

Table S2 Data mapped to the honeybee genome

Samples	Total Reads	Mapped Reads	Mapped Ratio%	Genome Size	Sequence Depth
CQ-1	44062202	30941071	70.22	214422117	21.64
CQ-2	50400498	36733405	72.88	214422117	25.70
CQ-3	49262244	34203016	69.43	214422117	23.93
NQ-1	59349720	43176744	72.75	214422117	30.20
NQ-2	43985428	31771053	72.23	214422117	22.23
NQ-3	48804460	35490757	72.72	214422117	24.83