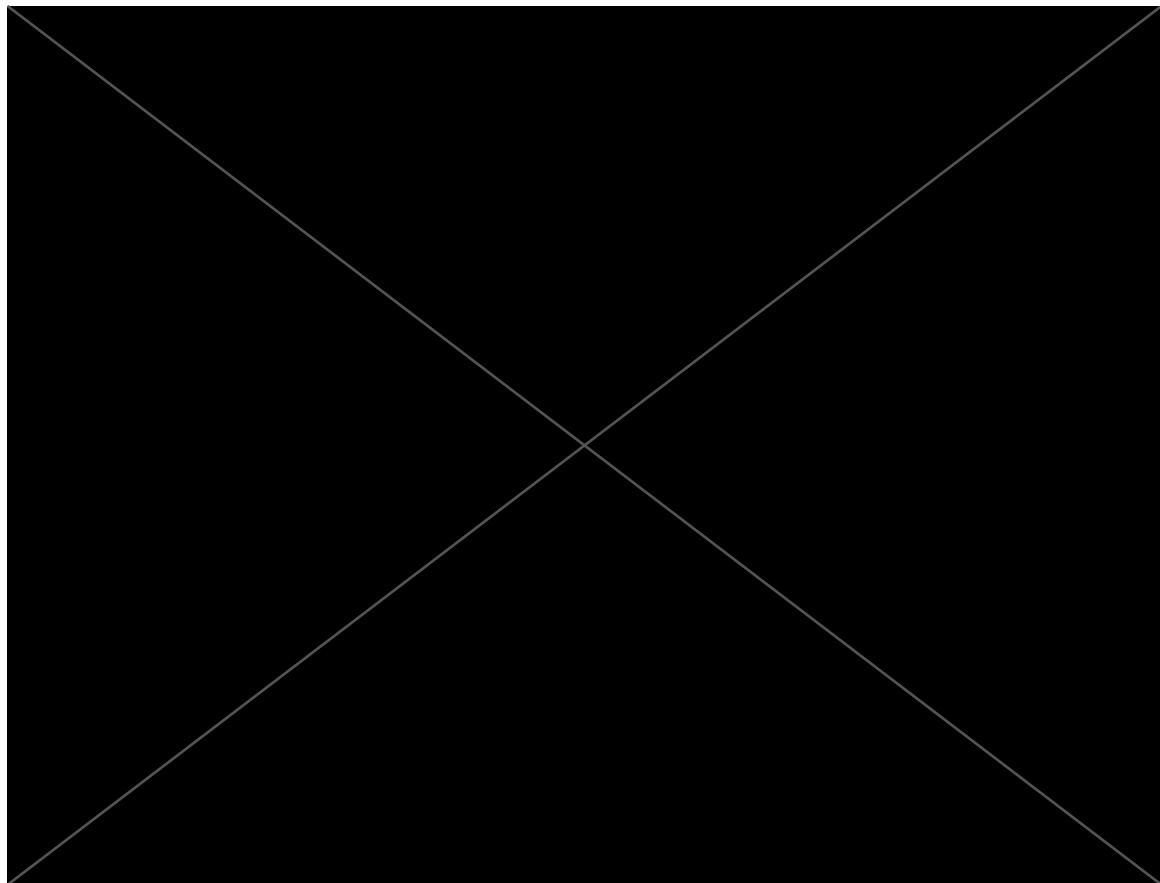


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<a href="#">▼Dataset Overview</a>
<a href="#">Status</a>
<a href="#">Thumbnails</a>
<a href="#">Display All</a>
<a href="#">►CCS Analysis Report</a>
<a href="#">►5mC CpG Report</a>
<a href="#">►Adapter Report</a>
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<a href="#">►Raw Data Report</a>
<a href="#">►Control Report</a>
<a href="#">►Barcodes</a>
<a href="#">►Analyses</a>
<a href="#">►Data</a>

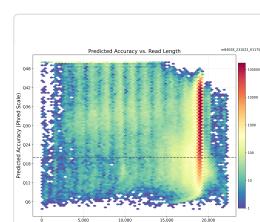
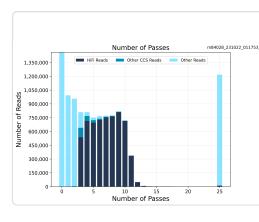
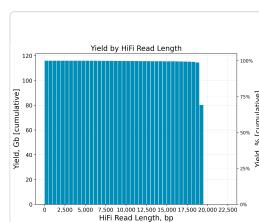
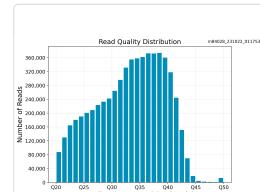
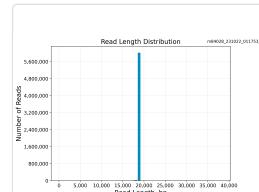
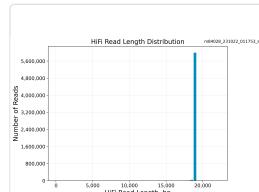
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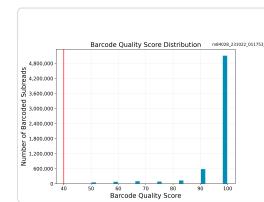
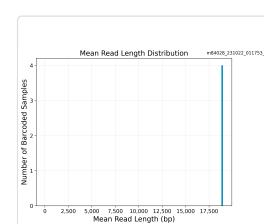
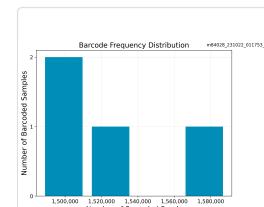
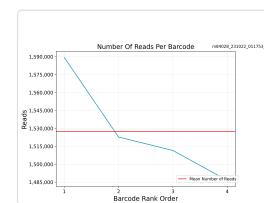
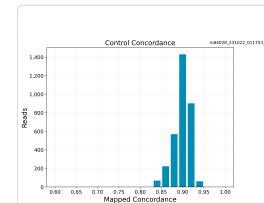
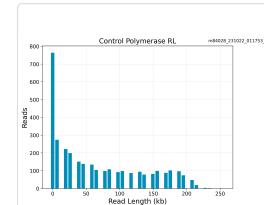
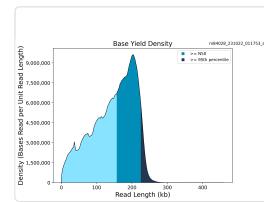
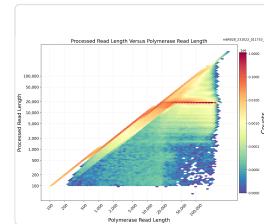
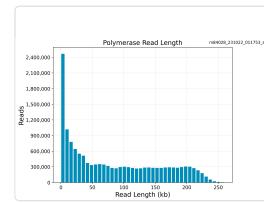
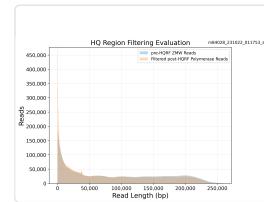
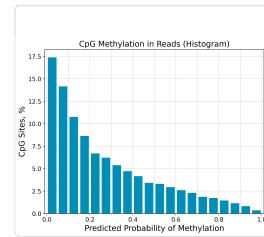
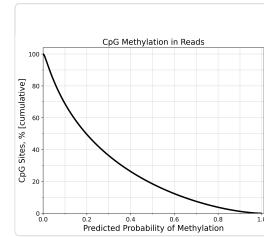


Movie Name m84028\_231022\_011753\_s3

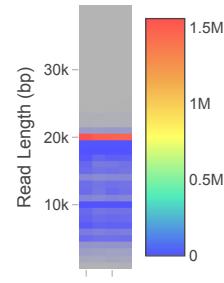
### ► Analysis Parameters

#### Thumbnails

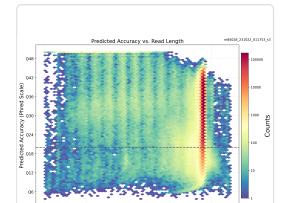
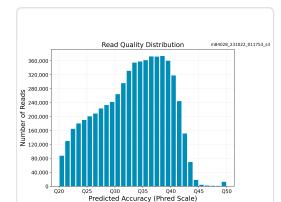
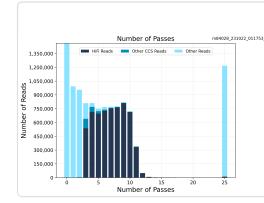
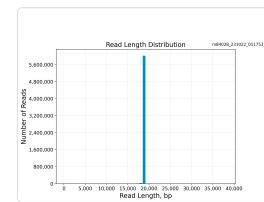
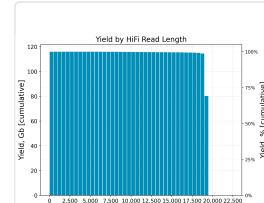
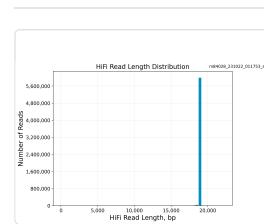
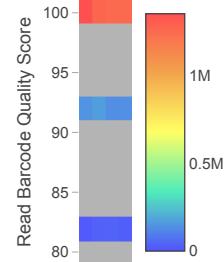


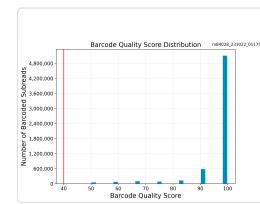
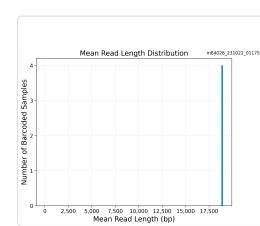
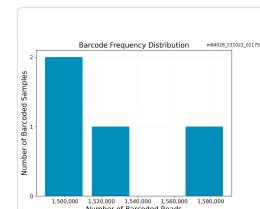
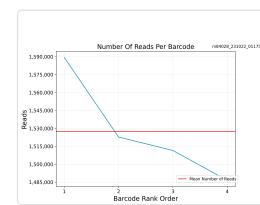
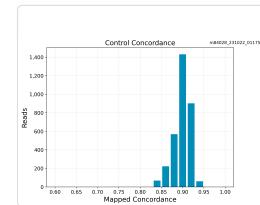
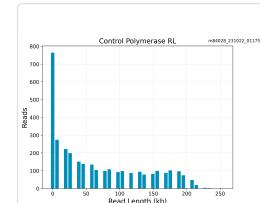
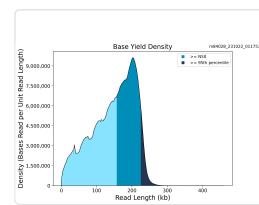
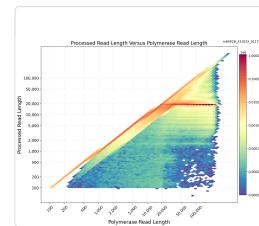
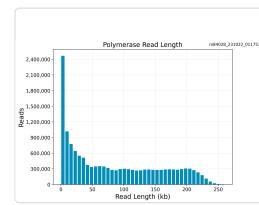
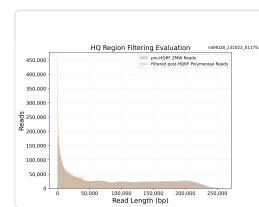
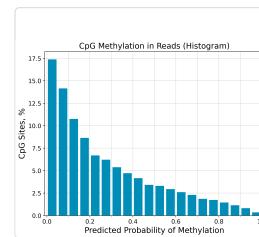
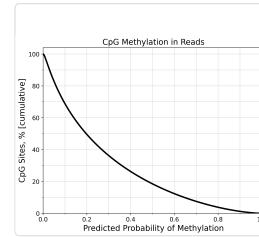


## Length Distribution By Barcode Rank Order

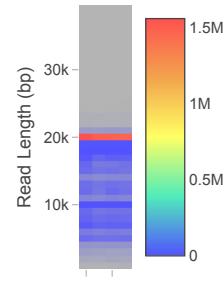


## Barcode Quality Distribution By Barcode Rank Order





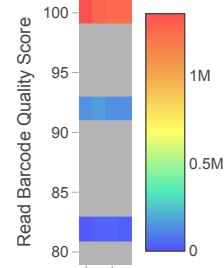
Length Distribution By Barcode Rank Order



bcdM0001-bcdM0001  
bcdM0002-bcdM0002

Barcode Rank Order By Read Count

Barcode Quality Distribution By Barcode Rank Order



bcdM0001-bcdM0001  
bcdM0002-bcdM0002

Barcode Rank Order By Read Count

## CCS Analysis Report

Value	Analysis Metric
6.1 M	HiFi reads
115.86 Gb	HiFi reads yield
18.91 kb	HiFi reads length (mean)
19,022	HiFi reads length (median, bp)
19,023	HiFi Read Length N50 (bp)
Q34	HiFi Read Quality (median)
93.58%	Base Quality $\geq$ Q30 (%)

Value	Analysis Metric
7	HiFi Number of Passes (mean)

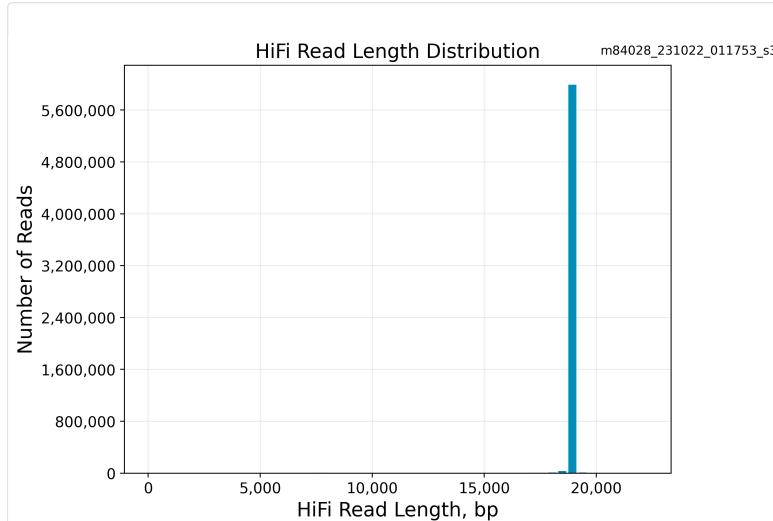
## HiFi Read Length Summary

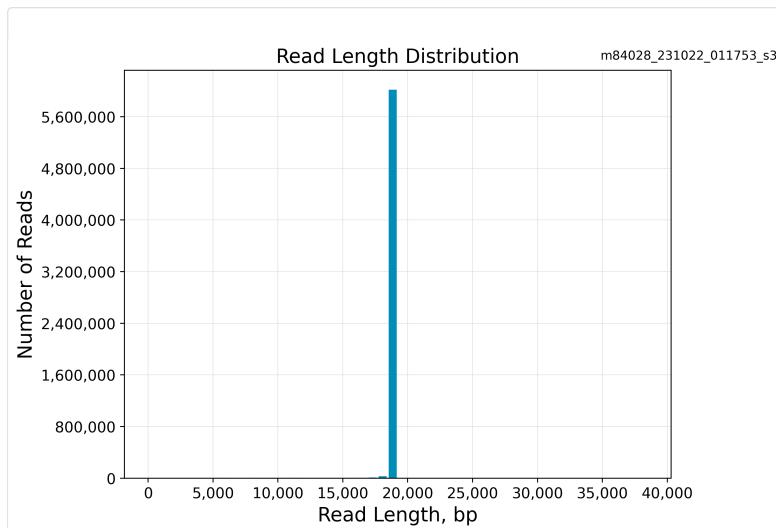
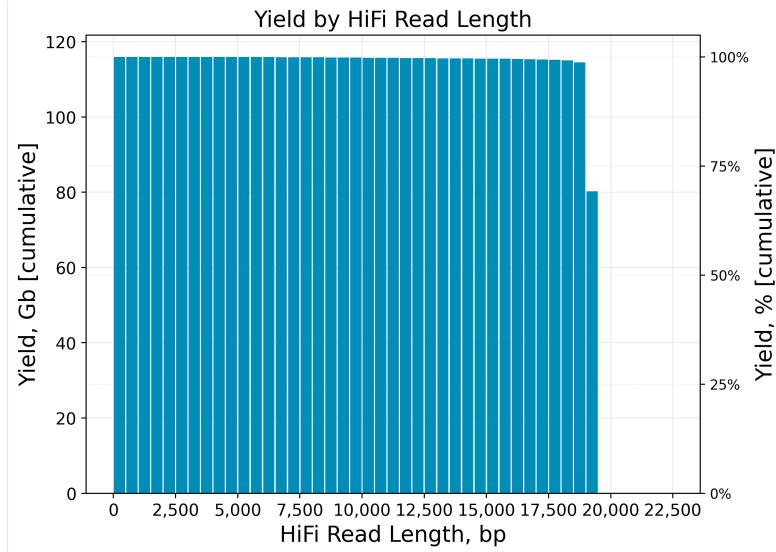
Read Length (kb) ↑	Reads ↓	Reads (%) ↓	Yield (Gb) ↑	Yield
≥ 0	6,127,402	100	115,855,626,177 Gb	100
≥ 5,000	6,117,614	100	115,819,966,821 Gb	100
≥ 10,000	6,091,975	99	115,623,620,586 Gb	100
≥ 15,000	6,072,156	99	115,376,651,115 Gb	100
≥ 20,000	1,442	0	30,479,508 Gb	0
≥ 25,000	53	0	1,608,988 Gb	0
≥ 30,000	24	0	829,210 Gb	0
≥ 35,000	11	0	403,983 Gb	0
≥ 40,000	0	0	0 Gb	0

## HiFi Read Quality Summary

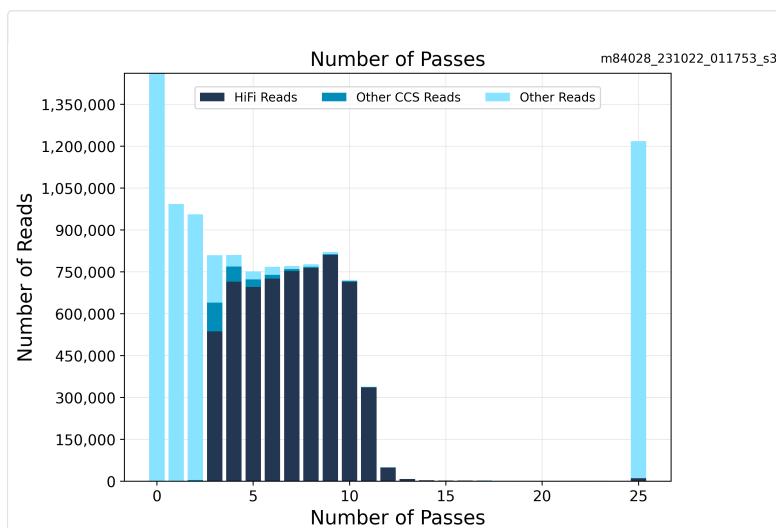
Read Quality (Phred) ↑	Reads ↓	Reads (%) ↓	Yield (Gb) ↑	Yield
≥ Q20	6,127,402	100	115.86 Gb	100
≥ Q30	4,267,968	70	80.68 Gb	70
≥ Q40	830,588	14	15.52 Gb	13
≥ Q50	11,685	0	0.07 Gb	0

## Read Length Distribution

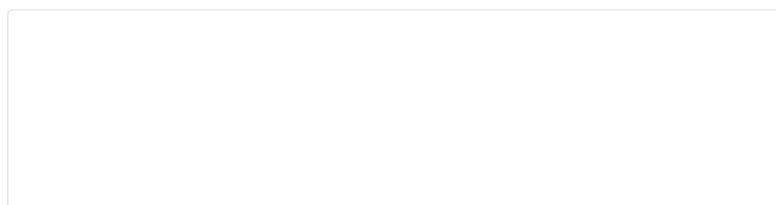


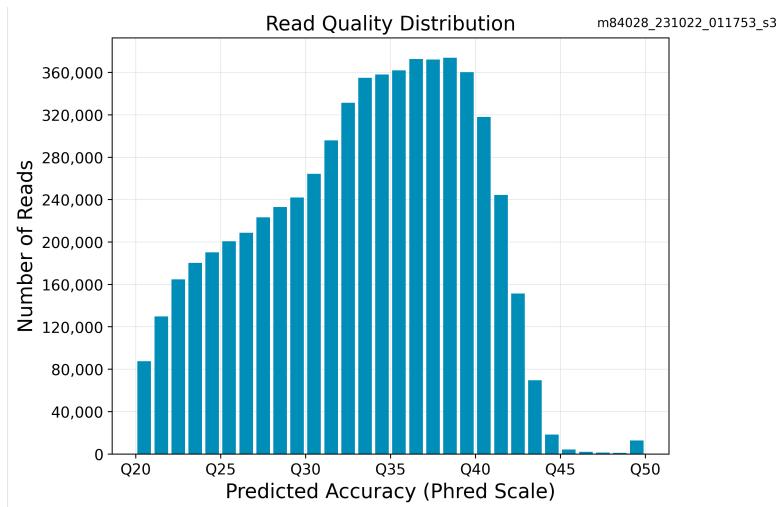


## Number of Passes

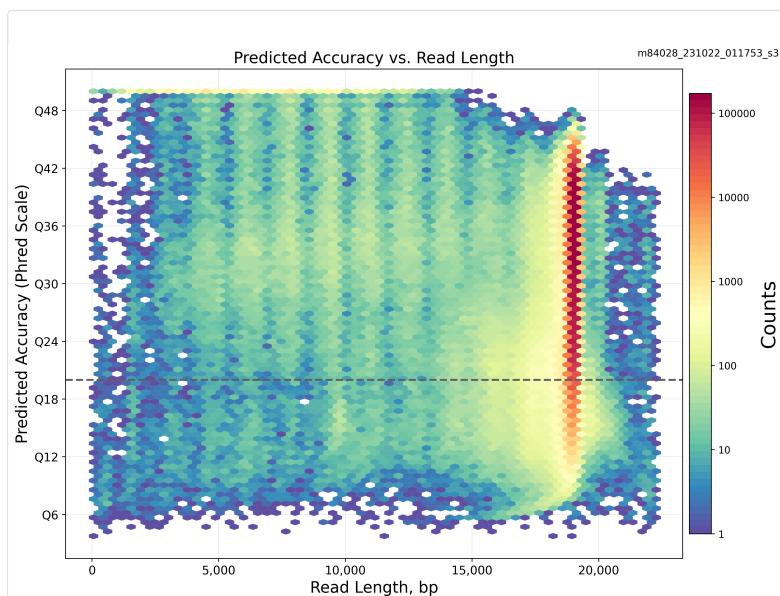


## Read Quality Distribution

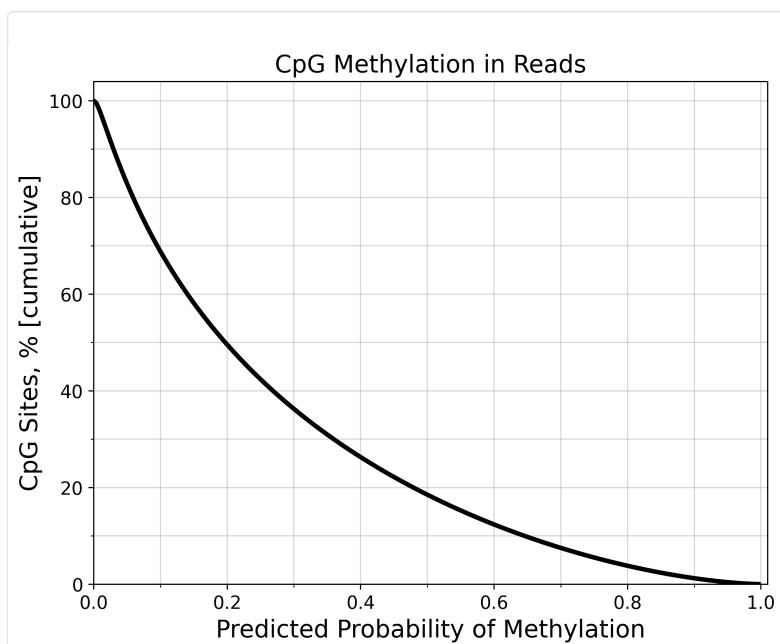


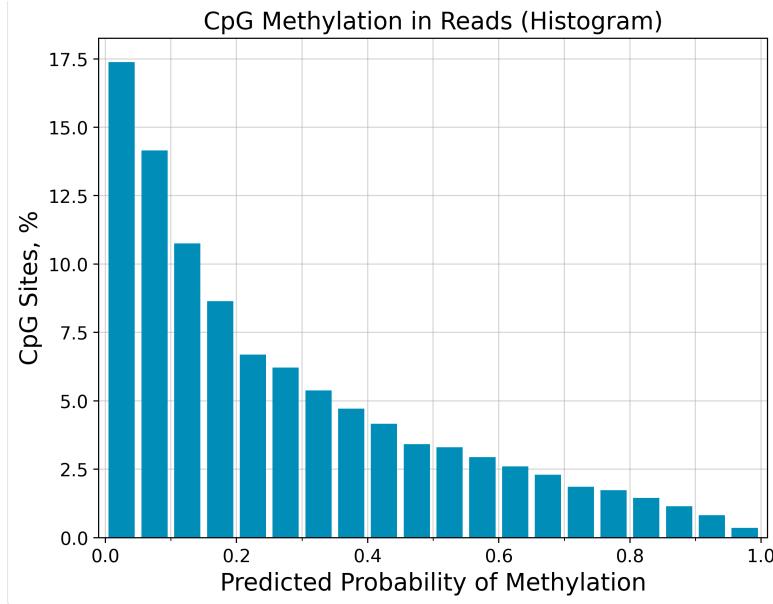


### Predicted Accuracy vs. Read Length



### Methylation Probability





## Adapter Report

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

## Loading Report

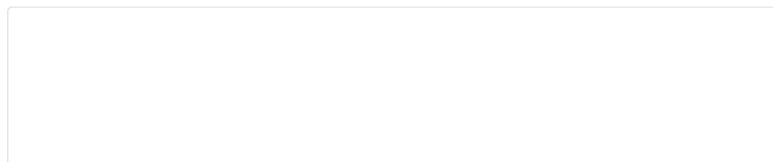
Value	Analysis Metric
25,165,824	Productive ZMWs
10,396,301	Productivity 0
14,678,327	Productivity 1
91,196	Productivity 2

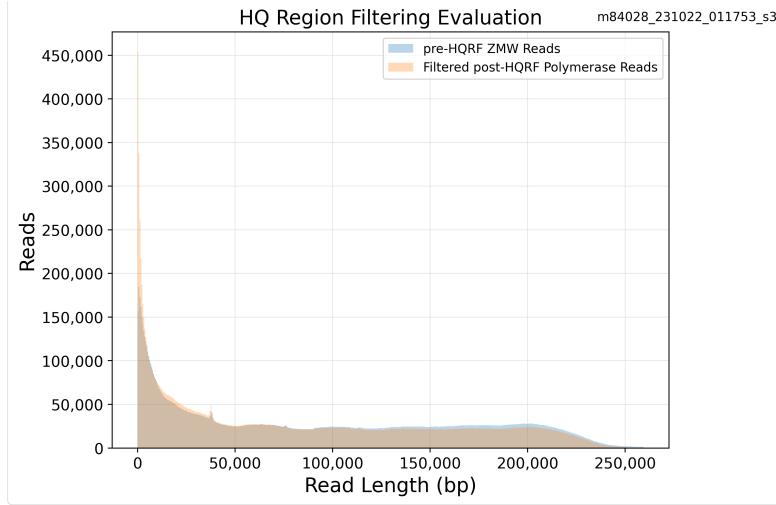
## Loading Statistics

Search ...

Collection	Total ZMWs	Productivity 0	P0 (%)	Productivity 1	P1 (%)	Productivity 2	P2 (%)
m84028_...	25,165,824	10,396,301	41.31	14,678,327	58.33	91,196	0.36

## HQ Region Filtering Evaluation

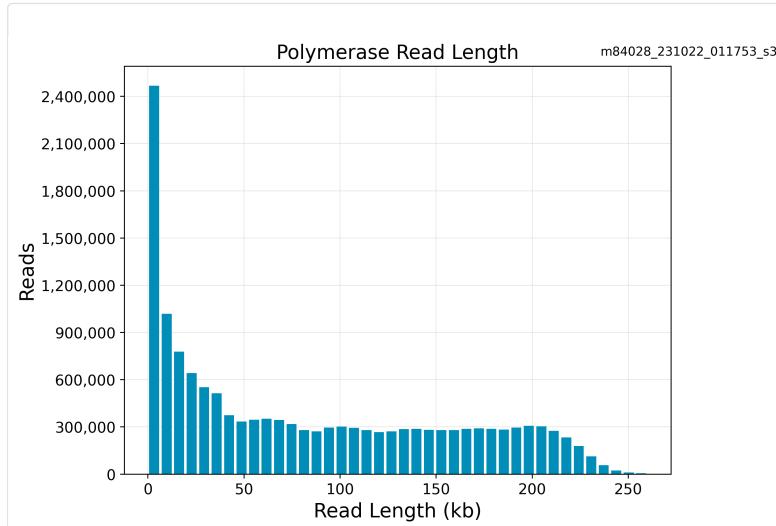




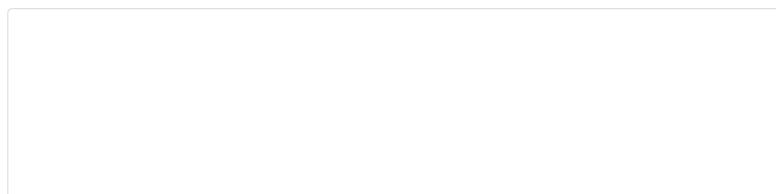
## Raw Data Report

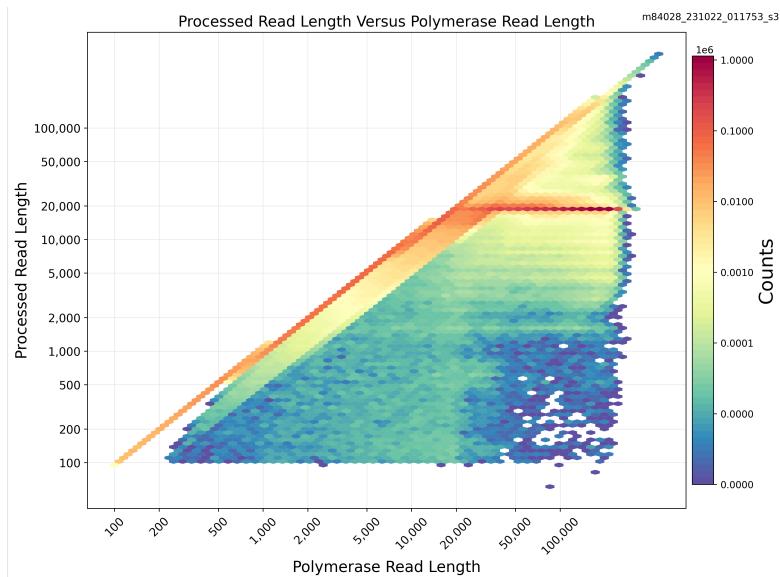
Value	Analysis Metric
1,190,965,740,710	Polymerase Read Bases
14,675,072	Polymerase Reads
81.16 kb	Polymerase Read Length (mean)
158.25 kb	Polymerase read length (N50)
17.20 kb	Polymerase read length longest subread length (mean)
19.75 kb	Polymerase read length longest subread length (N50)
238,684,094,464	Unique Molecular Yield

## Polymerase Read Length

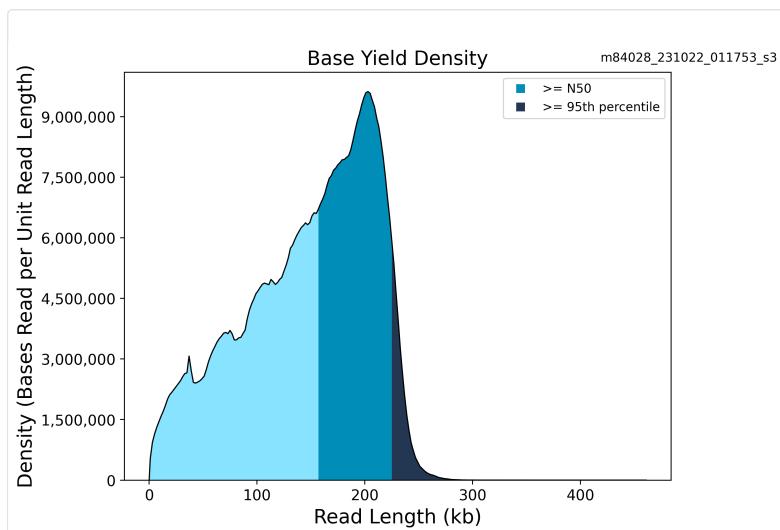


## Longest Subread Length Versus Polymerase Read Length





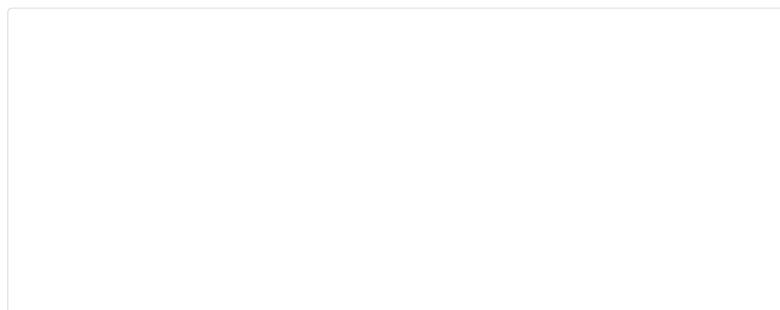
## Base Yield Density

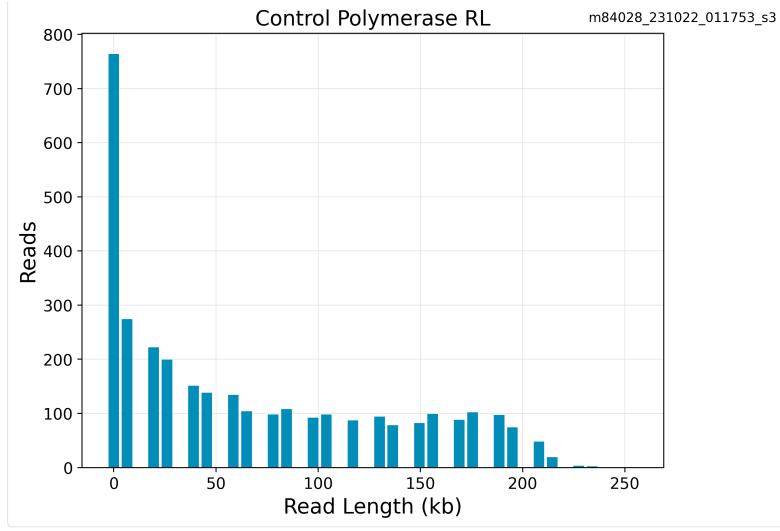


## Control Report

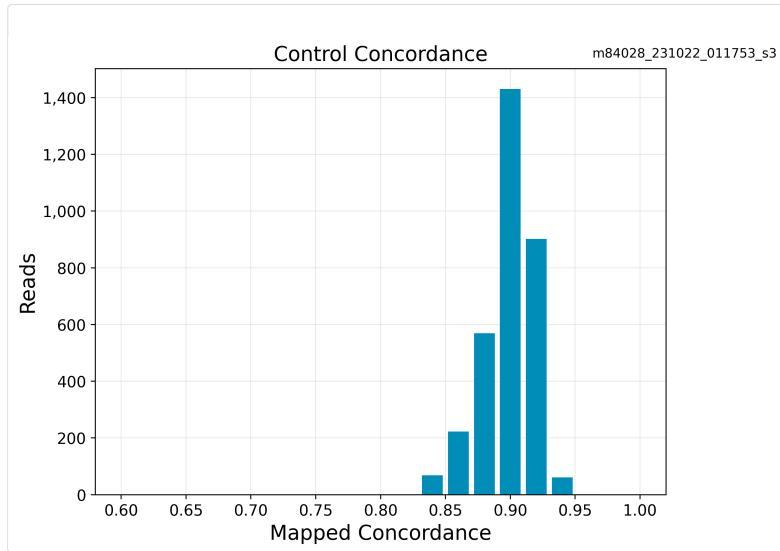
Value	Analysis Metric
3,255	Number of Control Reads
68,795	Control Read Length Mean
0.91	Control Read Concordance Mean
0.91	Control Read Concordance Mode

## Control Read Length





## Control Read Quality



## Barcodes

Value	Analysis Metric
4	Unique Barcodes
6,109,883	Barcoded HiFi Reads
17,519	Unbarcoded HiFi Reads
99.71%	Barcoded HiFi Reads (%)
115.55 Gb	Barcoded HiFi yield (Gb)
0.31 Gb	Unbarcoded HiFi yield (Gb)
99.73%	Barcoded HiFi Yield (%)
1,527,470	Mean HiFi Reads per Barcode
1,589,279	Max. HiFi Reads per Barcode
1,486,455	Min. HiFi Reads per Barcode

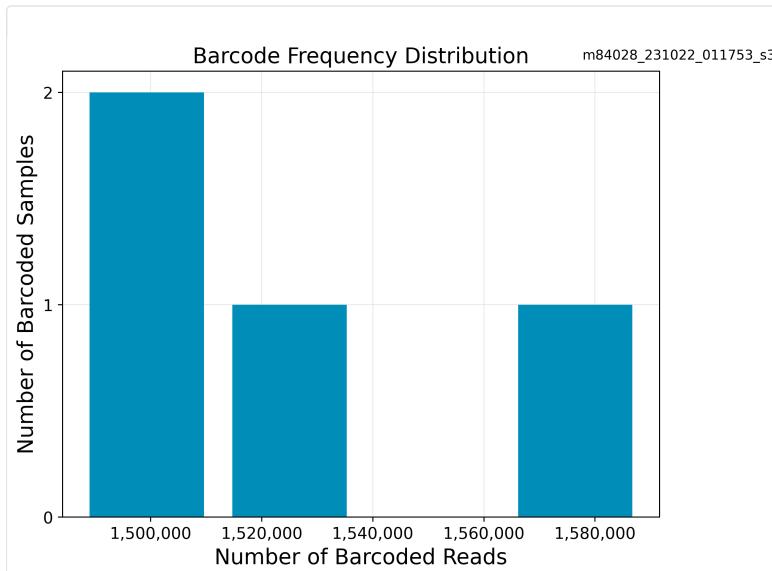
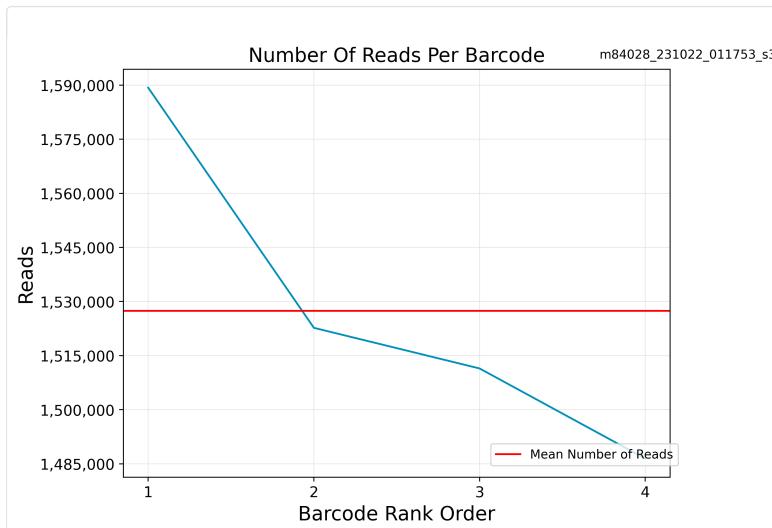
Value	Analysis Metric
18.91 kb	Barcoded HiFi read length (mean, kb)
17.46 kb	Unbarcoded HiFi read length (mean, kb)

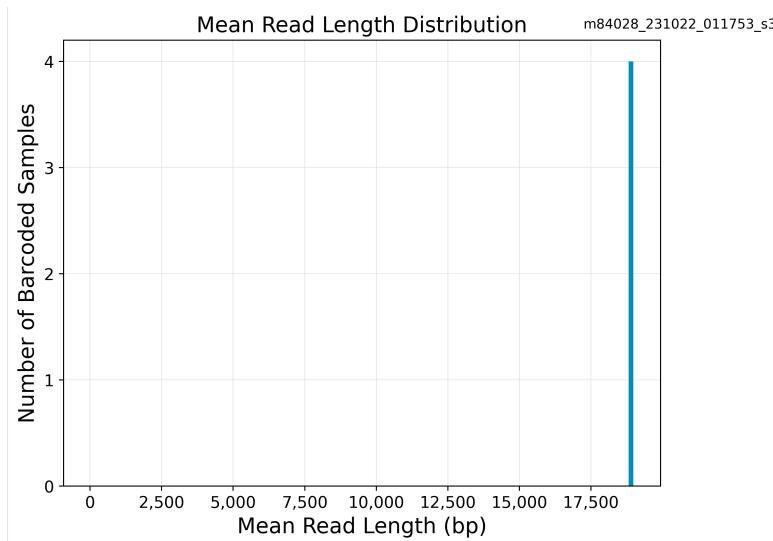
## Barcode Data

Search ...

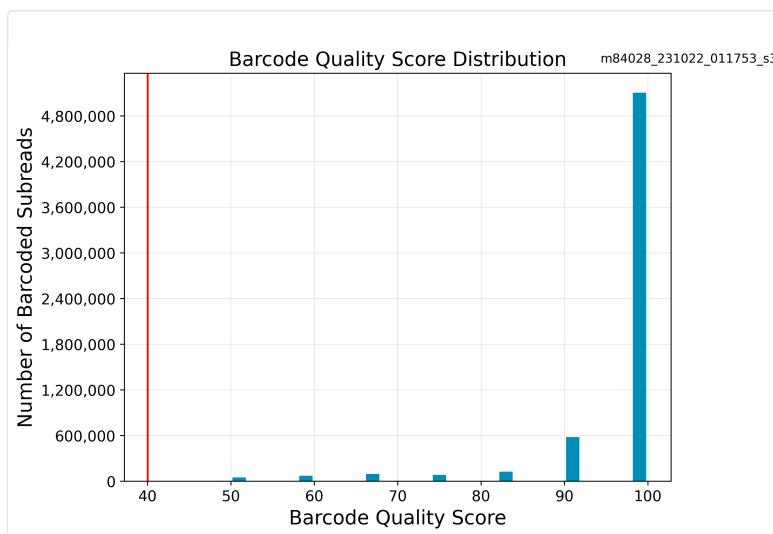
Sample N...	Barcode	Barcode Q...	HiFi Reads	HiFi Read ...	HiFi Read ...	HiFi Yield (..
MAS-seq_...	bcM0001--...	97.3	1,589,279	18,910	Q34	30,054,580
MAS-seq_...	bcM0002--...	96.6	1,511,444	18,910	Q34	28,581,526,
MAS-seq_...	bcM0003--...	97.0	1,486,455	18,909	Q34	28,108,847,
MAS-seq_...	bcM0004--...	97.2	1,522,705	18,916	Q34	28,804,726,
No Name	Not Barcod...	0.0	17,519	17,463	Q27	305,944,98

## Barcoded Read Statistics

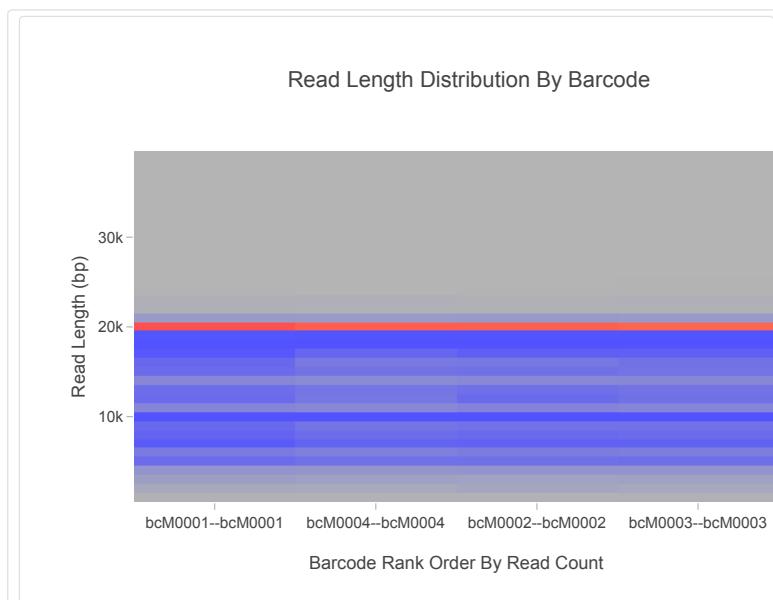




## Barcode Quality Scores



## Barcoded Read Binned Histograms



### Barcode Quality Distribution By Barcode

