

**A**

Statistics	Number
Short reads (Illumina)	28,672,142
Long reads (MinION)	409,847
Total base pairs (Illumina)	3,784,150,188
Total base pairs (MinION)	772,618,969
Coverage (Illumina)	114 X
Coverage (MinION)	23 X
Genome length	32,966,471 bp
Number of genes	7,720
% of genome covered by genes	61.2
GC%	47.73
Transposable elements (TEs)	206
% of genome covered by TEs	16.16
tRNAs	82

**B**

### BUSCO Assessment Results

