

Table S2: Polimorphisms detected in SOS1 regulatory regions

Polimorphism/Allele	Frequency*
promoter region	
rs368211853	
A	nd
G	nd
rs56263073	
-	0,1469
C	0,9853
rs189109845	
T	0,0013
G	0,9986
rs375493196	
C	nd
G	nd
rs368569135	
C	nd
A	nd
rs139592595	
G	0,9985
A	0,0014
rs201218035	
G	0,9963
C	0,0036
rs56360279	
G	nd
A	nd
rs372010415	
G	nd
A	nd
rs181491122	
A	0,8576
G	0,1423
rs149710810	
GGGGCGGAGCT	0,9063
-	0,0936
rs377094543	
CTGGGGCGGAG	nd
-	nd
rs376643748	
G	nd
T	nd
rs113095722	
C	nd
G	nd
rs184259916	
G	0,9958
A	0,0041

rs57378308	
G	0,8263
A	0,1736
rs146749262	
G	0,9981
A	0,0018
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miR-124 binding site	
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rs115465194	
A	0,9952
G	0,0047

*the allele frequencies are from 1000genomes database