

MYBL1	1	AKRSR----SEDEDD-DLQYADHDYEVPPQKGLKKLWNRVKWTRDEDDKLLKLLVEQHGTD	55
		A+R R S DEDD D++ DHDY+ K K+ + +WTR+ED+KLKLLVEQ+GTD	
C-MYB	2	ARRPRHSIYSSDEDEDEIEMCDHDYDGLLPKSGKRHLGKTRWTRREEDEKLLKLLVEQNGTD	61
MYBL1	56	DWTLIASHLQNRSDFCQCHRWQKVLNPELIKGPWTKEEDQRVIELVQKYGPKRWSLIAKH	115
		DW +IA++L NR+D QCCHRWQKVLNPELIKGPWTKEEDQRVIELVQKYGPKRWS+IAKH	
C-MYB	62	DWKVIANYLPNRTDVQCCHRWQKVLNPELIKGPWTKEEDQRVIELVQKYGPKRWSVIAKH	121
MYBL1	116	LKGRIGKQCRERWHNHLNPEVKKSSWTEEEEDRIIYEAHKRLGNRWAEIAKLLPGRTDNSI	175
		LKGRIGKQCRERWHNHLNPEVKK+SWTEEEEDRIIY+AHKRLGNRWAEIAKLLPGRTDN+I	
C-MYB	122	LKGRIGKQCRERWHNHLNPEVKKTSWTEEEEDRIIYQAHKRLGNRWAEIAKLLPGRTDNAI	181
MYBL1	176	KNHWNSTMRRKVEQEGYLQDGIK <b>ERSSSKLQHKPCATM</b> -----DHLQTQNFYIPVQIP-	230
		KNHWNSTMRRKVEQEGYLQ+ SK P AT +HL P Q+	
C-MYB	182	KNHWNSTMRRKVEQEGYLQ <b>E-----PSKASQTPVATSFQKNNHLMGFGHASPPSQLSP</b>	234
MYBL1	231	-----GYQVSPDGNCVEHV-----QTSAFIQPFVDEDPDKEKK	265
		Y +++ N HV +A IQ+ + DEDP+KEK+	
C-MYB	235	<b>SGQSSVNSEYPYHIAEAQNISSHVYPVALHANIVNVPQAAAAIQRHYNDEDPKEKR</b>	294
MYBL1	266	IKELELLLSAENEVR-RKRLPPQ--PGSFSSWSGSFLMDDSM----SNTLNNLEEHTTE	318
		IKELELLLS ENE++ ++ LP Q S+ W + ++D + S ++ L EH	
C-MYB	295	IKELELLLSSTENELKGQALPTQNHTCSYPGWHSTSIVDQTRPHGDSAPVSLGEGHAT	354
MYBL1	319	FYSMDENQTVSAQQNSPTKFLAVEANAVLSSLQTIPEFAETLELIES 365	
		+ ++ + SP + + V +L +++ + EFAETL+ I+S	
C-MYB	355	PSLPADPGSLPEESASPARCMIVHQGTILDNVKNLLEFAETLQFIDS 401	
MYBL1	67	NRSDFQCCHRWQKVLNPELIKGPWTKEEDQRVIELVQKYGPKRWSLIAKHLKGRIGKQCR	126
		NR+D QCCHRWQKVLNPEL KGPWTKEEDQRVIE VQKYGPKRWS IAKHLKGRIGKQCR	
v-MYB	1	NRTDVQCCHRWQKVLNPELNKGPWTKEEDQRVIEHVQKYGPKRWSIAKHLKGRIGKQCR	60
MYBL1	127	ERWHNHLNPEVKKSSWTEEEEDRIIYEAHKRLGNRWAEIAKLLPGRTDNSIKNHWNSTMRR	186
		ERWHNHLNPEVKK+SWTEEEEDRIIY+AHKRLGNRWAEIAKLLPGRTDN++KNHWNSTMRR	
v-MYB	61	ERWHNHLNPEVKKTSWTEEEEDRIIYQAHKRLGNRWAEIAKLLPGRTDNAVKNHWNSTMRR	120
MYBL1	187	KVEQEGYLQDGIK <b>ERSSS-----KLQHK-----PCATMDHLQTQNFY</b> ----	225
		KVEQEGY Q+ K+ S+ H P A L + +Y	
v-MYB	121	KVEQEGYPQESSK <b>AGPPSATTGFQKSSHLMAFAHNPPAGPLPAGAPLGSDYPPYHIAE</b>	180
MYBL1	226	---IPVQIPGYQYVSPDGNCVEHVQ-TSAFIQPFVDEDPDKEKKIKELELLLSAENEV	281
		+P QIP V+ N + Q +A IQ+ + DEDP+KEK+IKELELLLS ENE+	
v-MYB	181	<b>PQNVPVGGIP--Y</b> PVALHINIINVPQAAAAIQRHYTDEDPEKEKRIKELELLLSSTENEL	238
MYBL1	282	R-RKRLPPQ--PGSFSSWSGSFLMDDSMSNTLN-----NLEEHTTEFYSDENQTVSAQ	332
		+ ++ LP Q ++ W + + D++ ++ N H T +D + +	
v-MYB	239	KGQALPTQNHTANYPGWHSTTVADNTRTSGDNAPVSLGEGHHCTSPFPVDHG-CLPEE	297
MYBL1	333	QNSPTKFLAVEANAVLSSLQTIPEFAETLELIESDPVAWSDVTSFDLSDAAASPVKSTPV	392
		SP + + V + +L +++ + EFAETL+LI+S S+ + +L + A + STPV	
v-MYB	298	SASPARCMIVHQSNILDNVKNLLEFAETLQLIDSFLNTSSNHENLNLDNPA---LTSTPV	354