

*de novo* genome assembly (raw assembly from Velvet)

	number of scaffolds	min. length (bp)	N50 (bp)	N20 (bp)	max. length (bp)	sum (Mbp)
<i>SC</i>	358,309	200	4,522	59,233	965,826	508.82
<i>XT</i>	816,724	200	852	6,379	1,572,474	492.59

*de novo* genome assembly (final version)

	number of scaffolds	min. length (bp)	N50 (bp)	N20 (bp)	max. length (bp)	sum (Mbp)
<i>SC</i>	97,497	800	10,236	83,618	965,826	418.92
<i>XT</i>	97,640	800	4,078	29,235	1,572,474	257.96

*de novo* transcriptome assembly

	number of contigs	min. length (bp)	N50 (bp)	N20 (bp)	max. length (bp)	sum (Mbp)
<i>SC</i>	326,400	200	444	800	48,514	134.38
<i>XT</i>	346,768	200	491	958	48,514	151.8

annotated transcripts from the gene models

	number of contigs	min. length (bp)	N50 (bp)	N20 (bp)	max. length (bp)	sum (Mbp)
<i>SC</i> (eukaryotic)	26,967	63	1,176	2,184	18,322	25.29
<i>SC</i> (prokaryotic)	1,385	141	1,236	2,040	6,708	1.45
<i>SC</i> (unknown)	9,343	61	714	1,278	6,915	5.67
<i>XT</i> (eukaryotic)	22,337	93	1,312	2,427	21,282	24.09
<i>XT</i> (prokaryotic)	10,879	117	1,200	2,055	8,798	11.24
<i>XT</i> (unknown)	7,082	45	1,043	1,908	11,070	5.85