

	Nanopore Sequencing (this study)	Nanopore Sequencing (combined)	200M Genome Assembly	Post Control Hi-C Quality and Scaffold-ing
Number of Sequences/Contigs	633,844	1,043,600	1,098	302
Sum of Sequence/Contig Length	6,514,819,227	10,462,257,773	268,538,400	198,798,533
Average Sequence/Contig Length	10,278	10,025	244,570.49	658,273
Median Sequence Length	7,405	6,549	60,582.50	24,380
Longest Sequence Length	123,032	238,837	7,592,683	37,891,417
N50	–	–	1,274,151	34,538,105