

**Table S9** Functionally identified R2R3-MYBs in other plants.

Gene	Species/Cultivars	Genome id/Accession number/Protein Sequence
AcMYBF110	<i>Actinidia chinensis</i>	MESVTLGVRKGAWTEEDKLLKKCIEKYGEGKWHQVPLRSGLNRCRKSCRMRWLNLYRPNIINRGNFTAD EVLDIIRLHKLLGNRWSLIAGRLPGRTSDNVKNYWNTHLQKKLITYPRAQPIPKTQKTIVPKGTEAQPRAHPK SPPRPSPPSNNEILWWWDNKTVSPQIDNIGHWSIDGSIFEELGNLQSAGDSFLQQNQSDWSDIFLEDVNL RDLLGDD
AtMYB123	<i>Arabidopsis thaliana</i>	AT5G35550
AtMYB4	<i>Arabidopsis thaliana</i>	AT4G38620
AtPAP1	<i>Arabidopsis thaliana</i>	AAG42001.1
CsMYB75	<i>Camellia sinensis</i>	MDSVCCVPLGVRKGAWTGEEDSLLKNCIEQYGEGMWHQIPSAGLNRCRKSCRLRWLNLYRPNIKRGNF TMDEVLDIILKLQKLLGNRWSLIAGRLPGRTANDVKNYWNTNLQKKLITQREKVAKTQEKMETIIIRPRPRIF SKNQPRLMKDCTAIENIRTRDNLSEPFPLPLPPSRDDGISWDNIVVNSKINNGITWSANGLIEANIGNW DGKIRPGHTSCGNFIDEDQSDWSDIFFNNVNLWDLLGDE
FaMYB1	<i>Fragaria × ananassa</i>	AAK84064.1
FaMYB10	<i>Fragaria × ananassa</i>	ABX79947.1
FaMYB11	<i>Fragaria × ananassa</i>	AFL02461.1
LAP1	<i>Medicago truncatula</i>	ACN79541.1
LcMYB1	<i>Litchi chinensis</i>	MSHLLGASAYSATSAEGYIGVRKGAWTEEDNLLKKCIEIYGEQKWHQVFVRAGLNRCRKSCRLRWLNLYLK PNIKRGEFMEDEVLDLILRLHKLLGNRWSLIAGRLPGRTANDIKNYWNTHLRKKAVASKQESKMTSTTTQ STQKINVIPQPRTFAKKTSEWVNTKITMEENSRDHLGHPNLCKASPSTS LDTPDNNEIMMWERLLESQ QFFLETQNSMGWSERTMSNFWAISPLRTMCAGSTGFEVNQNCLDKHHDDFDATLWNLLSIEEDNA K
LhMYB12-lat	<i>Lilium hybrid 'Tiny Padhye'</i>	BAO04193.1
MdMYB1	<i>Malus × domestica</i>	DQ886414.1
MdMYB1	<i>Malus × domestica</i>	BAM84360.1
MdMYB10	<i>Malus × domestica</i>	ACQ45201.1
MdMYB16	<i>Malus × domestica</i>	ADL36756.1
MdMYB1A	<i>Malus × domestica</i>	BAF80582.1
PhAN2	<i>Petunia x hybrid</i>	ABO21074.1
PpMYB10.1	<i>Prunus persica</i>	Prupe.3G163100
PpMYB10.4	<i>Prunus persica</i>	XPO20420992.1
PpMYB17	<i>Prunus persica</i>	ALO81020.1
PpMYB18	<i>Prunus persica</i>	ALO81021.1
PpMYB19	<i>Prunus persica</i>	ALO81022.1
PpMYB20	<i>Prunus persica</i>	ALO81023.1
RsMYB1a	<i>Raphanus sativus</i>	MEGSSKGLRKGAWTAEDSLLRQCIDKYGEGKWHQVPLRAELNRCRKSCRLRWLNLYLKPSIKRGKLN SDE VDLLRLHKLLGNRWSLIAGRLPGRTANDVKNYWNTHSKKHEPGCKTCMKKEKRNIPCSSTTLACKIDVFK PRPRSFTVNNGCShIIGMPKPDVVPLCLRNNTKVCENIATCNKDDDKSELVCNLMGCGNMWWESLLD ESQDPAAALYPEATATKKAVTSEFDVDHLWSLLDGETGT