

Table 5 Statistics of SSRs identified in transcripts

SSR Mining:	
Total number of sequences examined:	7256
Total size of examined sequences (bp):	3226271
Total number of identified SSRs:	3107
Number of SSR containing sequences:	2191
Number of sequences containing more than one SSR:	747
Number of SSRs present in compound formation:	747
Distribution to different repeat type classes:	
Mono-nucleotide	2589
Di-nucleotide	235
Tri-nucleotide	243
Tetra-nucleotide	28
Penta-nucleotide	10
Hexa-nucleotide	2