

Table S3 The summary of assembly from transcriptome sequencing data in *A.squarrosom*.

<b>Length range (bp)</b>	<b>Transcripts</b>	<b>Unigenes</b>
200-500	88,664	81,733
500-1000	37,905	30,593
1000-2000	23,369	15,527
>2000	15,782	7,833
Total number	165,720	135,686
Total length	137,635,212	93,674,801
Mean length	830.53	690.38
N50 length	1349	933
GC%	39.85	39.73