

Table S2. Statistics of *de novo* transcriptome assembly.

Assembly	Total number	Base-pairs(Mbp)	N50(bp)	Mean(bp)	Max(bp)	Min(bp)
H6	108,450	137.9	1,790	1,271.2	16,429	200
SP3B	94,051	114.0	1,695	1,211.8	16,482	200
Combined	165,179	203.0	1,756	1,228.9	16,504	200