

Table 5. Statistics of SSRs identified in transcripts

SSR Mining:	
Total number of sequences examined:	22748
Total size of examined sequences (bp):	11444673
Total number of identified SSRs:	1150
Number of SSR containing sequences:	1055
Number of sequences containing more than one SSR:	92
Number of SSRs present in compound formation:	48
Distribution to different repeat type classes:	
Mono-nucleotide	362
Di-nucleotide	243
Tri-nucleotide	529
Tetra-nucleotide	10
Penta-nucleotide	3
Hexa-nucleotide	3
