

Table S5. Statistics of SSRs identified in kabuli chickpea transcripts.

SSR mining	
Total number of sequences examined	43,389
Total size of examined sequences (bp)	46,212,215
Total number of identified SSRs	5,409
Number of SSR containing sequences	4,398 (10.14%)
Number of sequences containing more than one SSR	743
Number of SSRs present in compound formation	478
Frequency of SSRs	One per 8.54 kb
Distribution of SSRs in different repeat types	
Di-nucleotide	2,216 (40.97%)
Tri-nucleotide	2,920 (53.98%)
Tetra-nucleotide	132 (2.44%)
Penta-nucleotide	48 (0.89%)
Hexa-nucleotide	93 (1.72%)