

Supplementary File 3. Final summary assembly statistics for chromosome-scale assembly.

Scaffold total	1,034
Contig total	7,930
Scaffold sequence total	306.5 Mb
Chromosome sequence	282.6 Mb
Contig sequence total	291.7 Mb (4.8% gap)
Scaffold N/L50	4/43.6 Mb
Contig N/L50	797/110.9 Kb
