

-2323 TGCTGAGACAGCCAAAGGACGCCTTCTCCACCCAGTCCACAAGABRE**GTGGC**CCCTCAATTCCGGACCTACATT

-2253 AAAuxRE**CAGCAA**CAAAAGATGAGAATGCCATCCGGAAGAGTCAACAAAAAATCAACAAAGTAAAAACCGGACAA

-2183 TAGTCACAAAGCAAACCTACCCGGAACCACCAAGGAGTAGTTTGGAGAGAAAGGCCTCGCCGGATGCTCC

-2113 AAAAGCCLRE**CCGCCC**ATGCACCCCGGABRE**ACCACCACGTGTCCCGTCG**ABRECCCCTCCCTTTGTGCAATCTGGCAGCT

-2043 JARE JARE**CGTCACC**AGTTGGAGGGAGGGCAGGTGTGCGGACATGCTGAAGATGTCATTTTTGACCTTCTTCAAAGA

-1973 ATACACAAAAAACACCTCCAGCAGCGTGAGATCGAGGTTGTACAGCATGTTTATGATGCTGCATCCCATC

-1903 AGCACCCGGACGATGTGARE**TGGGAT**SURE**GAATAAA**GACGGGTGGAATCTGGGTGAAGTGGAGGAACTCCTTGAACA

-1833 ACGCCAGCAGAGGGAACCGGAGCCCCGCGTTGAATTGTTCCCTTCGAGAAGATGATAGTTTTTTCTTCTGC

-1763 TTTCTCAGAAGGAACAAGCACCCCTCCCATTAuxRE**CAGCAA**TCGATAGCCJARE**CGTCATTTGG**ABRE LREGATACAGAACCCT

-1693 TCCCAGAAATTCCTTCGCGTCCAACLRE**TTATCTAT**CGCTTTTCCACGAACCTCGGTACCCCGAGAAGATGAAA

-1623 CAGTCTTTTTTGCAGCCATTLRE**TCTTAC**CACAAACACAAGCCCCAAAACCTACACACAAGAAGCGCAATGGGTA

-1553 AAACACAAGTAACCCAGAAAATACACAAACCCTAACCCGAACAAATCAGAAACCAAACCAACAAAACGC

-1483 AAGARE**CAAAAGGAA**AuxRE**CAGCAA**TAGCAAAAGTAABRE**CACGTACC**LREABREAAATGAAGCTCTGAAGAAGAAAGGTTCGTCGTTTTC

-1413 GAATACAAGAACACCAGAuxRE**CAGCAA**ACAAATGTGCGGACACAAAATCACCGGTACAAAAGTGTTCGAGTT

-1343 TCTTTACTCAGAAAAAGCAAAGGACGCAACAAAAGCAAGAAGAAGAAAGGATCTCAGGAAAATGCAGTAG

-1273 AAAAAATGSARE**AAAAAGAAGG**TACAGTACGATATTTATAGGGGGACCAGCCCTCGGAAAGCCAAACGTCCAAA

-1203 CAGCGCTATCATTGAAACAACATGCTGCCTGGGAATCCCAGAGTCCCGGCTCATATAAATGCCTTTT

-1133 ATGGCTTCCGCACCCCTCTCGABRE LRE**CCACGTGGCTCAG**ABRELRETCAGACGAAGAGACCTCTTCAAATTCAAAAGCC

-1063 AGSURE**TTATTTT**LRE**TTAACCC**CGCCATTTTTTGGCAAAATAGGCAAGTTAAAAGGGGGCAATGTAGGGACCCC

LRE ABRE LRE LRE ABRE
-0993 TCCCTCTGGGAAA**CACGTGGCAC**GCACCTCACAGTGACACGCAG**CACGTGTTAT**CAGCCGGACCATCATC
-0923 ATCCGGATTCCCTTAAGGATACGCATGATGATGGTTCLRE**TCCTATC**CGGACCGCCTCAAGGAAAAGCACATLRE**TG**
-0853 JARE**ACG**TTTTTTCAGCTTCTTCTGTCCAAGGAAGAGCAAAuxRE**ACGAC**GCTGACAGAGCATAGACATCCGGACAACCTTC
-0783 ATAATGTATCTGCTCCACTATAACAATCCGGATAGTCAGCATGTGACCATCCGGATTTAATCGTCCGGATC
-0713 ATCAATTAAGTAAAGCAAGLRE**TCTTAC**ACGCTATLRE**CACGAC**AACCAGCCATGGCCCACATCCCATCATCTGC
-0643 AGAGTGAAAGGACGGGTCGAGGTGACAACAAGTCACTTCCCACGATCATTCTACATGATCATTTCACG
-0573 ATATCTAGACAGCAGCATCACCTACCACGGTTTCTGACAERE**GCCGCC**AGTAGGABRE**GTGGC**GATGACCATGCTGC
-0503 CTCCGAATGTCATCATGACAAACATAAAATATCTCCTCGCCLRE**ATTAAT**GAGAGGAACAGTACCCCTGAAGC
-0433 TGTATATATATGCCTTCGCACGAAGAAGAAGGGGATCCTCCTGGTAACTTCTTAATACCTGGTAAAAGGC
-0363 CAACTAATTTATATTCTCTCTAACCATGGCTAACAAAACCATCGGAGGATGCGTCCAGACACCCTGT
-0293 CCGGATGCCTTCTTGCAGGAAJARE**TGACG**ACTGGATCAAAAACCTTTATGAGTTGAGATCACGCGTCCATCCA
-0223 TCTGGTTACABRE/LRE**TACGTG**GACCGCCAAAGACGCGAGGTAACAACAACACACCCTTTGTCCATGAACTCCAG
-0153 CGCLRE**ATTTGG**AAGCCAGTAATGCACCATAAABRE LRE SURE**GAAACGTGTCGAATAAA**CCAATTAGGGGTCTGGTGTCCGAG
-0083 TCATGAGATAGAACAGGTTTCGAGGTTGTTATA Box**TATA**TATCAATCAATAATTAGAGAAGGAGCCGGTCTCTTTGTG
-0013 TTGAGTTGACTCG

Supplementary Figure 4. Different cis-elements detected in the *MYBA1* promoter region from -2323 to +1. Sequence was analyzed using the PlantCARE website (<http://bioinformatics.psb.ugent.be/webtools/plantcare/html/>) (Lescot et al., 2002) and PLACE website (<https://www.dna.affrc.go.jp/PLACE/?action=newplace>) (Higo et al., 1998). Number of lines indicates the nucleotide location relative to the start codon. Cis-elements sequence are indicated in bold letters and abbreviations of the different responsive elements are ABA (ABRE), ethylene (ERE), light (LRE), sugar (SURE), methyl jasmonate (JARE), gibberellic acid (GARE), auxin (AuxRE) and salicylic acid (SARE).