

Table S4. Summary statistics of filtered scaffolds of the *Cacopsylla jukyungi* genome for microsatellite marker identification (from mono-mer to Hexa-mer).

Type	Counts	Length (bp)	Percent (%)	Average length (bp)	Relative abundanc e (loci/Mb)	Relative density (bp/Mb)
Mono	307,755	5,654,103	48.8	18.37	351.98	6,466.6
Di	105,092	2,970,530	16.66	28.27	120.19	3,397.4
Tri	137,256	3,737,724	21.76	27.23	156.98	4,274.84
Tetra	63,228	1,314,648	10.03	20.79	72.31	1,503.56
Penta	10,728	260,815	1.7	24.31	12.27	298.29
Hexa	6,596	240,798	1.05	36.51	7.54	275.4