

Table 5 Statistics of SSRs identified in transcripts

SSR Mining:	
Total number of sequences examined:	34678
Total size of examined sequences (bp):	27932177
Total number of identified SSRs:	5228
Number of SSR containing sequences:	4391
Number of sequences containing more than one SSR:	681
Number of SSRs present in compound formation:	337
Distribution to different repeat type classes:	
Mono-nucleotide	2019
Di-nucleotide	1271
Tri-nucleotide	1818
Tetra-nucleotide	78
Penta-nucleotide	17
Hexa-nucleotide	25