Table S1. Sequences of oligonucleotides used in this study

Primer	**Sequence** [a]
full-length 4HA-myb2	
bgl2-ha-myb2-5'	AGATCTTACCCATACGATGTTC
N-terminal deletion	
bamh1-myb2(48)-5'	TTTTTTGGATCCAAAAAGCAAAAATTCACACCAG
bamh1-myb2(55)-5'	TTTTGGATCCGAAGAAGATGAAATGCTCAAGC
not1-myb2(48)-5'	GCGGCCGCATGAAAAAGCAAAAATTCACACCAG
C-terminal deletion	
myb2(T133ochre)-5'	CCTGGAAGATAAGATATTCATATCAAGAATAGATGGGT
myb2(T133ochre)-3'	TTGATATGAATATCTTATCTTCCAGGGAAAAACTTTGC
myb2(I144ochre)-5'	GATGGGTTACATAATCAAATAAGCTTGGAATACCTCA
myb2(I144ochre)-3'	GCTTATTTGATTATGTAACCCATCTATTCTTGATATGAA
TetR fusion protein	
myb2(143)-tetR-5'	TGGGTTACAATGTCTAGATTAGATAAAAGTAAAGTGATT
myb2(143)-tetR-3'	TCTAATCTAGACATTGTAACCCATCTATTCTTGATATGA
not1-SV40-tetR-5'	GCGGCCGCATGAAGAAGAAGCGCAAGGTGTCTAGATTAGATAAA AGTAAAG
neo-3'	GCCGAATAGCCTCTCCACCCAA
Luciferase fusion protein	
xho1-luc-5'	TCTCTCGAGATGGAAGACGCCAAAAACATAAAG
bamh1-SV40-tetR-5'	AAAGAAGAAGCGTAAGGTCGGATCCATGTCTAGATTAGATAAAA G
bgl2-SV40-ha-3'	TTCTTCTTTGGAGATCTAAGAGCGTAATCTGGAACATCGTATGGG
site-directed mutagenesis	
myb2(K51A)-5'	AAAAAAGCAAGCATTCACACCAGAAGAAGATGAAA
myb2(K51A)-3'	TGGTGTGAATGCTTGCTTTTTTGCAGCTTTTAA
myb2(I74A)-5'	GGAAAATGGCTGCTGCTACTTTTCCAAATAGAAA
myb2(I74A)-3'	GTAGCAGCAGCCATTTTCCAATCGCTTCCAT
myb2(I74P)-5'	GGAAAATGCCTGCTGCTACTTTTCCAAATAGAAA
myb2(I74P)-3'	GTAGCAGCAGGCATTTTCCAATCGCTTCCAT
myb2(KK48~49AA)-5'	AAAGCTGCAGCAGCGCAAAAATTCACACCAGAAGAA
myb2(KK48~49AA)-3'	GAATTTTTGCGCTGCTGCAGCTTTTAAATTCACTTGTAC
myb2(KR61~62AA)-5'	GAAATGCTCGCGGCTGCTGTCGCTCAACATGG
myb2(KR61~62AA)-3'	AGCGACAGCAGCCGCGAGCATTTCATCTTCTTCTGGT
myb2(K138A)-5'	ATTCATATCGCGAATAGATGGGTTACAATTTCAAATA
myb2(K138A)-3'	CCCATCTATTCGCGATATGAATATCTGTTCTTCCAGGG
myb2(N139A)-5'	TCATATCAAGGCTAGATGGGTTACAATTTCAAATAAG
myb2(N139A)-3'	AACCCATCTAGCCTTGATATGAATATCTGTTCTTCCAG
recombinant protein	
sac1-myb2(48)-5'	TTTTGAGCTCAAAAAGCAAAAATTCACACCAG
myb2(148)-bamh1-3'	TTTTGGATCCAAGCTTATTTGAAATTGTAACCCA

[a] The restriction site as designed into the name of the primer is underlined, where as the site to be mutated is boxed.