

Table S3 Mapping statistics of RNA-seq data

Sample	Raw reads	Clean reads	Clean Ratio	Q20(%)	Q30(%)	GC content(%)
OE:: <i>llm1</i> _1	56132880	55763098	0.993412381	98.36	95.09	54.22
OE:: <i>llm1</i> _2	63294932	62772572	0.991747207	98.4	95.18	54.32
OE:: <i>llm1</i> _3	52481180	52067124	0.992110391	98.33	95.02	54.36
WT_1	52014002	51659080	0.993176414	98.41	95.24	54.22
WT_2	57594222	57246210	0.993957519	98.43	95.26	54.19
WT_3	58672768	58282756	0.993352759	98.42	95.25	54.26