

Figure S5 Alignment of the lignin repressor MYB members from gymnosperm and angiosperm species: MYB3-like and MYB4-like copies from *Picea abies* (Pa) and *Pinus taeda* (Pt) along with GbMYB1 from *Ginkgo biloba* (Gb); CfMYB5 from *Cryptomeria fortune* (Cf); MYB3, MYB4, MYB7 and MYB32 from *Arabidopsis thaliana* (At); MYB156 and MYB221 from *Populus trichocarpa* (Potri); EgMYB1 from *Eucalyptus gunnii* (Eg); ZmMYB31 and ZmMYB42 from *Zea mays* (Zm) and PvMYB4 from *Panicum virgatum* (Pv). Alternative EAR domain in Pt_AtMYB4-like1, Pt_AtMYB4-like3, Pt_AtMYB4-like4 and Pa_AtMYB4-like1 are marked with box and, bold and underlined.

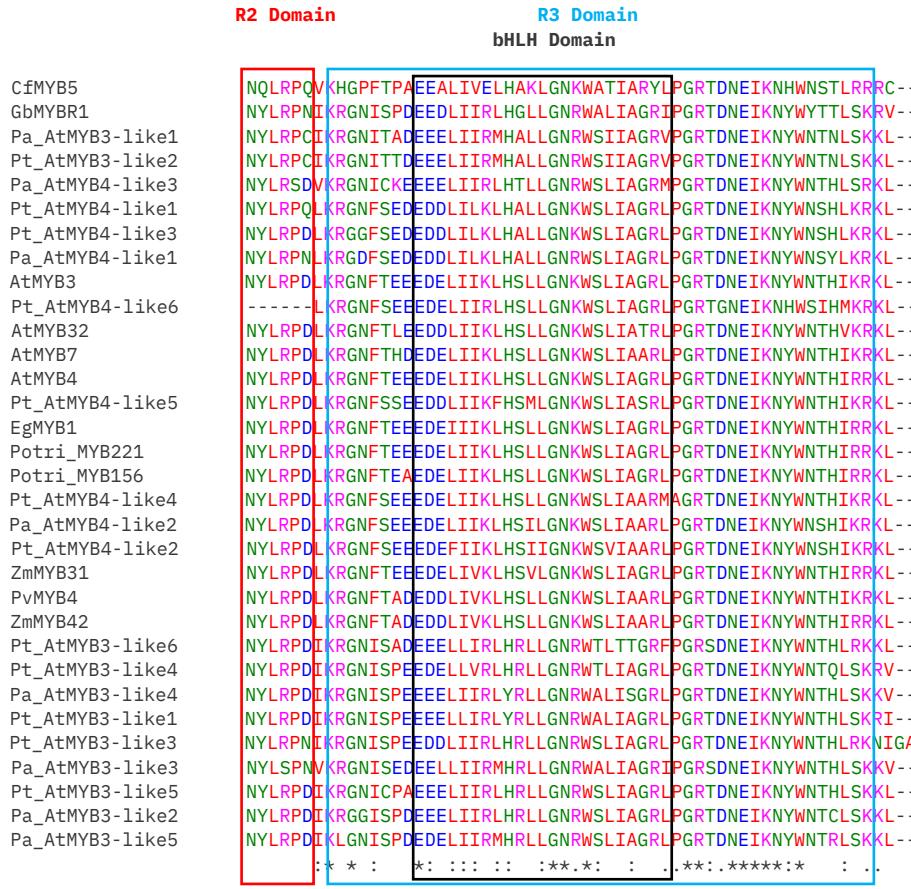
CfMYB5	-----	-MGRS
GbMYB1	-----	-MGRS
Pa_AtMYB3-like1	-----	-MGRA
Pt_AtMYB3-like2	-----	-MGRA
Pa_AtMYB4-like3	-----	-MGRA
Pt_AtMYB4-like1	-----	-MARA
Pt_AtMYB4-like3	-----	-MGRA
Pa_AtMYB4-like1	-----	-MGRG
AtMYB3	-----	-MGRS
Pt_AtMYB4-like6	-----	-MKR-
AtMYB32	-----	-MGRS
AtMYB7	-----	-MGRS
AtMYB4	-----	-MGRS
Pt_AtMYB4-like5	-----	-MGRS
EgMYB1	-----	-MGRS
Potri_MYB221	-----	-MGRS
Potri_MYB156	-----	-MGRS
Pt_AtMYB4-like4	-----	MSLFPLSLPSGIPVVAFLSMKMRKY
Pa_AtMYB4-like2	-----	-MGRS
Pt_AtMYB4-like2	-----	-MGRT
ZmMYB31	-----	-MGRS
PvMYB4	-----	-MGRS
ZmMYB42	-----	-MGRS
Pt_AtMYB3-like6	MRKFVIKSITGRSRLPSIYLLAITGKRSTEKKWSLRDFIEAVGREFVGLVRSYIMGRS	
Pt_AtMYB3-like4	-----	-MMGRS
Pa_AtMYB3-like4	-----	-MGRS
Pt_AtMYB3-like1	-----	-MGRS
Pt_AtMYB3-like3	-----	-MGRS
Pa_AtMYB3-like3	-----	-MGRP
Pt_AtMYB3-like5	-----	-MPVAGLGPAGGGEHQPWSYRPGKYKMGRS
Pa_AtMYB3-like2	-----	-MGRS
Pa_AtMYB3-like5	-----	-MGRS

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R2 Domain

CfMYB5	AECE-----KVHTNKGWPSSPEEDALLQCYVQKYG-ADNWSLISKGAGLRLCGKSCRLRWC	
GbMYB1	PMCS----KVGLLNKGAWTVEEDNLITKYFQTHDEEGGWKSVPKKAGLKRCKGKSCRLRWM	
Pa_AtMYB3-like1	PCCA----KVGLNKGAWSAEEEDSLLGKYIQTGH-EGNWRSLPKKAGLRRCGKSCRLRWL	
Pt_AtMYB3-like2	PCCT----KVGLNKGAWSAEEEDSLLGKYIQTGH-EGNWRSLPKKAGLRRCGKSCRLRWL	
Pa_AtMYB4-like3	PCCS----KVGLNRGRWTAAEDEILTKYIQTGH-EGSWRSLPQNAGLLRCGKSCRLRWI	
Pt_AtMYB4-like1	PCCP----IGDRNKGAWTREEDDKLIQYIQTGH-EGGWNSLPKAAGLLRCGKSCRLRWI	
Pt_AtMYB4-like3	PCCA----NGDRSKGAWTKEEDDRLTQYIQAHG-EGCWRSLPKAAGLLRCGKSCRLRWI	
Pa_AtMYB4-like1	PCCA----NGDRNKGAWTREEDDRLIQYIQAHG-EGCWRSLPNAAGLLRCGKSCRLRWI	
AtMYB3	PCCE----KAHMNKGAWTKEEDQLLVDYIRKHG-EGCWRSLPRAAGLQRCGKSCRLRWI	
Pt_AtMYB4-like6	-----KLLIKGFDAH-----	
AtMYB32	PCCE----KDHTNKGAWTKEEDDKLISYIKAHG-EGCWRSLPRSAGLQRCGKSCRLRWI	
AtMYB7	PCCE----KEHMNKGAWTKEEDERLVSYIKSHG-EGCWRSLPRAAGLRLRGKSCRLRWI	
AtMYB4	PCCE----KAHTNKGAWTKEEDERLVAIYIKAHG-EGCWRSLPKAAGLRLRGKSCRLRWI	
Pt_AtMYB4-like5	PCCE----KAHTNKGAWSEEDQVLIAYIRAHG-EGCWRSLPKAAGLQRCGKSCRLRWI	
EgMYB1	PCCE----KAHTNKGAWTKEEDDKLIAYIRAHG-EGCWRSLPKAAGLRLRGKSCRLRWI	
Potri_MYB221	PCCE----KAHTNKGAWTKEEDDRLIAYIRTHG-EGCWRSLPKAAGLRLRGKSCRLRWI	
Potri_MYB156	PCCE----KAHTNKGAWTKEEDDRLVAYIRAHG-EGCWRSLPKAAGLRLRGKSCRLRWI	
Pt_AtMYB4-like4	SCCE----KAHTKKKGAWTQQEDARLVAYIQAHG-EGGWRYLPKAAGLRLRGKSCRLRWI	
Pa_AtMYB4-like2	PCCE----KSHTNKGAWTKEEDDLRLIAHIRAHG-EGCWRSLPKDAGLRLRGKSCRLRWI	
Pt_AtMYB4-like2	PCCE----KGHTNKGAWTKEEDDLRLIAHIRAHG-EGRWRSLPKAAGLMRCGKSCRLRWI	
ZmMYB31	PCCE----KAHTNKGAWTKEEDDRLVAHIRAHG-EGCWRSLPKAAGLRLRGKSCRLRWI	
PvMYB4	PCCE----KAHTNKGAWTKEEDDRLVAYIRAHG-EGCWRSLPKAAGLRLRGKSCRLRWI	
ZmMYB42	PCCE----KAHTNKGAWTKEEDERLVAHYRAHG-EGCWRSLPRAAGLRLRGKSCRLRWI	
Pt_AtMYB3-like6	PSHFSAKNDIAGLKRGAWTPSEDQILSDCVKJTHG-VGQWKSMAKKTGLRRSAKSCRFRWL	
Pt_AtMYB3-like4	PGYSSYSKHEEHLKGAWTTPSEDKILIEYIKTHG-IGQWRDLPRKAGLRLRGKSCRLRWL	
Pa_AtMYB3-like4	PSCSSYSKHEDGLNRGAWTAGEDKILSEYVKTHG-VGRWRSLPKKAGLRLRGKSCRLRWL	
Pt_AtMYB3-like1	PSLWSCSKQEDGICNRGAWTASEDKILSEYVKTHG-VGRWRSLPIKTGLKRCAKSCRRLWL	
Pt_AtMYB3-like3	RRCS----TDEGSNRAAWTTKEDMILRDYINIHGVQGGWTSVRDQTALNRSGKSCRLRWL	
Pa_AtMYB3-like3	QSCS----GLKLKGAWTSEEDLLLKKCIEVHG-EGSWPLLPQKTGLQRSGKSCRLRWL	
Pt_AtMYB3-like5	PCCS----KEGLNRGSWTKREDMILSEYVRIHG-DGGWRNLPEKAGLKRCKGKSCRLRWL	
Pa_AtMYB3-like2	PCCS----KEGLNRGAWTKTEDIILCEYIRIHG-DGGWRTLPKKAGLKRCKGKSCRLRWL	
Pa_AtMYB3-like5	-CCS----KEGLNRGAWSRKEDMILSEYIRIHG-DGGWTNLQPQAGLKRCKGKSCRLRWL	

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CfMYB5	-----Q-----	-DSNP
GbMYB1	-----ALK-----	-GNEA
Pa_AtMYB3-like1	-----AVR-----	-GIDP
Pt_AtMYB3-like2	-----AVR-----	-GIDP
Pa_AtMYB4-like3	-----ANR-----	-GIDT
Pt_AtMYB4-like1	-----ISM-----	-GVDP
Pt_AtMYB4-like3	-----ISM-----	-GVDP
Pa_AtMYB4-like1	-----ISM-----	-GVDP
AtMYB3	-----LSR-----	-GIDP
Pt_AtMYB4-like6	-----SSR-----	-GLHP
AtMYB32	-----LRK-----	-GIDP
AtMYB7	-----LSK-----	-GIDP
AtMYB4	-----INR-----	-GIDP
Pt_AtMYB4-like5	-----LEK-----	-GIDP
EgMYB1	-----LNR-----	-GIDP
Potri_MYB221	-----LNR-----	-GIDP
Potri_MYB156	-----LNR-----	-GIDP
Pt_AtMYB4-like4	-----LSR-----	-GLDP
Pa_AtMYB4-like2	-----LSR-----	-GLDP
Pt_AtMYB4-like2	-----LSR-----	-GIDP
ZmMYB31	-----LSR-----	-GIDP
PvMYB4	-----LSR-----	-GIDP
ZmMYB42	-----LGS-----	-GIDP
Pt_AtMYB3-like6	-----QM-----	-GM-P
Pt_AtMYB3-like4	-----QKG-----	-EFDP
Pa_AtMYB3-like4	-----RMG-----	-ESDP
Pt_AtMYB3-like1	-----QKG-----	-EFKH
Pt_AtMYB3-like3	LNKDRDRSEAKMTEEDM	GVSPRLDLVGANQIAVNIENNCVGRQICVWGSQRCIRGLRRALS
Pa_AtMYB3-like3	-----NRSR-----	-SDDF
Pt_AtMYB3-like5	-----SSQ-----	-ESQD
Pa_AtMYB3-like2	LLSMNK	-SQSK
Pa_AtMYB3-like5	-----L-----	-SIDD

CfMYB5	-GLAAAVTH-	NED-
GbMYBR1	-KEHKTVPM-	KRS-
Pa_AtMYB3-like1	-KTHKKVTT-	DSINRASDRFNQ
Pt_AtMYB3-like2	-KTHKKITT-	DGTNRVNGDGFN
Pa_AtMYB4-like3	-ATHKPLGE-	VAEIFIQAKP-
Pt_AtMYB4-like1	-LTHRPFQK-	ICR-
Pt_AtMYB4-like3	-LTRRPFQR-	NSR-
Pa_AtMYB4-like1	-LNHRPFQK-	NSR-
AtMYB3	-NSHRLINE-	SVVSPSSLQND-
Pt_AtMYB4-like6	-QTHRSLRP-	PH-
AtMYB32	-ATHRPINE-	TKTSQDS-
AtMYB7	-ATHRGINE-	AKI-
AtMYB4	-TSHRPPIQE-	SSASQD-
Pt_AtMYB4-like5	-NTHLPILP-	DHGS-
EgMYB1	-ATHRLINE-	PAQD-
Potri_MYB221	-ATHRPLNE-	PA-
Potri_MYB156	-ATHRPLNE-	PAV-
Pt_AtMYB4-like4	-QTHRSCLCP-	PH-
Pa_AtMYB4-like2	-GSHRPLCF-	PH-
Pt_AtMYB4-like2	-RSHRPLCP-	P-
ZmMYB31	-VTHRNPVTE-	H-
PvMYB4	-VTHRPIAD-	
ZmMYB42	-VTHRRVAG-	
Pt_AtMYB3-like6	-IDSRNYNE-	KEVQDRY-
Pt_AtMYB3-like4	-EYQEPEFKE-	RAIPAPSIYDKG
Pa_AtMYB3-like4	-RFQNLCKR-	RDVPPYPNYD-
Pt_AtMYB3-like1	-KFQNHFKR-	KHMPSDSNYG-
Pt_AtMYB3-like3	RDL SARL RWL NYL RPD IKRGN ISP DEE ELI IIR MH RLL GNR WS LIAG RL PG RT DNE IKN YW	
Pa_AtMYB3-like3	-NCNKE-	
Pt_AtMYB3-like5	-KTSKKLSA-	SSKSPVP-
Pa_AtMYB3-like2	-VTKNVKST-	SK-
Pa_AtMYB3-like5	-SQSKSTAR-	

CfMYB5	-NSSSLKRSSCSNDDGS-	STLDENCGFEAA-
GbMYBR1	-RGHSACKQLIMPDSNT-	KMQDLLSASPTRKELEVSTNQS-
Pa_AtMYB3-like1	RKGKLY-DCSQRSQLERNVVRAGQSTGLVIPNVHNLKADLKQYIAGPREIQSSNSIR	
Pt_AtMYB3-like2	QRKGKEIYDSPQKPRQPERNVARAAQSTGVVIPNVHNLKADLKAQYIARIREFKSSNTIS	
Pa_AtMYB4-like3	-SIITNVRNTSLNDIKP-	VRCSENIEASMVCIDSALKKISK
Pt_AtMYB4-like1	-HDQPLSENVSEPEIAS-	SGKFVQDFFFHC
Pt_AtMYB4-like3	-HDLSLPEKEVPEIAR-	PREIVENFFHSP-
Pa_AtMYB4-like1	-HRPSASGNVGEPEGTAR-	SREIVPDFFHCP-
AtMYB3	-VVETIHLD FSGPVKPE-	PVREEIGMVNNCE-
Pt_AtMYB4-like6	-NSSTTLNSPPPDPHQI-	LAFESRRRAEIADFFQLEGLKS
AtMYB32	-SDSSKTEDPLVKILSFG-	PQLEKIANFGDER-
AtMYB7	-SDLKKTQDQIVKDVSF-	VTKEETDKSGDQ-
AtMYB4	-SKPTQLEPVTSNTINISF-	TSAPKVETFHESI-
Pt_AtMYB4-like5	-FGSSNENHVFSKRSRC-	KPSEVLTDFFQQQ-
EgMYB1	-HHDEPTISFAANSKEI-	-KEMKN-
Potri_MYB221	-QEASTTISFSTTS-	VKEESLSSVKEES-
Potri_MYB156	-QEATTISFTTTTSV-	LEEESLGSIKEE-
Pt_AtMYB4-like4	-NSNSTLSSPSVPEHEI-	SPLQNPRTADIADFFQQD-
Pa_AtMYB4-like2	-HNNNTSPSRPAPDHEF-	PAFQNLKTPETDFFQQH-
Pt_AtMYB4-like2	-SNTTSLSWSAPDREF-	PAIQNLRTPETDFFQHN-
ZmMYB31	-HASNITISFETEAAA-	ARDDKKGAVFRLE-
PvMYB4	-AARNVTISFQPDA-	PSQQQLSDDAEAP-
ZmMYB42	-GAATTISFQPSPNTAV-	AAAAE-
Pt_AtMYB3-like6	-DGVTQNTPLKTMPLRYWEKEE-	VDVENLDCNYDC-
Pt_AtMYB3-like4	FLHRD-NDSYDILPFKTRAVRH-	CGRTIVDGDNLDSYDHDC-
Pa_AtMYB3-like4	-EDIYASQNAPKTTGVRGYGRAV-	AELNFDSYNPDH-
Pt_AtMYB3-like1	-EDRVVDKNINAGEDTPSKATGEISCARAVGKVVDNLDKYNQDN-	
Pt_AtMYB3-like3	NTHLSKKLTMSSNPKFPMAPADPEQTASSSESEDVVRSAHYRVIHSGMRЛАPAKIG	
Pa_AtMYB3-like3	-NVVQTLTSEEKG-	GQEIGAENYSSH-
Pt_AtMYB3-like5	-VQNRVFKATAVKITTA-	VRHTEIIRPNACN-
Pa_AtMYB3-like2	-SSPVQNRVFKTTPVKV-	TTAVRLSETIRPK-
Pa_AtMYB3-like5	-NLETSSKSPPPPNHW-	FKTTPIKITTAVKFSETVGPKRCN-

CfMYB5	-----DS-----	KR-----	LKS-----
GbMYBR1	-----	-----	LSE-----
Pa_AtMYB3-like1	SSSPINTPIQPKSNDFMQPQPVVSSDAGKQ-----	-----	TDD-----
Pt_AtMYB3-like2	SSSRLNAQIEPKSRELSTEDPIFRCSSASEKTRETTHPDFMEPHPVASSDAGKQ-----	-----	TND-----
Pa_AtMYB4-like3	KSSKKRSVWKEK-----QM-----	-----	PFE-----
Pt_AtMYB4-like1	-----	-----	PSE-----
Pt_AtMYB4-like3	-----SEL-----CG-----	-----	DSE-----
Pa_AtMYB4-like1	-----PE-----	-----	LST-----
AtMYB3	-----SSG-----	-----	TT-----
Pt_AtMYB4-like6	-----	-----	SPI-----
AtMYB32	-----	-----	IQK-----
AtMYB7	-----KQ-----NK-----	-----	YIR-----
AtMYB4	-----SFPGKSEK-----	-----	ISM-----
Pt_AtMYB4-like5	-----TPG-----GT-----	-----	SSV-----
EgMYB1	-----	-----	NAE-----
Potri_MYB221	-----NKE-----KT-----	-----	ISA-----
Potri_MYB156	-----NKE-----KI-----	-----	ISA-----
Pt_AtMYB4-like4	-----RS-----ER-----	-----	SPV-----
Pa_AtMYB4-like2	-----	-----	RSE-----
Pt_AtMYB4-like2	-----RS-----ES-----	-----	SPI-----
ZmMYB31	-----EEEERNKATMVVGRD-----	-----	RQS-----
PvMYB4	-----	-----	PPP-----
ZmMYB42	-----	-----	TAA-----
Pt_AtMYB3-like6	-----	-----	PPD-----
Pt_AtMYB3-like4	-----	-----	VPD-----
Pa_AtMYB3-like4	-----	-----	VPD-----
Pt_AtMYB3-like1	-----	-----	VPV-----
Pt_AtMYB3-like3	GTSISAGVPAPVSSDEVINMKSWKQLLEDLSMSDFDDSEDDTSLMGIELSDDVLGSEAPL-----	-----	QPK-----
Pa_AtMYB3-like3	-----	-----	PGE-----
Pt_AtMYB3-like5	-----GDGCSNFI-----	-----	GSM-----
Pa_AtMYB3-like2	-----	-----	SAE-----
Pa_AtMYB3-like5	-----GYGRSNCS-----	-----	-----

**EAR
suppression domain**

CfMYB5	HGNDCLEES-----	KQAPQVVYKPVPVASC-----	GFLKPPQQEAT-----
GbMYBR1	SVVSNTDDVRTDSNVQSGSPGLQEIRASRIFLPS-----	-----	FRGNQVSSHSELMAPANP
Pa_AtMYB3-like1	STVYCSSDSAASCALIDHLSSEDDQYLSLEGNSNECYSTSVAEESGTLKSSNPQTHSEA-----	-----	-----
Pt_AtMYB3-like2	STVYCSSDSAASCSLIEHFSSSEDDHHYLSEMEGSNECYSQTLAEDYGSLKPKSTPHTESEP-----	-----	-----
Pa_AtMYB4-like3	EQIPSGQSSSCESCSGNLSSPDLLMAACCNPPSCDQVKKLSFEKTPCQDRSTDHVLYRE-----	-----	-----
Pt_AtMYB4-like1	LSVESAQLSNAAGGLSARDGQPDLNLDLGLTV-----	-----	TCPSIHSEG-----
Pt_AtMYB4-like3	QVSDAASES-----LARD-EPS-TLNLLNLISINW-A-----	-----	SMHVAVKEEP-----
Pa_AtMYB4-like1	KSDQVSDAASGLARKELPRFLNLNLISITRPS-----	-----	IHAAGKEAVVNS-----
AtMYB3	EKDYGNEED-----	-----WVLNLNLISVGPSY-----	-----RYESTRKVSVVDSAEST-----
Pt_AtMYB4-like6	ESAASTDEE-----	HPCPDINLDLCMGLSS-----	-----NSAPA-----
AtMYB32	RVEYSVVEE-----	RCLDLNLNLRLRISPPW-----	-----QDKLHDE-----
AtMYB7	NGLVCKEER-----	VVVEEKIGPDNLNLRLRISPPW-----	-----
AtMYB4	LTFKEEKDE-----	CPVQEKFDPDLNLNLRLISLPD-----	-----DVDR-----
Pt_AtMYB4-like5	LKVIAKNQD-----	HDYDKDVNLDSITLPS-----	-----LCSSQGSSSSRELSSGI-----
EgMYB1	LNFMCNLLEESADVASSARERC-----	CPDNLNLNLGISPPS-----	-----HQLHOPEP-----
Potri_MYB221	AAFICKEEK-----	TPVQERCPDNLNLRLRISLPC-----	-----QNOQDRH-----
Potri_MYB156	TAFCVKEEK-----	TQVQERCPDNLNLRLGISLPS-----	-----QNOQDPDH-----
Pt_AtMYB4-like4	ENAACKEEE-----	HPDNLNLNL CISLPS-----	-----NSTLAAN-----
Pa_AtMYB4-like2	SSPIVPEAS-----	DAEEHPDNLNLNL CISLPS-----	-----NSPPRRE-----
Pt_AtMYB4-like2	LPAAPDAEE-----	DPDNLNLNL CISLPS-----	-----NSSPAANTAQS-----
ZmMYB31	QSQSHSHPAGEWGQGKRPLKCPDNLNLCLISPPC-----	-----	-----QEEEEMEEAAMR-----
PvMYB4	PPPQQQQQQ-----	LKPPPRCPDNLNLNL CISLPC-----	-----HKEEEDQELIKPA-----
ZmMYB42	QAPIKAET-----	AAVKAPRCPDNLNLNL CISLPC-----	-----QHEDDGEEEEELDLTICKPAV-----
Pt_AtMYB3-like6	SHDGINPEE-----	EGMKGDDHVSEKDASETWCQLLEDCTGYYLYDRLEFVQGL-----	-----
Pt_AtMYB3-like4	THDGINQEE-----	GLKATNHHVVEIDTSKSWCQLLEDCTGDYQYDRLEEISGLQPN-----	-----
Pa_AtMYB3-like4	THDGSTQEG-----	LKAADDHVVVEIDTSKSWSQLLLEHCMGDYQYDRLQPNASS-----	-----
Pt_AtMYB3-like1	TNEGGINQEG-----	GLRAADDNVVEIDTSKSWLQLLEDCMGEYHYNLLQPNIMNTNK-----	-----
Pt_AtMYB3-like3	MEIQTNVSCSTSGPNEPVHSHGHLADATSNSRSSESNVLYSQDCNVLSSLPTDNTLDFR-----	-----	-----
Pa_AtMYB3-like3	SRISINTDH-----	NQHADMLEEPSTGLTK-----	-----CQSYRNTPTEQTDL-----
Pt_AtMYB3-like5	ALKLCNVEN-----	TTKSSSCGLLVNNNSSTSQPERNAFATTDNLVQTEATNCTSSMLNF-----	-----
Pa_AtMYB3-like2	NGYGCSNRS-----	SDEAFKLCAKETTNSSKSWCDLLGNDTEGRISNSSMSLALERS-----	-----
Pa_AtMYB3-like5	AIKLCNIKE-----	NTNSIPHDDSEHIVFDMTDVNLL-----	-----ETEAGNCTSIAWSL-----

**EAR
suppression domain**

CfMYB5	-VSSSASVPPT-----	SLCLSLPGLPEPTTPSSKPVDAQTRKDSPSPSPTP
GbMYBR1	MIESNIORKLFLSLVDD-----	YLSVSTELSLGFSGMNC-----
Pa_AtMYB3-like1	I----CDSRERDNGGPVQKHQDFPEYDVFSFFDVRNAENEICC-----	SADQWVHEQE-----
Pt_AtMYB3-like2	I----CDSRERDNDSHVQKHQDFPEYDVFSFFDVRNAENEICC-----	IKEEDCNSDDPVNNSK-----LWEESFNSINILNSDMENMMFGYLETEVELFGCLA-----
Pa_AtMYB4-like3	IKEEDCNSDDPVNNSK-----LWEESFNSINILNSDMENMMFGYLETEVELFGCLA-----	KQNVVNSEQGE-----NDLGEFGNGLAYHSGECA-----
Pt_AtMYB4-like1	KQNVVNSEQGE-----NDLGEFGNGLAYHSGECA-----	I----LKSXPGEEEGA-----GGRFDEKGYL-FRYEECP-----
Pt_AtMYB4-like3	I----LKSXPGEEEGA-----GGRFDEKGYL-FRYEECP-----	EQGESNLSEGK-----GYLCRHGECPPQSPTDG-----
Pa_AtMYB4-like1	EQGESNLSEGK-----GYLCRHGECPPQSPTDG-----	RRWGSELFGAHESDAV-----CLCCRIGLFRNESCNR-----
AtMYB3	RRWGSELFGAHESDAV-----CLCCRIGLFRNESCNR-----	
Pt_AtMYB4-like6		
AtMYB32		--RNLRFGGRVKYR-----CSACR-FGFGNGKECSCN-----NVKC-----
AtMYB7		QNQRE-----ISTCTASRFYMENDMECS-----SETVKCQ-----
AtMYB4		LQGHGKSTTPR-----CFKCS-LGMINGMERCRCG-----
Pt_AtMYB4-like5	R----LSNDCTQQDHK-----EKICCHIGLRTVGECCS-----	
EgMYB1	LLRFTGRKSDL-----CXECN-LGLKNSQNCRCS-----	
Potri_MYB221	QAFKTGGSTSL-----CFACCS-LGLQNSKDCSCS-----	
Potri_MYB156	QPFKTGGSRSL-----CFACCS-LGLQNSKDCSCN-----	
Pt_AtMYB4-like4	RAMDANSNSGL-----GVWCH-TGPQINS DPRCE-----	
Pa_AtMYB4-like2		
Pt_AtMYB4-like2	R----KTVDSNLNSGS-----SEVCYPVGLQINYGKYCD-----	
ZmMYB31	VRPAVKREAGL-----CFGCS-LGLPLTADCKCS-----	
PvMYB4	AVKREMLQAGHGTGLG-----CFGCSGLQKGAAAGCTCS-----	
ZmMYB42	VKREALQAGHGHGHGL-----CLGCGLGGQKGAAAGCSCS-----	
Pt_AtMYB3-like6	QTDRINLLPPS-----VSPSGSEILQEKQSSCN-----	
Pt_AtMYB3-like4	SMTNSSNVTTDOHQHQPQS-----PFVQSIIILQENHSSCD-----	
Pa_AtMYB3-like4	SAVQLPPQLSV-----NLSCPRLFIDLPQRSHTD-SKKKQITMDGAFEIE-----	
Pt_AtMYB3-like1	CGTDYDQNQFPS-----PTLGRDILLQENGASSCN-----	
Pt_AtMYB3-like3	IQDFSLESLLPSVYN-----FDEICPVQSQLNSSDHVSCFAVAAEQAMTDIYNNT-----	
Pa_AtMYB3-like3	IHFHDY-----KFSCQNSADDLFQAGNCS-----	
Pt_AtMYB3-like5	GDQQSPYSDYFEGDAA-----NLAESLLFLTDLSSPDCN-----LLASPSDCST-----	
Pa_AtMYB3-like2	PNSYYFGTDA-----TLADSLLDLNELSSPDCS-----LLSPPSNCSA-----	
Pa_AtMYB3-like5	EEERSPHSYFAVDTA-----TLDESLSELNVLSSPDCN-----	

CfMYB5	VPTVPVPVPVQVQIPVPASAFGYVRA-----	DEAMSWMSAVVRATVAHTLAPILNSP-----
GbMYBR1	VSFKSTNSHNLYLMGSSLNT-----	DHHMPCGDQSVCRDQYRWMT-----
Pa_AtMYB3-like1	NDDQWVHEQEMPQLHSWDNQIIDQGK-----	EHFISHVNNDVTAMSWEASFWF-----
Pt_AtMYB3-like2	AEFMQQKDQEMAQLGSWEKQIDDQE-----	ENFESHVNNDVTAMSWEASFWF-----
Pa_AtMYB4-like3	DPPMPAPNADEGNMETLSSDADYNT-----	DELWSTFLSNSEKTTLPSPSSLSA-----
Pt_AtMYB4-like1	PPMLLL-----R-----	
Pt_AtMYB4-like3	VGK-----SLLR-----	
Pa_AtMYB4-like1		
AtMYB3	VSDVRTH-----	
Pt_AtMYB4-like6		
AtMYB32	QTEDSSSSSYSTSIDISSSIGYDFLGL-----	NNTRV-----LDFSTLEMK-----
AtMYB7	TENSSSISYSSIDISSSNVGYDFLGL-----	KTRI-----LDFRSLEMK-----
AtMYB4	-RMRCDVVGSSKGSDMSNGDFFLGL-----	AKKETTSL-----LGFRSLEMK-----
Pt_AtMYB4-like5	GCWD-----	
EgMYB1	VGVISETSVGYDFLGL-----	KASV-----LDYRS-----
Potri_MYB221	VIVGTTIGSSSSAGSKTGYDFLGM-----	KSGV-----LDYRGLEMK-----
Potri_MYB156	VIVSTVGGSSGTSTKTGYDFLGM-----	KSGV-----LDYRSLEMK-----
Pt_AtMYB4-like4	NRYCERNSTFELDS-----	SSGSIHGI D --- L Q ₁ ALGVQ-----
Pa_AtMYB4-like2	ESFQRL-----	
Pt_AtMYB4-like2	SGYCEQNASCFSQF-----	RLVL-----
ZmMYB31	SSSFLGL-----	RTAM-----LDFRSLEMK-----
PvMYB4	SNSHFLGL-----	RVGML-----LDFRGLEMK-----
ZmMYB42	NGHHFLGL-----	RTSV-----LDFRGLEMK-----
Pt_AtMYB3-like6	LLEEEFLQT-----	ENFNSTYF-----
Pt_AtMYB3-like4	-LIQQESP H KDFISSYFSSFLDLGE-----	QSILLP-----
Pa_AtMYB3-like4	NLELLDQEKGGRNQDWVCOGGESHDT-----	TSQEQRGMPRDLGRAESRSQQLDHGS-----
Pt_AtMYB3-like1	-FFDQEFP H SKEFISP Y SDNCLDLGE-----	PSFVIP-----
Pt_AtMYB3-like3	LEPDWLTPVEYEKQSTQSDQMNLGNFFLSEENWEEEVPTSLSKIQEELYRAILVDISNK-----	
Pa_AtMYB3-like3	-ASDCSETEILQQLLHPCSDM-----	EEYWREVGDAQLFAFDVWNMV-----
Pt_AtMYB3-like5	DFASEDLYRGPAALPEETNGFEDMC-----	DQNNMQRVNSS-SMDQLEGFYNN-----
Pa_AtMYB3-like2	DFGLQDFCGEPVLTAEANEFEVVSS-----	DQSIMQRDKGFTVLDQGIEEFYDHT-----
Pa_AtMYB3-like5	LLYSASGSSTD F ELEEFYRE-----	AATLATGINDSVFT-----

CfMYB5	PPRGSASDHG-----
GbMYBR1	-----
Pa_AtMYB3-like1	-----
Pt_AtMYB3-like2	-----
Pa_AtMYB4-like3	-----
Pt_AtMYB4-like1	-----
Pt_AtMYB4-like3	-----
Pa_AtMYB4-like1	-----
AtMYB3	-----
Pt_AtMYB4-like6	-----
AtMYB32	-----
AtMYB7	-----
AtMYB4	-----
Pt_AtMYB4-like5	-----
EgMYB1	-----
Potri_MYB221	-----
Potri_MYB156	-----
Pt_AtMYB4-like4	-----
Pa_AtMYB4-like2	-----
Pt_AtMYB4-like2	-----
ZmMYB31	-----
PvMYB4	-----
ZmMYB42	-----
Pt_AtMYB3-like6	-----
Pt_AtMYB3-like4	-----
Pa_AtMYB3-like4	RSSPS-----
Pt_AtMYB3-like1	-----
Pt_AtMYB3-like3	-----
Pa_AtMYB3-like3	-----
Pt_AtMYB3-like5	-----
Pa_AtMYB3-like2	QDEDWIHELDCLENSGAQPLSSLLLSEDEWEERSIGKLAEDHVQI
Pa_AtMYB3-like5	QPLDWIHELDYIEKSSPRALSFLLESEDEREEKLGDHTQIPDVV-----
Pa_AtMYB3-like5	-----