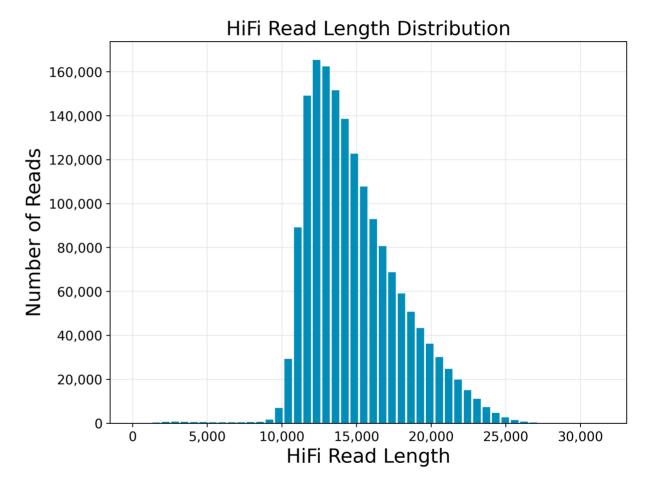
CCS Analysis Report

Value	Analysis Metric
1,682,414	HiFi Reads
25,038,087,630	HiFi Yield (bp)
14,882	HiFi Read Length (mean, bp)
Q34	HiFi Read Quality (median)
12	HiFi Number of Passes (mean)
273,468	<q20 reads<="" td=""></q20>
4,337,584,600	<q20 (bp)<="" td="" yield=""></q20>
15,861	<q20 (mean,="" bp)<="" length="" read="" td=""></q20>
Q16	<q20 (median)<="" quality="" read="" td=""></q20>



Raw Data Report

Value	Analysis Metric
411,009,987,641	Polymerase Read Bases
4,154,317	Polymerase Reads
98,935	Polymerase Read Length (mean)
215,103	Polymerase Read N50
14,006	Subread Length (mean)
14,902	Subread N50
18,623	Longest Subread Length (mean)
20,342	Longest Subread N50
71,037,124,608	Unique Molecular Yield

