Table S2. Statistical table of RNA-seq data.

Sample	Total reads	Total mapped	Multiple mapped	Uniquely mapped
C96_1	50801914	48812447(96.08%)	2744756(5.4%)	46067691(90.68%)
C96_2	42586554	40856104(95.94%)	2196587(5.16%)	38659517(90.78%)
C96_3	41335472	39364345(95.23%)	2191190(5.3%)	37173155(89.93%)
D96_1	45119030	43293614(95.95%)	2289748(5.07%)	41003866(90.88%)
D96_2	44093210	42280961(95.89%)	2221617(5.04%)	40059344(90.85%)
D96_3	45920218	43974094(95.76%)	2324046(5.06%)	41650048(90.7%)
R96_1	45730762	43895433(95.99%)	2212941(4.84%)	41682492(91.15%)
R96_2	50864774	48596561(95.54%)	2445482(4.81%)	46151079(90.73%)
R96_3	49818742	47767077(95.88%)	2699682(5.42%)	45067395(90.46%)
C163_1	43464170	41289016(95.0%)	2302457(5.3%)	38986559(89.7%)
C163_2	48033256	45456188(94.63%)	2504451(5.21%)	42951737(89.42%)
C163_3	43481320	41487312(95.41%)	2391918(5.5%)	39095394(89.91%)
D163_1	50922110	48366005(94.98%)	2745527(5.39%)	45620478(89.59%)
D163_2	48521900	45996295(94.79%)	2580383(5.32%)	43415912(89.48%)
D163_3	48172766	45576488(94.61%)	2404490(4.99%)	43171998(89.62%)
R163_1	45766292	43492962(95.03%)	2697161(5.89%)	40795801(89.14%)
R163_2	45012922	42653550(94.76%)	2425358(5.39%)	40228192(89.37%)
R163_3	50920950	48260376(94.78%)	2674438(5.25%)	45585938(89.52%)