

TableS5 Information of RNA-seq data

Sample	CleanBases (bp)	CleanQ30(%)	Total reads	Read mapping rate	Multiple mapping rate
DPL971+1R1	7,168,622,700	95.38	23895409	91%	5%
DPL971+1R2	6,412,138,500	95.4	25321633	91%	9%
DPL971+3R1	7,571,760,300	95.31	25239201	91%	6%
DPL971+3R2	7,265,123,100	95.46	23930032	91%	5%
DPL971+5R1	7,042,378,500	95.27	23474595	91%	5%
DPL971+5R2	8,017,725,900	95.28	27254966	91%	6%
DPL972+1R1	7,596,489,900	95.41	21373795	91%	7%
DPL972+1R2	8,608,806,300	95.46	28696021	91%	7%
DPL972+3R1	7,179,009,600	95.25	24217077	91%	4%
DPL972+3R2	7,619,923,500	95.24	25399745	91%	5%
DPL972+5R1	8,176,489,800	95.51	26725753	92%	6%
DPL972+5R2	7,619,780,400	95.35	25399268	91%	7%