**Table S1. Result of the *de novo* transcriptome assembly**

|  |  |
| --- | --- |
| Type | Resource |
| Total transcripts number | 180,769 |
| Total unigenes number | 126,979 |
| Total sequence base | 146,455,511 |
| Largest (bp) | 13,536 |
| Smallest (bp) | 201 |
| Average length (bp) | 810.18 |
| N50 (bp) | 1,176 |
| GC percent (%) | 42.55 |
| Mean mapped reads | 839.46 |