

Table 5 Statistics of SSRs identified in transcripts

SSR Mining:	
Total number of sequences examined:	6999
Total size of examined sequences (bp):	2110290
Total number of identified SSRs:	1405
Number of SSR containing sequences:	1304
Number of sequences containing more than one SSR:	86
Number of SSRs present in compound formation:	64
Distribution to different repeat type classes:	
Mono-nucleotide	1218
Di-nucleotide	87
Tri-nucleotide	90
Tetra-nucleotide	7
Penta-nucleotide	2
Hexa-nucleotide	1