

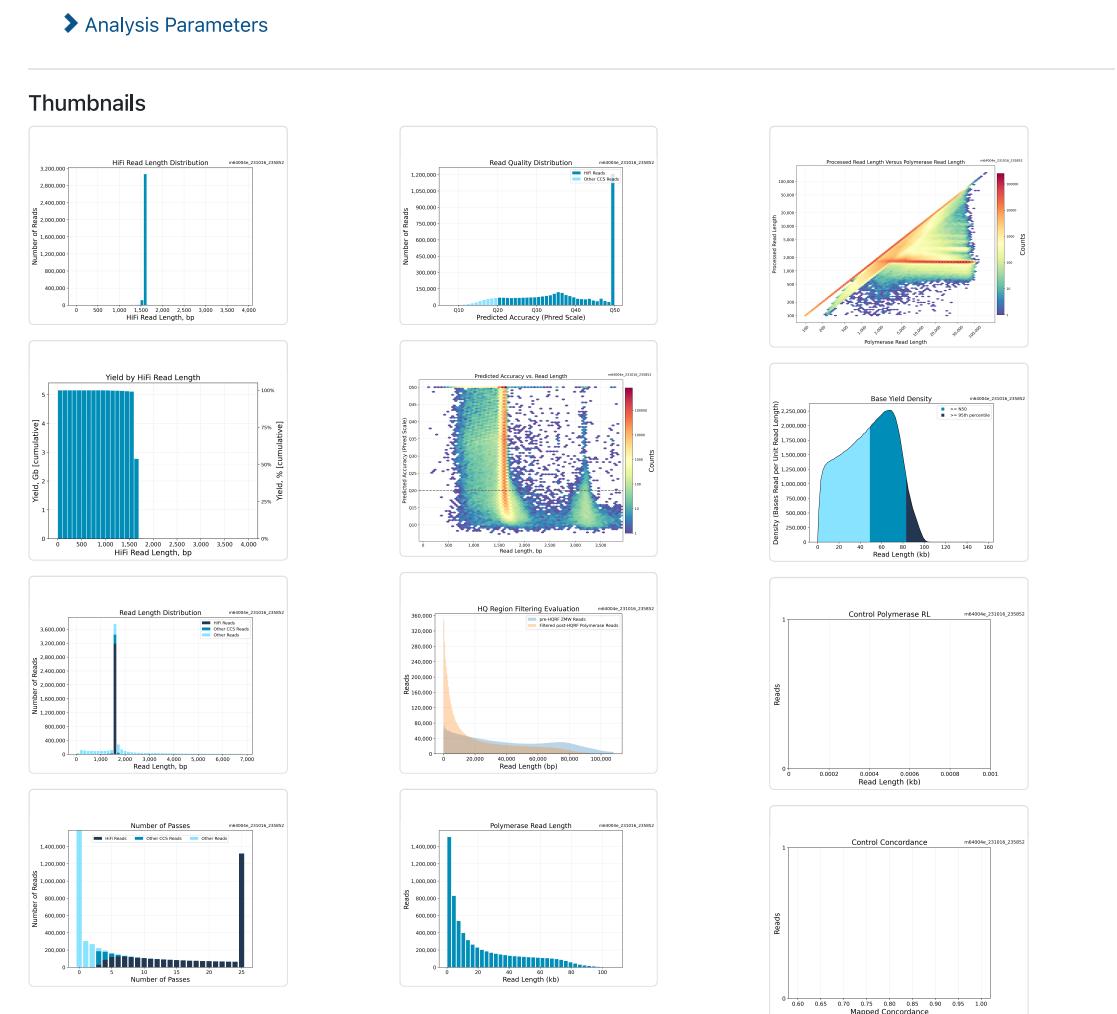
import-dataset

SUCCESSFUL

Delete

Analysis Overview	
Status	Thumbnail
Display All	
CCS Analysis Report	
Adapter Report	
Loading Report	
Raw Data Report	
Control Report	
Data	

Display All		
Status	Analysis	import-dataset
Analysis ID	109729	
Status	SUCCESSFUL: 1 task finished	
Created By	smrtlinktest	
Date Created	2023-12-13, 05:47:14 AM	
Date Updated	2023-12-13, 05:49:30 AM	
Application	Generate Dataset Reports	
SMRT Link Version	13.0.0.207600	
Inputs		
Data Type ↑	Name ↑	
		No Rows To Show
Path	/pbi/flash/smrtlink/smrtlink-alpha/smrtlink/userdata_root/0000/0000109/0000109729	



CCS Analysis Report

Value	Analysis Metric
3.2 M	HiFi reads
5.15 Gb	HiFi reads yield
1.59 kb	HiFi reads length (mean)
1,600	HiFi reads length (median, bp)
1,600	HiFi Read Length N50 (bp)
Q40	HiFi Read Quality (median)
18	HiFi Number of Passes (mean)

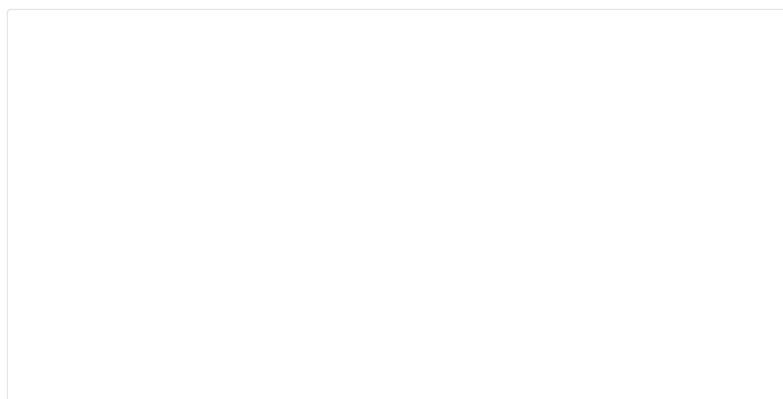
HiFi Read Length Summary

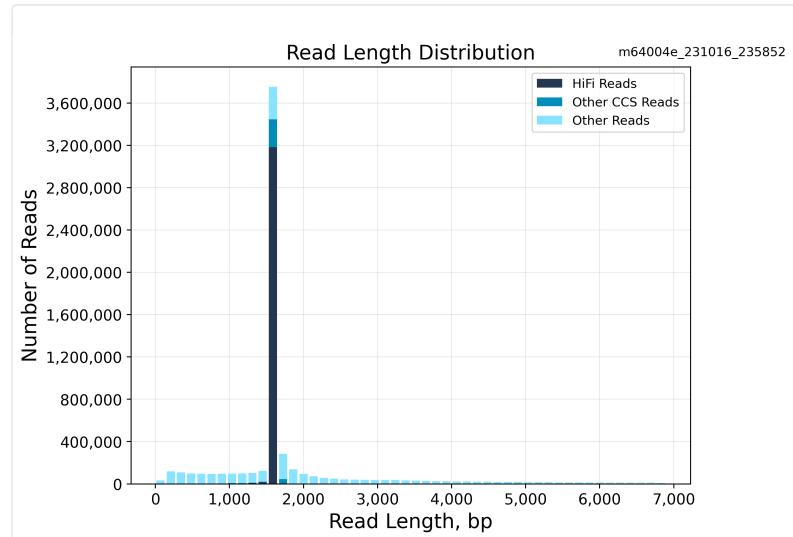
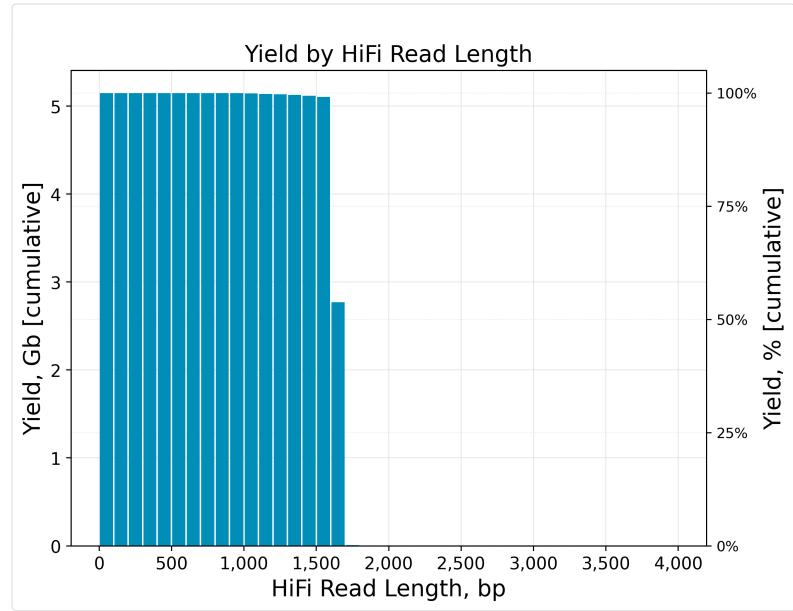
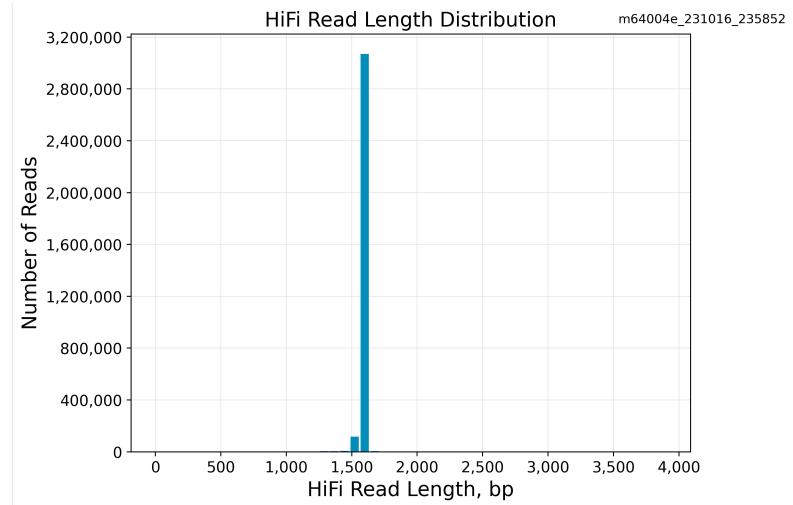
Search ...			
Read Length (kb) ↑	Reads ↓	Reads (%) ↓	Yield (Gb) ↑
≥ 0	3,228,520	100	5,149,944,597 Gb
≥ 5,000	227	0	2,286,313 Gb
≥ 10,000	187	0	2,041,350 Gb
≥ 15,000	0	0	0 Gb
≥ 20,000	0	0	0 Gb
≥ 25,000	0	0	0 Gb
≥ 30,000	0	0	0 Gb
≥ 35,000	0	0	0 Gb
≥ 40,000	0	0	0 Gb

HiFi Read Quality Summary

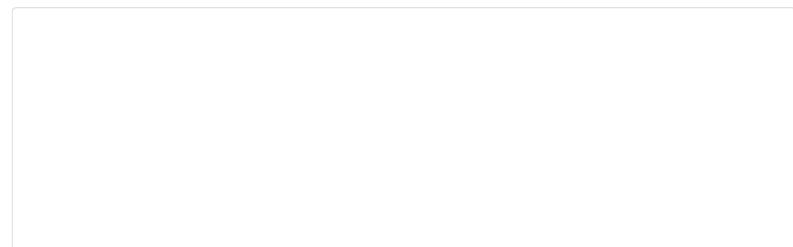
Search ...			
Read Quality (Phred) ↑	Reads ↓	Reads (%) ↓	Yield (Gb) ↑
≥ Q20	3,228,520	100	5.15 Gb
≥ Q30	2,547,767	79	4.06 Gb
≥ Q40	1,639,362	51	2.61 Gb
≥ Q50	1,176,174	36	1.87 Gb

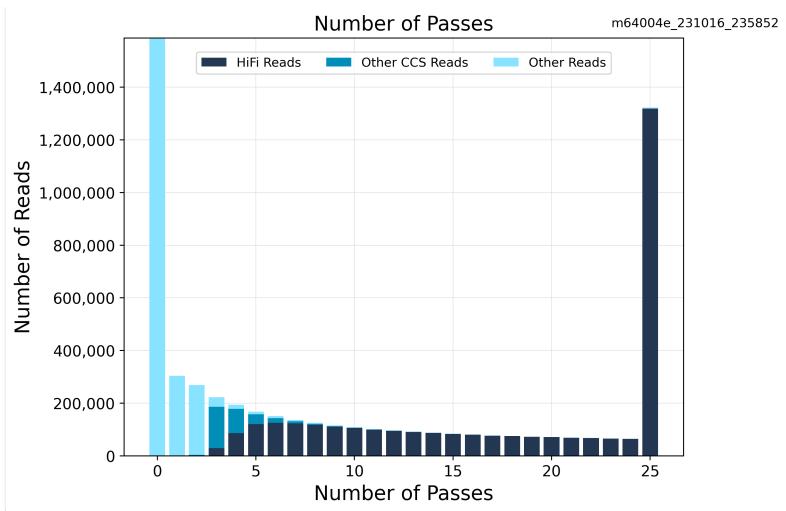
Read Length Distribution



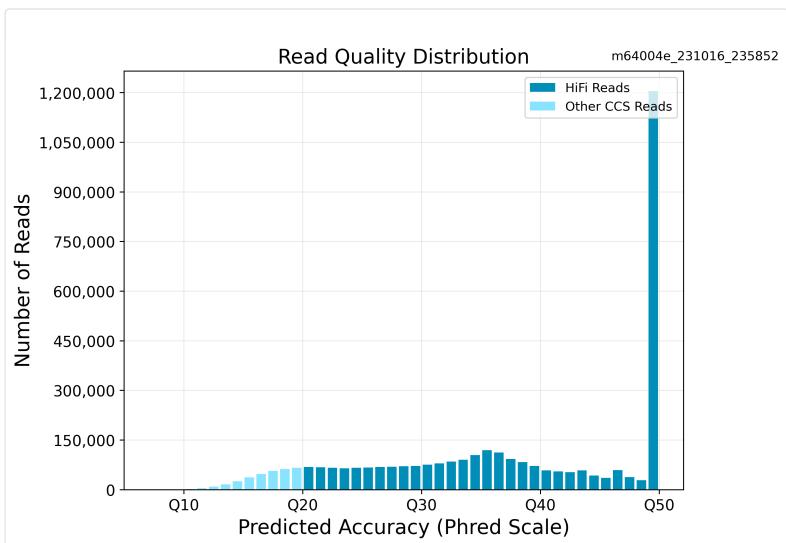


Number of Passes

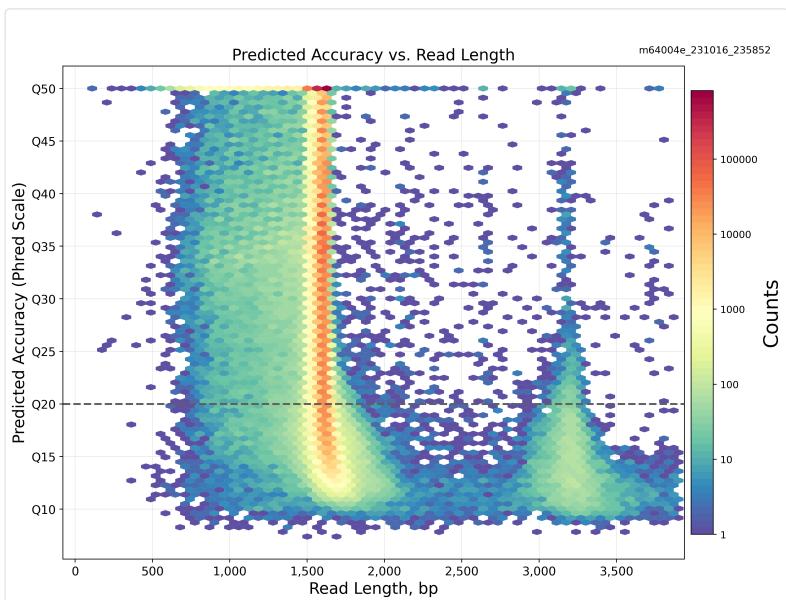




Read Quality Distribution



Predicted Accuracy vs. Read Length



Adapter Report

Value	Analysis Metric
0.00 %	Adapter Dimers (0-10bp) %
0.00 %	Short Inserts (11-100bp) %
2.11	Local Base Rate

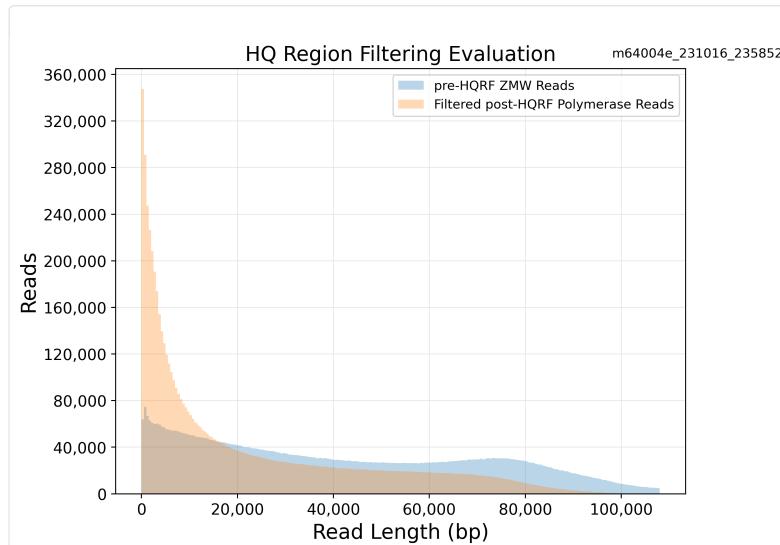
Loading Report

Value	Analysis Metric
8,014,671	Productive ZMWs
632,634	Productivity 0
6,676,707	Productivity 1
705,330	Productivity 2

Loading Statistics

Collection C...	Total ZMWs	Productivit...	P0 (%)	Productivit...	P1 (%)	Productivit...
m64004e_2...	8,014,671	632,634	7.89	6,676,707	83.31	705,330

HQ Region Filtering Evaluation

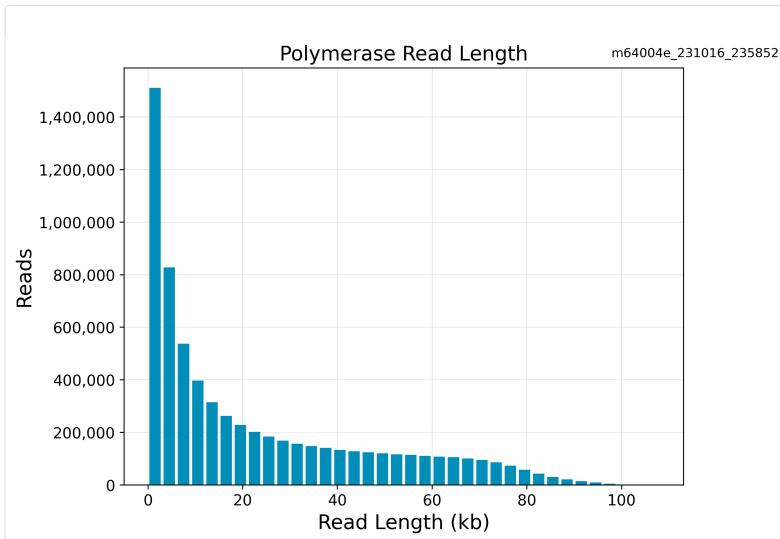


Raw Data Report

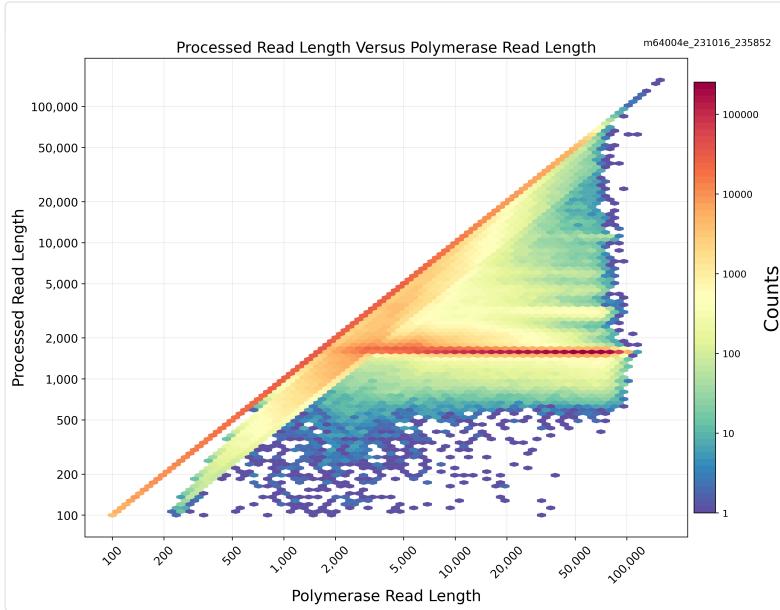
Value	Analysis Metric
150,484,296,061	Polymerase Read Bases
6,676,707	Polymerase Reads

Value	Analysis Metric
22.54 kb	Polymerase Read Length (mean)
49.75 kb	Polymerase read length (N50)
3.44 kb	Polymerase read length longest subread length (mean)
5.25 kb	Polymerase read length longest subread length (N50)
18,716,176,384	Unique Molecular Yield

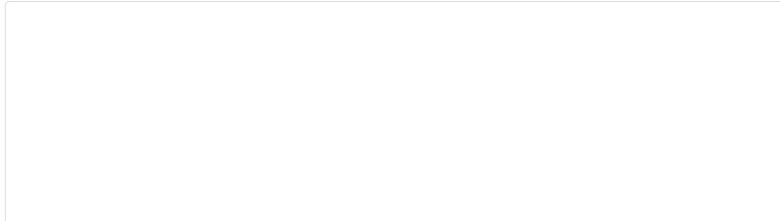
Polymerase Read Length

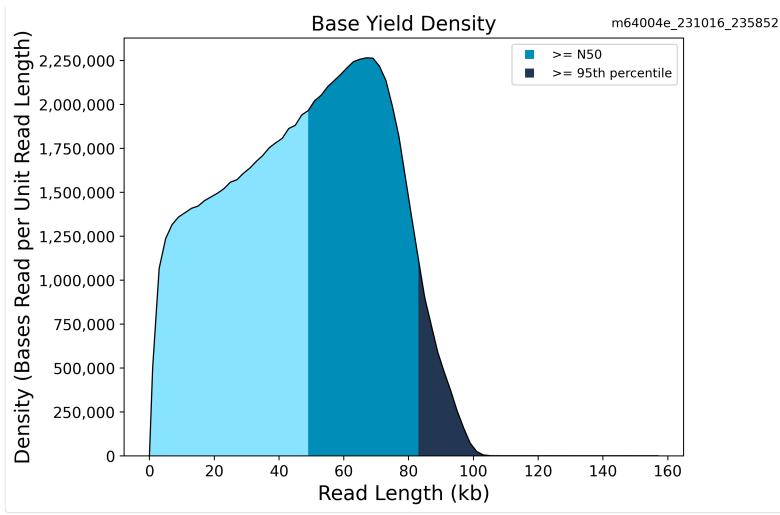


Longest Subread Length Versus Polymerase Read Length



Base Yield Density

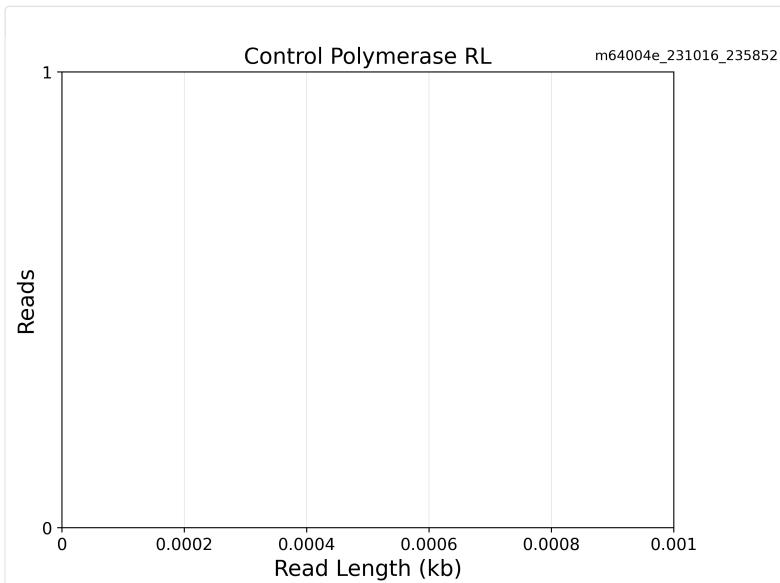




Control Report

Value	Analysis Metric
0	Number of Control Reads
NA	Control Read Length Mean
NA	Control Read Concordance Mean
NA	Control Read Concordance Mode

Control Read Length



Control Read Quality

