

# Supplementary Information

**Table S1.** Oligonucleotide primers used for normalized cDNA construction, PCR detection and gene expression analysis ( $N = A, C, G$ , or  $T$ ;  $K = A, G$ , or  $C$ ).

Primer name	Primer sequence (5' to 3')
SMART oligo IV	AAGCAGTGGTATCAACGCAGAGTGGCCATTACGGCCrGrGrG
CDS-3M adapter	AAGCAGTGGTATCAACGCAGAGTGGCCGAGGCGGCC(T)20 KN
CapM primer	AAGCAGTGGTATCAACGCAGAGT
M13	TGTAAAACGACGCCAGT
	CAGGAAACAGCTATGACC
<i>AsMDAS-box1</i>	GCTGCAGAGTAGTCATCAAGAGT
	GAUTGTGCAAGGTATCCATTG
<i>AsMDAS-box2</i>	GGTCGCCCTCATCATCTTCT
	GCCGGCTGTATCCCACCATT
<i>AsMDAS-box5</i>	GCTCAAGTACGAAGGAGATTATTG
	CACGACTTAATCCAGTTCAAGAG
<i>AsMDAS-box6</i>	GCCCTCGTCATCTCTCCAACC
	GCCGGCTGTATCCCACCATT
<i>18S rRNA</i>	GGCCTTCGGGATCGGAGTAAT
	CTAAGAACGCCATGCACAC
<i>AsKNOX</i>	CCGCCGACGACCTCCACAC
	GATCCCCACTTCGCCTTTA

**Figure S1.** Alignment of *MADS-box* amino acid sequences between *A. sisalana* and *A. tequilana*.

MADS-box_1.pro	MGRGRVELKRI ENKI NRAVTFAKDRNGLNNKKAYE LSVL CDAEVALVI FSNRSKLYFCCSS. SSMMKTLERYGKCSY GAP	79
AEX92976.pro	MGRGRVELKRI ENKI NRCVTFAKRRNGLLNKKAYE LSVL CDAEVALVI FSNRGKL YFCCSS. SSMMKTLERYGKCSY GAP	79
MADS-box_2.pro	MGRGRVELKRI ENKI NRCVTFAKRRNGLLNKKAYE LSVL CDAEVALVI FSNRGKL YFCCSS. SSMMKTLERYGKCSY GAP	79
AEX92975.pro	MGRGRVELKRI ENKI NRCVTFAKRRNGLLNKKAYE LSVL CDAEVALVI FSSRGRLL EFCCSS. SSMMKTLERYGKCSY GAP	79
MADS-box_6.pro	MGRGKIEI KKI ENPTNRCVTFYSKDRSGI MAAKAKAEATVL CDAETSIVMFSSACKFSEYCDPSTDTKSFDRYGCATG. . . . .	78
AEX92972.pro	MGRGKIEI KKI ENPTNRCVTFYSKRRSGI MAAKAKAEATVL CDAETSIVMFSSACKFSEYCDPSTDTKSFDRYGCATG. . . . .	78
MADS-box_5.pro	MA. REKI NR RKR DNTTARCVTFSKRRRGLFLKKAELSI CDAEVGLI FSATGKLF EFSSS.. STKEI EDHSMHSKKI L	77
AEX92969.pro	MA. REKI NR RKR DNTTARCVTFSKRRRGLFLKKAELSI CDAEVGLI FSATGKLF EFSSS.. STKEI ERHSMHSKKI L	77
 MADS-box_1.pro	DNSVCI RENQNLCSHSHQEYLKIKARVEALQRSQRNLLGNDLGPLSSKEI LEQLERQDSSUQI RSTRTQYWLDCGLADLQR	159
AEX92976.pro	DNSVCI RENQNLCSHSHQEYLKIKARVEALQRSQRNLLGNDLGPLSSKEI LEQLERQDSSUQI RSTRTQYWLDCGLADLQR	159
MADS-box_2.pro	KANASSKET. . . CNSNEEYLKIKARFELLQLSQRNLLGNDLGPLSSKEI LEQLERQDSSUQI RSTRTQYWLDCGLADLQR	156
AEX92975.pro	KANASSKET. . . CNSYEELYLKIKARFELLQLSQRNLLGNDLGPLSSNEI LEQESQIAMSUKI RSSKTCNVLGCCLCDLKR	156
MADS-box_6.pro	NLWTAQYEK. . . . NGNTLNHKEI NYNLRKEI RQRMCEEIDGNDVKDLRGEQCNDEALKLVRHRKYHVITCTETYKK	153
AEX92972.pro	NLWTAQYEK. . . . NGNTLNHKEI NYNLRKEI RQRMCEEIDGNDVKDLRGEQCNDEALKLVRHRKYHVITCTETYKK	153
MADS-box_5.pro	SPEQPSLDLN. . . LCNDNYARSKQVDETSRCLEKWRGEI CGQLTI ESLCNLESTI ETGUSDVLGRKSEQI MEQI NGLQQ	154
AEX92969.pro	SPEQPSLDLN. . . LCNSNYARSKQVDETSRCLRKWRGEI CGQLTI EELCNLEKT ETGUSRVLGRKSEQI MEQI NGLQQ	154
 MADS-box_1.pro	RECMILEANRSIRKRCVQLEETSGANCVWEANPN. . . . AMVYYSRPCN. CPQGDEFHPLECPPTLG. MGVC. . PDCN	230
AEX92976.pro	RECMILEANRSIRKRCVQLEETSGANCVWEANPN. . . . AMVYYSRPCN. CPQGDEFHPLECPPTLG. MGVC. . PDCN	230
MADS-box_2.pro	EECMIQDANRAIRSKLQEI GPE. NPLCLSVQNGGGGGGGCACTSAHCNRCPCPEGFFQPLGRDPSSQ. TGFSRVSMDH	234
AEX92975.pro	EECMIQDANRAIRSKLQEI GPE. NPLCLSVQNGGGGGGGCACTSAHCNRCPCPEGFFQPLGRDPSSQ. TGFSRVSMDH	234
MADS-box_6.pro	KFKNSCEAHRNLRELEMKDEHPAVYGFVDEDPSN. . . . YECALALAN. GGSCINYAFRVXPSCPMLHGWGYG. . . SQD	223
AEX92972.pro	KFKNSCEAHRNLRELEMKDEHPAVYGFVDEDPSN. . . . YECALALAN. GGSCINYAFRVXPSCPMLHGWGYG. . . SQD	223
MADS-box_5.pro	KGLCMEENTRIRCCVVDDNSQVGKQVVTGLENGSH. . . . EECQSSDSVTNASNSDAPCDYHDSSDT. LKLC. . LPWN	225
AEX92969.pro	KGLCMEENTRIRCCVVDDNSQVGKQVVTGLENGSH. . . . EECQSSDSVTNASNSDAPCDYHDSSDT. LKLC. . LPWN	225
 MADS-box_1.pro	AGPSVSAFNLC. . . . VIL	243
AEX92976.pro	AGPSVSAFNLG. . . . VIL	243
MADS-box_2.pro	LNSAVTNGNVNSFRHVM	251
AEX92975.pro	LNSAVTNGNVNSFRHVM	251
MADS-box_6.pro	LRLA. . . . .	227
AEX92972.pro	LRLA. . . . .	227
MADS-box_5.pro	LNTNF. . . . .	230
AEX92969.pro	LNTNF. . . . .	230