Table S3. Cloning primers

	Sequence (5′ to 3′)	
Construct	Forward	Reverse
pLMI2:LMI2	aaaaagcaggctGAAACGTGTCTCCACCCAAT	agaaagctgggttCTAGAATTTGGAAACCATGGA
LMI2:GUS*	GTCGACACTTGTAACTGTGCATGAAAC	CCCGGGGATTGTTCCTCACCCCACTAACA
LMI2:GUS†	GAGCTCGTCTTATGAGAGCCTAATATC	CAATTGAATTTTCTCAAGCATTGTCAC
LMI2 in situ	CTTTTATCTATGGGTCTTGATCCC	GAATGGTTAATTGTTTAATGTTCTGCAA
AP1 in situ	CGGAATTCCTTACGCCGAAAGACAGCTT	CGGGATCCCGTTCATTCTCTCTGACCTTCA
pLMI2:LMI2-HA	aaaaagcaggctGAAACGTGTCTCCACCCAAT	agaaagctgggttGAATTTGGAAACCATGGAAAC
GST-LFY	TAGAATTCATGGATCCTGAAGGTTTCAC	ATGCGGCCGCCTAGAAACGCAAGTCGTCGC
LMI2N	GAGGTCGACCATGGGAAGAACACCTTGTTG	TTAGCGGCCGCCTATGATTCTCTAGAAAGCCTTGC
LMI2C	GAGGTCGACCAGAGAATCAATGCTCTTTAGC	ATAGTTTAGCGGCCGCCTAGAATTTGGAAACCATGGA
LFY‡	aaaaagcaggctacATGGATCCTGAAGGTTTC	agaaagctgggttCTAGAAACGCAAGTCGTC
LFY§	CACCATGGATCCTGAAGGTTTCACGAG	GAAACGCAAGTCGTCGCC
2xmCherry§	CACCATGGTGAGCAAGGGCGAGGAG	CTTGTACAGCTCGTCCATGCCG

Lower case sequences are *att*B1 and *att*B2 sequence specific.
*Primers used to amplify *LMI2* upstream intergenic region.
†Primers used to amplify *LMI2* downstream intergenic region.
‡Primers used for yeast-two-hybrid constructs.
§Primers used for bimolecular fluorescence complementation (BiFC) constructs.