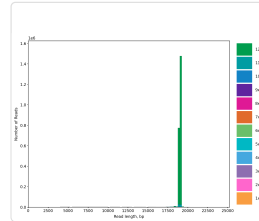
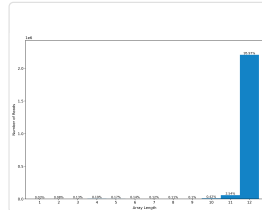
Data Type ↑↓	Name ↓↓
BarcodeSet	<a href="#">Barcode Sets: MAS-Seq Adapter v2 (MAS12)</a>
ConsensusReadSet	<a href="#">HiFi Reads: 16S_ATCC1003_DL_HB001 (demux1)_(Bio Sample 1)</a>

### Path

/pbi/flash/smartlink/smartlink-alpha/smartlink/userdata/jobs\_root/0000/0000109/0000109620

### ► Analysis Parameters

### Thumbnails

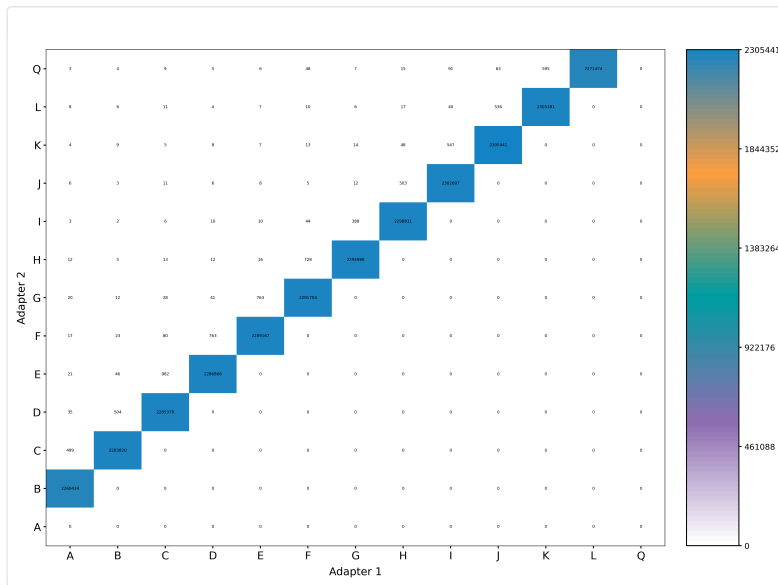
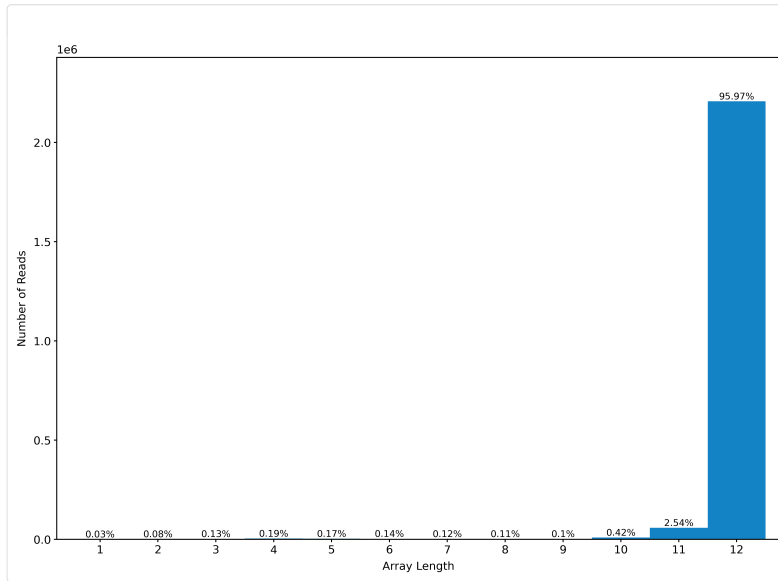


### Read Segmentation

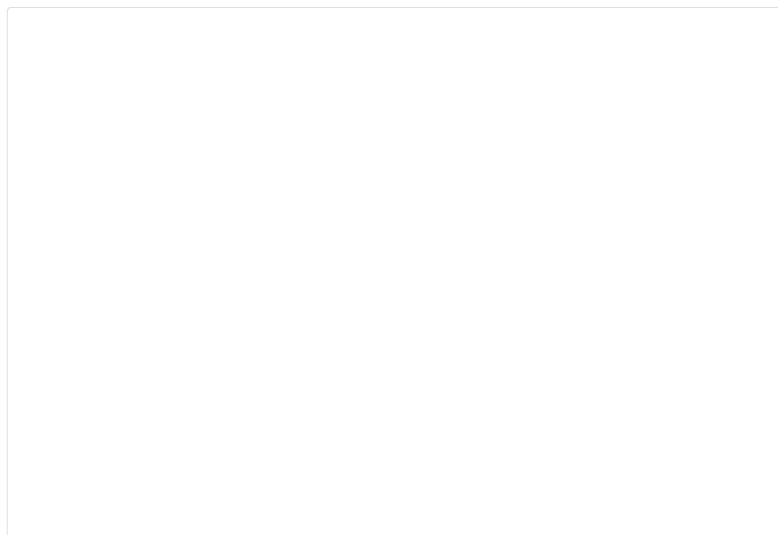
Value	Analysis Metric
2,310,268	Reads
27,476,179	Segmented reads (S-reads)
1,567	Mean length of S-reads
95.97 %	Percent of reads with full arrays

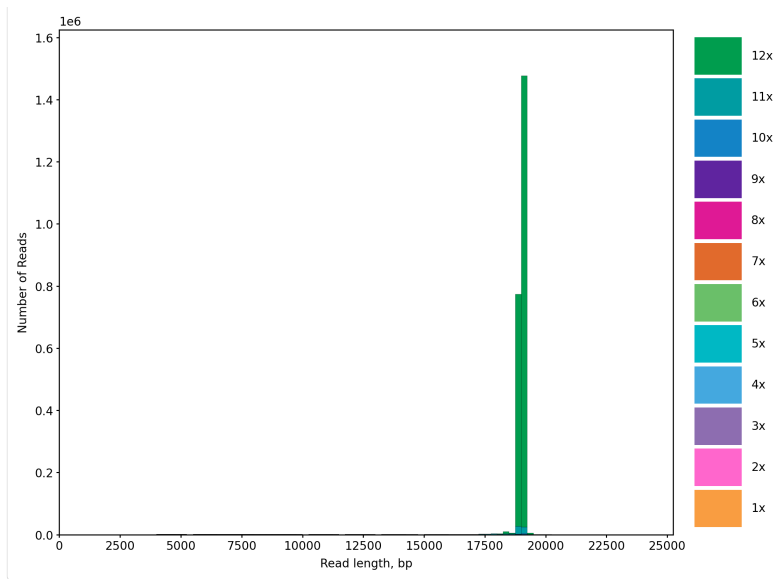
Value	Analysis Metric
11.89	Mean array size (concatenation factor)

## Segmentation Statistics

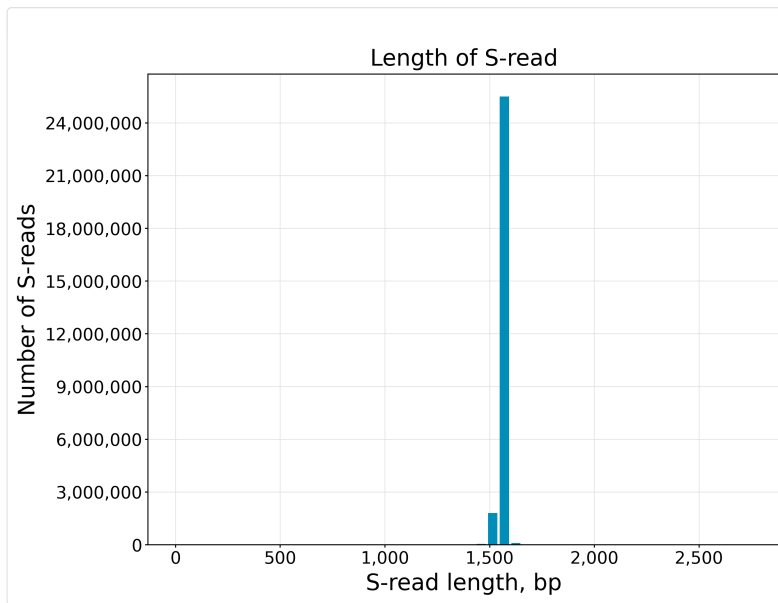


## Length of Reads





### S-read Length Distribution



### File Downloads

**Example:**analysis-Bio Sample 1-109620