

Table S2. Length distribution of assembly

Transcript length interval	200-500bp	500-1kbp	1k-2kbp	>2kbp	Total
Number of transcripts	65364	20873	15818	15814	117869
Number of unigenes	52931	12775	8104	7657	81467

Table S3. Assembly statistics of reads

Item	Min Length	Mean Length	Median Length	Max Length	N50	N90	Total Nucleotides
Transcripts	201	968	430	29472	2004	332	114057975
Unigenes	201	786	352	29472	1636	276	64048359