

Supplemental Table S1. The information of reads obtained by RNA-seq.

Sample	Processed fragments	Mapping rate
Untreated_1	17,500,000	78.68%
Untreated_2	21,500,000	80.99%
Untreated_3	18,500,000	78.98%
Treated_1	22,000,001	75.63%
Treated_2	20,500,000	75.97%
Treated_3	18,000,000	76.19%