

Figure S1

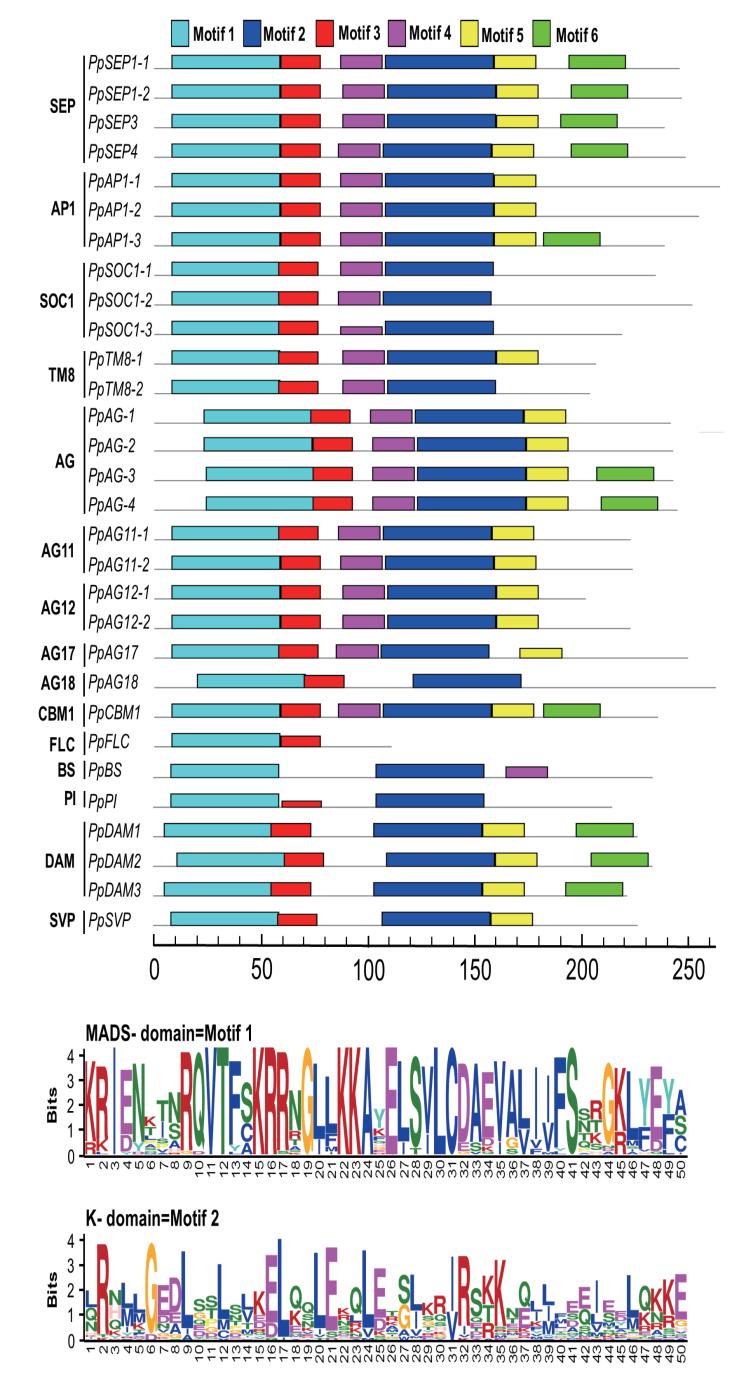


Figure S2

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- GAATTTTGTT TGGTTGCGGA TTTGTTTTAG TTGTAAGTCA GCGTACTTCT CTAAAGTCTC TTCTCCTTCC
  + GAGGAATAGA AGTACCGACC ATAATTTATT TATTTATTAT AAATAAATAA ATAAATAAA GCATATGAAG
- CTCCTTATCT TCATGGCTGG TATTAAATAA ATAAATAATA TTTATTTATT TATTTATTTT CGTATACTTC
  + AGATTGCACT GCAGGGAAGA CGAGAGGGCC GGCCCGGGATG GCAACGGTGG AAGAGGGTTT GGCCGACGGT
- TCTAACGTGA CGTCCCTTCT GCTCTCCCGG CCGCCCTAC CGTTGCCACC TTCTCCCAAA CCGGCTGCCA
  + GGATGGCCGG TGTTGCAGCG ATTGCTGCTA CCGCTGCTCT TTAGGATAAA GAAGTGAGAG AGACGAGGTG
- CCTACCGGCC ACAACGTCGC TAACGACGAT GGCGACGAGA AATCCTATTT CTTCACTCTC TCTGCTCCAC
  + GATGETAC A CACCTAAAAA GAAATAAACC GAGAATCCCC ACTCAGTTAG ATATGAACAA ATCTAAGAGA
    CTACACTGCT GTGGATTTTT CTTTATTTGG CTCTTAGGGG TGAGTCAATC TATACTTGTT TAGATTCTCT
  + TTCCATATTT TTACTGTCGT CCATTCCACA TATATTATGA ACGGTGGTGA TCGCATTCAT CACTCATCAC
- AAGGTATAAA AATGACAGCA GGTAAGGTGT ATATAATACT TGCCACCACT AGCGTAAGTA GTGAGTAGTG
   + GCCAACTGAT TTGCGTTCCA CGTTTCTTAC CTTAAAAATA TCCAAAAATA GGAAAGAGAA ACAAACAGTT
  - CGGTTGACTA AACGCAAGGT CAAAGAATG GAATTTTTAT AGGTTTTTAT CCTTTCTCTT TGTTTGTCAA
   + GATATTAAAG CGAGATATAT CTTATTTTC TATAAATAGG AAATTTAATT CGCCAACCTA TCCTTTCTAG
  - CTATAATTTC GCTCTATATA GAATAAAAAG ATATTTATCC TTTAAATTAA GCGGTTGGAT AGGAAAGATC
  + AACAACCGGC AGCACCCGAA CTCTCTCTCT CCTCTCTGTT CTTCTCTTTC CAAAATATCT TTGGTTTCCT
  - TTGTTGGCCG TCGTGGGCT<mark>T GAGAGA</mark>GAGA GGAGAGACAA GAAGAGAAAG GTT<mark>TTATAGA AAC</mark>CAAAGGA
  + TTCACTGCTC TGCAAATTTC CCAAAACTTC AGACCTGAAA CTCCCCGACG AAGGTAAACA AATAACAACG
   - AAGTGACGAG ACGTTTAAAG GGTTTTGAAG TCTGGACTTT GAGGGGCTGC TTCCATTTGT TTATTGTTGC
  + GAAAACCCAG TAAATAATTT ATGTATGGTT TTTTTTTTC TTCTTCTCTT CGATTGGTGG TAAATACAAA
     CTTTTGGGTC ATTTATTAAA TACATACCAA AAAAAAAAA AA AAGAAGAGAA GCTAACCACC ATTTATGTT
    + ATACTTTAGG CTTGTTGGGC AGTTGTTTGT GGTTGGTTGG TTTTTCTGGG TTTTGATTTT CTTATTTAGA
                   AAATC GAACAACCG TCAACAACA CCAACCAACC AAAAGACCC AAAACTAAAA GAATAAATCT
  + TCTGAGATGC TAGGGTTTGA GGT<mark>CAAT</mark>ATC TTTAAACTTA TAAAAATACC CAGAAAATAT ACGGTAGTTT
   - AGACTCTACG ATCCCAAACT CCAGTTATAG AAATTTGAAT ATTTTTATGG GTCTTTTATA TGCCATCAAA
  + TTGAGTTTGT GAACAGTTAT ATATGTGAAA AGTGCATGGT TGGTTTATTT TTATTTATTT GAATTAATTG
- AACTCAAACA CTTGTCAATA TATACACTTT TCACGTAC<mark>CA ACCAAATAAA</mark> AATAAATAAA CTTAATTAAC
   + TTTTTGAAGG GGTAGGAAGA
  - AAAAACTTCC CCATCCTTCT
  + 5UTR Py-rich stretch
  + AAGAA-motif
+ ACE

        Organism
        Position Strain
        Matrix score.
        sequence

        Petroselinum hortense
        437
        -
        7
        ACGTGGA

        Petroselinum crispum
        770
        -
        9
        CTAACGTATT

                                                                                                                                                                                     cis-acting element involved in light responsiveness cis-acting element involved in light responsiveness
 + AE-box
                                               Organism Position Strand Matrix score.

Arabidopsis thaliana 477 + 8 AGAAACAA
                                                                                                                                                                                                      part of a module for light response
 + ARE
                                                                                          Position Strand Matrix sequence score.
             Site Name
                                                          Organism
                                                                                                                                                                                                                             function
                                                           Zea mays
Zea mays
Zea mays
Zea mays
Zea mays
                                                                                                                                              TGGTTT
TGGTTT
TGGTTT
TGGTTT
TGGTTT
                                                                                                                                                                         cis-acting regulatory element essential for the anaerobic induction cis-acting regulatory element essential for the anaerobic induction
  ARE
ARE
ARE
ARE
 + Box 4
                                               Organism Position Strand Matrix sequence
Fetroselinum crispum 973 + 6 ATTAAT
                                                                                                                                                                                                                            function
                                                                                                                                                                          part of a conserved DNA module involved in light responsiveness
  Вож 4
 + Box-W1
                                              Organism Position Strand Matrix sequence score. Fetroselinum crispum 861 - 6 TTGACC
           Site Name
  Box-W1
                                                                                                                                                                                                     fungal elicitor responsive element
 + C-repeat/DRE
                                               Organism Position Strand Matrix sequence arabidopsis thaliana 200 + 8 TOGCCGAC
   C-repeat/DRE
                                                                                                                                                                        regulatory element involved in cold- and dehydration-responsiveness
+ CAAT-box
                                                                                         Position Strand Matrix sequence
           Site Name
                                                      Organism
                                                                                                                                                                                common cis-acting element in promoter and enhancer regions common cis-acting element in promoter and enhancer regions
                                                Hordeum vulgare
Arabidopsis thaliana
Brassica rapa
Hordeum vulgare
Brassica rapa
Brassica rapa
Brassica rapa
Hordeum vulgare
Hordeum vulgare
Glycine max
                                                                                                                                              CAAT
CCAAT
CAAAT
CAAAT
CAAAT
CAAAT
CAAAT
CAAAT
CAAT
CAAT
CAAT
CAAT
CAAT
                                                                                              143
753
643
977
338
967
689
864
 CAAT-box
CAAT-box
  CAAT-box
+ CATT-motif
                                                      Organism Position Strand Matrix score.
           Site Name
                                                                                                                                                                                                                           function
                                                         Zea mays
                                                                                                                                                                                                    part of a light responsive element
 CATT-motif
+ CCAAT-box
                                                                                        Position Strand Matrix sequence
         Site Name
                                                      Organism
                                                                                                                                                                                                                          function
+ CGTCA-motif
                                                                                         Position Strand Matrix sequence
        Site Name
                                                       Organism
                                                                                                                                                                                                                           function
                                                                                                                                              CGTCA
                                                    Hordeum vulgare
                                                                                             285
                                                                                                                                                                        cis-acting regulatory element involved in the MeJA-responsiveness
 CGTCA-motif
+ G-Box
                                                                                        Position Strand Matrix score. sequence
                                                      Organism
                                                     Pisum sativum
                                                                                            439
                                                                                                                                            CACGTT
                                                                                                                                                                          cis-acting regulatory element involved in light responsiveness
+ G-box
                                                        Organism
                                                          Zea mays
                                                                                                                                                                            cis-acting regulatory element involved in light responsiveness
+ GAG-motif
                                                                                      Position Strand Matrix score.

580 - 7 AGAGAGT
                                                       Organism
                                               Arabidopsis thaliana 580 -
+ L-box
                                                                                Position Strand Matrix sequence serum 949 - 11 AAATTAACCAAC
                                            Lycopersicon esculentum 949
                                                                                                                                                                                                        part of a light responsive element
+ MBS
                                                         Organism
                                                                                        Position Strand Matrix sequence
                                               Arabidopsis thaliana
Arabidopsis thaliana
Arabidopsis thaliana
Arabidopsis thaliana
Arabidopsis thaliana
                                                                                                                                                                                        MYB binding site involved in drought-inducibility
+ 02-site
                                                        Organism Position Strand Matrix score.

Zea mays 417 - 9 GTTGACGTGA
          Site Name
                                                                                                                                                                                                                            function
+ sp1
                                                                                        Position Strand Matrix sequence score.
    Site Name
                                                         Organism
                                                                                                                                                                                                                            function
                                                         Zea mays
                                                                                                                                                                                                               light responsive element
 Sp1
+ TATA-box
                   TCA-element

        Organism
        Position Strand Score.
        Matrix sequence score.

        dictional tabacum
        353
        +
        9
        CCAICTITIT

        reassica cleracea
        739
        -
        9
        GAGAAGATA

        ressica cleracea
        597
        -
        9
        GAGAAGATA

                                                                                                                                                                                                                           function
                                                                                                                                                                   cis-acting element involved in salicylic acid responsiveness
cis-acting element involved in salicylic acid responsiveness
cis-acting element involved in salicylic acid responsiveness
 + TCCACCT-motif
               TCT-motif

ite Name Organism Position Strand Matrix sequence score, sequence score, for the format of the format o
      Site Name Organism Position Summer Position Su
                                                                                                                                                                                                                            function
                                                                                                                                                                                                    part of a light responsive element
 TCT-motif
+ TGACG-motif
                                                  Organism Position Strand Strand Store: function score: function score: function score:
 TGACG-motif
+ Unnamed_1 + Unnamed_11
                  Unnamed__11
                  Unnamed__2
   + Unnamed_3
+ Unnamed_4
    + W box
                                              Organism Position Strand Matrix sequence score.

Brassica oleracea 528 - 9 AAATITCCT
+ circadian
                                                                                    Position Strand Matrix sequence function

um 614 - 9 CAAAGATATC cis-acting regulatory element involved in circadian control
        Site Name
                                                     Organism
                                          Lycopersicon esculentum 614 -
```

```
- TAAATAAATT TAATTATTTT AGTATACTTC TATAACGAAA CGTCTCTTCT GCTTCCCCGG CCCTACCGTT
+ TGGTGGAAGA GGGTTTGGAG AATGCGGCTA CTGAAACTCG GGGGTTGGGG TTTGGGGAGT TTTGTTTCAC
- ACCACCTTCT CCCAAACCTC TTACCCCGAT GACTTTGAGC CCCCAACCCC AAACCCCTCA AAACAAAGTG
+ AAGTGGGAGT GAGTATTGCC AGCAAGTGTT GCAGCGATCG CTGCTACCGC TGCTGTTTGG GATAAAGAAG
- TTCACCCTCA CTCATAACGG TCGTTCACAA CGTCGCTAGC GACGATGGCG ACGACAAACC CTATTTCTTC
+ TGAGAGAGAC GAAGTGGGTG AGGAAAAT<mark>CA ACGG</mark>TGTGGT ACGTAAAAAG AATTAAACGT AGAATCCCCA
- ACTCTCTCTG CTTCACCCAC TCCTTTTAGT TGC ACACCA TGCATTTTTC TTAATTTGCA TCTTAGGGGT
+ GCCTTACTGT CGTCCAGTCC ACATATTATA GACGGTCGTG ATCGCATTCA TGACTCATCA CACCCAACTGA
- CGGAATGACA GCAGGTCAGG TGTATAATAT CTGCCAGCAC TAGCGTAAGT ACTGAGTAGT GTGGTTGACT
+ TTTGTGTTCC GCGTTTCTAA CCTTATGTTC CCAAAAAATA TCTAAAAAGA CAAAAAGGCA CAAACAGTGA
- AAACACAAGG CGCAAAGATT GGAATACAAG GGTTTTTTAT AGATTTTTCT GTTTTTCCGT GTTTGTCACT
+ TAACCAAAGG AAGATATGTC CTATTTTCC ATAATTAGGA AAATTAATTT AGCAACCTTT CCTTTCTTGA
- ATTGGTTTCC TTCTATACAG GATAAAAAGG TATTAATCCT TTTAATTAAA TCGTTGGAAA GGAAAGAACT
+ ACTGCCGGCA GCATCCCAAC TCTTTCTCAT CTCTTCTT CTCTTTCCAG AATTGCTGGC TTTTCTTTCG
 TGACGGCCGT CGTAGGGTTG AGAAAGAGTA GAGAGAAGAA GAGAAAGGTC TTAACGACCG AAAAGAAAGC
+ CTGTTCTGCA AATTTC<mark>CCGA AA</mark>CTTCAGAG ACCTAAAA<mark>CC CCCCAACG</mark>AA GGTAAACAAG TAA<mark>CAACGG</mark>A
 GACAAGACGT TTAAAGGGCT TTGAAGTCTC TGGATTTTGG GGGGTTGCTT CCATTTGTTC ATTGTTGCCT
+ AAACCCAGTG TAATTAATTT ATGTATGGTC TTCCTCCTCC TCCCCCTTCT TCTTCTTCTT ATGGTAGATA
 TTTGGGTCAC ATTAATTAAA TACATACCAG AAGGAGGAGG AGGGGGAAGA AGAAGAAGAA TACCATCTAT
+ \hspace{1.5cm} \texttt{CTGTAGGCTT} \hspace{1.5cm} \textbf{GTTGGACAGT} \hspace{1.5cm} \textbf{TGATTGTGGT} \hspace{1.5cm} \textbf{GGGTTTTTCT} \hspace{1.5cm} \textbf{GGGGTTTGTT} \hspace{1.5cm} \textbf{TTTCTTTATA} \hspace{1.5cm} \textbf{GATCTGAGAT} \\
- GACATCCGAA CAACCT<mark>GTCA AC</mark>TAACACCA CCCAAAAAGA CCCCAAACAA AAAGAAATAT CTAGACTCTA
+ GCTAGGGTTT GAGGGCAATA ATCTCTTATT TAGATCTGAG ATTCTAGGGT TTGAGGTCAA GAATCTTTAA
- CGATCCCAAA CTCCCGTTAT TAGAGAATAA ATCTAGACTC TAAGATCCCA AACTCCAGTT CTTAGAAATT
+ AATTATAAAT AAAAAAAAA ATATGCTAGT TTTTGAGTTT GTGGACGATT ATATATATGT GAAAAGTGGT
- TTAATATTTA TTTTTTTTA TATACGATCA AAAACTCAAA CACCTGCTAA TATATATACA CTTTTCACCA
+ TGTTTTATTT TATTTATTTG AATTAATTGT TTTTGCAGGG GTAGCAAGAA ATGGTGGAAA GGATGAAAGA
- ACAAAATAAA ATAAATAAAC TTAATTAACA AAAACGTCCC CATCGTTCTT TACCACCTTT CCTACTTCT
+ GAAGATAAAG ATCAGGAGGA
- CTTCTATTTC TAGTCCTCCT
+ 5UTR Py-rich stretch
       AAGAA-motif
+ Box 4
                         Organism Position Strand \frac{Matrix}{score}.

        Petroselinum crispum
        11
        +
        6
        ATTAAT

        Petroselinum crispum
        643
        +
        6
        ATTAAT

        Petroselinum crispum
        463
        +
        6
        ATTAAT

                                                                                part of a conserved DNA module involved in light responsiveness
part of a conserved DNA module involved in light responsiveness
part of a conserved DNA module involved in light responsiveness
                      Petroselinum crispum
                                                                   ATTAAT
                                                                                 part of a conserved DNA module involved in light responsiveness
+ Box-W1
                     Organism Position Strand Matrix sequence score.
   Site Name Organism Posici...
Petroselinum crispum 825 -
                                                                                                        function
                                                                  TTGACC
                                                                                              fungal elicitor responsive element
+ CAAT-box
+ CATT-motif
   Site Name Organism Position Strand Matrix sequence
CATT-motif
                          Zea mays
Zea mays
                                                                  GCATTC
                                                                                              part of a light responsive element part of a light responsive element
+ CCAAT-box
                                          Position Strand Matrix sequence score.
   Site Name Organism
                                                                                                        function
                                          239 + 6
624 + 6
                                                                                                    MYBHv1 binding site
                       Hordeum vulgare
                                                                  CAACGG
CCAAT-box
                       Hordeum vulgare
+ G-box
                                         Position Strand Matrix sequence score.
                         Organism
   Site Name
                                                                                                        function
                           Zea mays
                                                                  CACGAC
                                                                                 cis-acting regulatory element involved in light responsiveness
+ GA-motif
                                        Position Strand Matrix sequence score.
                         Organism
Glycine max
                                           427 +
GA-motif
                                                                 AAGGAAGA
                                                                                              part of a light responsive element
+ GAG-motif
                                         Position Strand Matrix sequence score.
                        Organism
                                                                                                        function
                                                                                              part of a light responsive element
+ GC-motif
                                         Position Strand Matrix score.

109 - 6 CCCCCG
                         Organism
                         Zea mays
                                                                                 enhancer-like element involved in anoxic specific inducibility
GC-motif
+ GCN4_motif
                                        Position Strand Matrix sequence
   Site Name Organism
                                                                                                       function
                       Oryza sativa 331 -
                                                                 TGAGTCA
GCN4_motif
                                                                                   cis-regulatory element involved in endosperm expression
+ HSE
                                        Position Strand Matrix sequence score.
  Site Name
                         Organism
                                                                                                        function
HSE
+ LTR
                                         Position Strand Matrix sequence score.
                         Organism
  Site Name
                                          577 +
                                                                  CCGAAA
                        Hordeum vulgare
                                                                                 cis-acting element involved in low-temperature responsiveness
LTR
+ MBS
                         Organism Position Strand Matrix sequence score.
                     Arabidopsis thaliana 344 + 6 CAACTG
Arabidopsis thaliana 717 - 6 CAACTG
                                                                                      MYB binding site involved in drought-inducibility MYB binding site involved in drought-inducibility
+ Skn-1_motif
  Site Name Organism Position Strand Matrix sequence score.
                       Oryza sativa 330 -
Skn-1 motif
                                                                  GTCAT
                                                                              cis-acting regulatory element required for endosperm expression
+ Sp1

        Organism
        Position Strand score:
        Matrix sequence score:

        Zea mays
        599
        +
        5
        CC(6/A) CCC

        Zea mays
        669
        +
        5
        CC(6/A) CCC

  Site Name
                       Organism
                                                                                                       function
                                                                                                  light responsive element
+ TATA-box
+ TATC-box
                      Organism Position Strand \frac{\text{Matrix}}{\text{score}} sequence Oryza sativa 198 - 7 TAICCCA
Site Name
                                                                                                       function
                                                                                 cis-acting element involved in gibberellin-responsiveness
+ TC-rich repeats
                        Organism Position Strand Matrix sequence function

Nicotiana tabacum 229 - 9 ATTTCTTCA cis-acting element involved in defense and stress responsiveness
  Site Name
                      Nicotiana tabacum 229 -
+ TCA-element
                     Organism Position Strand score. sequence function

Brassica oleracea 524 - 9 GAGAAGAATA cis-acting element involved in salicylic acid responsiveness
TCA-element
 + Unnamed_4
       W box
  + chs-Unit 1 m1
   Site Name Organism Position Strand Matrix sequence score.

-Unit 1 ml Arabidopsis thaliana 244 - 10 ACCTACCACAC
chs-Unit 1 m1
                                                                                            part of a light responsive element
```

+ ATTTATTTAA ATTAATAAAA TCATATGAAG ATATTGCTTT GCAGAGAAGA CGAAGGGGCC GGGATGGCAA

```
+ CAGGTGTGGC CGACTCCACG TGAACTCGGA TAAACACCGT ACCGTATAAC TGAAAAACCG GTTCGGTTTT
- GTCCACACCG GCTGAGGTGC ACTTGAGCCT ATTTGTGCA TGGCATATTG ACCTTTTGGC CAAGCCAAAA
+ GCACCGTACC AAAGTAAATT AAGAAAAAAA AAACAAACCA AACCGATCAG TTATTTTCCG GTTCGATACT
- CGTGGCATGG TTTCATTTAA TTCTTTTTT TTTG<mark>TTTGGT T</mark>TTGGCTA<mark>GTC AAT</mark>AAAAGGC CAAGCTATGA
+ ATATCTCCAT TCCCTCCACC GCCCCATTTC CTTTCCTTCT ATCAGCCACC TGTTACGTGT CAACTGCACA
- TATAGAGGTA AGGGAGGT<mark>GG CGGG</mark>GTAAAG GAAAGGAAGA TAGTCGGTGG ACA<mark>ATGCAC</mark>A GTTGACGTGT
+ AGCATCGACT GAGGAAGAAG CAGCTTTAGC AACTAGCAAG TGATGCTAGC ATCTCTCAAA AAAGTGGGAC
- TCGTAGCTGA CTCCTTCTTC GTCGAAATCG TTGATCGTTC ACTACGATCG TAGAGAGTTT TTTCACCCTG
+ ACGTGAAAAG AAATAGGCTA AACCGACAAT AAACGGTGGG ATAGCATTCA TCACACCAAC CGCATCGATC
- TGCACTTTTC TTTATCCGAT TTGGCTGTTA TTTGCCACCC TATCGTAAGT AGTGTGGTTG GCGTAGCTAG
+ ACCGTTGATT ATCGTTCCAT GTTTCTTACC TTATTTGCCC AACAAAATAT CCAAAATATC TAAAAATAGA
- T<mark>GGCAACTAA TAGCA</mark>AGG<mark>TA CAAAGA</mark>ATGG AATAAACGGG TTGTTTTATA GGTTTTATAG ATTTTTATCT
+ AAAAGCAACA AACAGCGGAT ATTAATAAAA GATTTATCCA AGGTCTTCCA TATATGGGAA AATTCATTTA
- TTTTCGTTGT TTGTCGCCTA TAATTATTTT CTAAATAGGT TCCAGAAGGT ATATACCCTT TTAAGTAAAT
+ GCGACCTTTC CTTTCTTGAA CAGCCTGCAG CACCAACTCT CTCTCTCCC TCAGTCTTCT CCTCTTTCCA
- CGCTGGAAAG GAAAGAACTT GTCGGACGTC GTGGTTGAGA GAGAGAGAG AGTCAGAAGA GGAGAAAGGT
+ AAATTGCTCC GAAAATTTCC AGTTCAGACC TAAACTTCAG ACCTAAGACC CCTCAACGAA GGTAAACAAG
  TTTAACGAGG CTTTTAAAGG TCAAGTCTGG ATTTGAAGTC TGGATTCTGG GGAGTTGCTT CCATTTGTTC
+ AGAACCAGAG GTGAACCCAG TACTTAGTTT ATCTATGTTT TTCCTTCTT TTTTTTCTTT TTTCCTTCTA
- TCTTGGTCTC CACTTGGGTC ATGAATCAAA TAGATACAAA AAGGAAGAAA AAAAAAGAAA AAAGGAAGAT
+ TTAATGGTGA ATACTAAAAA CTCTAGGCTT GTCTGGCAGT TGATTGTGGT TGGTGGGTTG ATCTGTGTTT
- AATTACCACT TATGATTTTT GAGATCCGAA CAGACC<mark>GTCA AC</mark>TAACACCA ACCACCCAAC TAGACACAAA
+ TATTTTACTT ATTTAGATCT GAGATATGTT AGGGTATGAG GTCAATAATC TTAAAATTAT CAGAAACCCA
- ATAAAATGAA TAAATCTAGA CTCTATACAA TCCCATACTC CAGTTATTAG AATTTTAATA GTCTTTGGGT
+ GAAAATATAC TGTAGTTTGT GAAACATTTA TATATATATG TGGAAATTTA AAGTGAT<mark>TGG TTT</mark>ATTTTAT
 CTTTTATATG ACATCAAACA CTTTGTAAAT ATATATATC ACCTTTAAAT TTCACTAACC AAATAAAATA
+ TTTATTTGAA TTATTGTTTT
- AAATAAACTT AATAACAAAA
+ 4cl-CMA2b
                      Organism Position Strand Matrix sequence
Petroselinum crispum 401 + 10 TCTCACCAACC
                                                                                                                    function
4c1-CMA2b
                                                                                                              light responsive element
+ AAGAA-motif
+ ABRE
                           Organism Position Strand \frac{Matrix}{score}.
   Site Name
                     Arabidopsis thaliana 87 +
Arabidopsis thaliana 350 +
Arabidopsis thaliana 264 +
                                                                                          cis-acting element involved in the abscisic acid responsiveness cis-acting element involved in the abscisic acid responsiveness cis-acting element involved in the abscisic acid responsiveness
                                                                          CACGTG
                                                                           CACGTG
                                                                           TACGTG
+ AC-I
 + ACE
                           Organism Position Strand \frac{Matrix}{score}.
                                                                                                                   function
     Site Name
                       Petroselinum hortense
                                               85 -
                                                                                               cis-acting element involved in light responsiveness
                                                                         ACGTGGA
+ AE-box
                                           Position Strand Matrix sequence score.
   Site Name
                            Organism
                                                                                                                   function
                       Arabidopsis thaliana 439 -
                                                                                                       part of a module for light response
AE-box
+ ARE

        Position Strand
        Matrix score.
        sequence

        175
        -
        6
        TGGTTT

        968
        +
        6
        TGGTTT

     Site Name
                            Organism
                                                                                                                   function
                                                                                         cis-acting regulatory element essential for the anaerobic induction cis-acting regulatory element essential for the anaerobic induction
+ Box 4
                           Organism Position Strand Matrix sequence core: sequence stroselinum crispum 770 + 6 ATTAAT attactions attack.
                                                                                                                   function
                        Petroselinum crispum
Petroselinum crispum
                                                 511 +
770 +
                                                                                          part of a conserved DNA module involved in light responsiveness part of a conserved DNA module involved in light responsiveness
+ Box-W1
                           Organism Position Strand Matrix score.
                                                                                                                   function
                      Petroselinum crispum 880 -
Box-W1
                                                                                                        fungal elicitor responsive element
+ C-repeat/DRE
                                            Position Strand Matrix sequence score. 8 TGGCCGAC
                            Organism
    Site Name
                                                                                                                   function
C-repeat/DRE
                        Arabidopsis thaliana
                                                                                        regulatory element involved in cold- and dehydration-responsiveness
+ CAAT-box
+ CATT-motif
                                              Position Strand Matrix sequence
                           Organism
   Site Name
                                                                                                                    function
                            Zea mays
CATT-motif
                                               394 +
                                                                          GCATTC
                                                                                                        part of a light responsive element
+ CCAAT-box
                                              Position Strand Matrix sequence
    Site Name
                            Organism
                                                                                                                   function
                         Hordeum vulgare
                                                 422 -
                                                                          CAACGG
                                                                                                               MYBHv1 binding site
CCAAT-box
+ G-Box
                                              Position Strand Matrix sequence
                            Organism
                                                                                                                   function
                                                                          CACGTG
CACGTG
CACGTA
                          Pisum sativum
Pisum sativum
Antirrhinum majus
                                                                                          cis-acting regulatory element involved in light responsiveness cis-acting regulatory element involved in light responsiveness cis-acting regulatory element involved in light responsiveness
G-Box
G-Box
+ G-box
                                              Position Strand Matrix sequence score.
                                                                                           cis-acting regulatory element involved in light responsiveness cis-acting regulatory element involved in light responsiveness cis-acting regulatory element involved in light responsiveness cis-acting regulatory element involved in light responsiveness
                           Brassica napus
                                               86 – 7
264 + 6
                       Daucus carota
Arabidopsis thaliana
Arabidopsis thaliana
G-box
G-box
                                                                           CACGTG
G-box
                                                                           CACGTG
+ GA-motif
                                           Position Strand Matrix sequence
score. sequence
                            Organism
                                                                                                                   function
                      Arabidopsis thaliana 729 -
GA-motif
                                                                                                        part of a light responsive element
+ GAG-motif
                                              Position Strand Matrix score.
  Site Name
                             Organism
                                                                                                                    function
                                                                        AGAGATG
                                                                                                        part of a light responsive element part of a light responsive element
                          Spinacia oleracea
GAG-motif
                         Arabidopsis thaliana
                                                                          AGAGAGT
+ GT1-motif
                             Organism Position Strand \frac{Matrix}{score}, sequence
   Site Name
                                                                                                                    function
                         Solanum tuberosum
                                              813 +
                                                                        ATGGTGGTTGG
                                                                                                             light responsive element
+ I-box
                                            Position Strand Matrix sequence
                                                                                                                    function
                            Organism
                                                                         ACGATAATC
                                                                                                        part of a light responsive element
                      Arabidopsis thaliana 427 -
+ L-box
                                              Position Strand Matrix score.
401 + 10 TCTCACCAACC
                                                                                                                    function
   Site Name
                            Organism
                      Petroselinum crispum 401 +
+
                                              Position Strand Matrix sequence score.
     Site Name
                             Organism
                          Hordeum vulgare
+ MBS
                                              Position Strand Matrix sequence score.
                         Arabidopsis thaliana
Arabidopsis thaliana
Arabidopsis thaliana
                                               117 + 6
271 + 6
188 - 6
807 - 6
                                                                           TAACTG
CAACTG
TAACTG
                                                                                                 MYB binding site involved in drought-inducibility
MBS
                         Arabidopsis thaliana
                                                                           CAACTG
+ Sp1
                                              Position Strand \frac{\text{Matrix}}{\text{score.}} sequence
                             Organism
                                                                                                                   function
                              Zea mays
                                                                    5 CC (G/A) CCC
                                                                                                             light responsive element
light responsive element
light responsive element
                            Oryza sativa
Zea mays
                                                                   5.5 CC (G/A) CCC
                                                  42
                               Zea mays
Zea mays
                                                                                                             light responsive element
light responsive element
                                                                         CC (G/A) CCC
+ TATA-box
+ TATC-box
                           Organism Position Strand Matrix sequence score.

Oryza sativa 387 - 7 TAICCCA
                                                                                                                    function
TATC-box
                           Oryza sativa
                                                                                             cis-acting element involved in gibberellin-responsiveness
+ TCA-element
                          t
Organism Position Strand Matrix sequence score.
                                                                                          cis-acting element involved in salicylic acid responsiveness
TCA-element
+ TCT-motif
                           Organism Position Strand \frac{Matrix}{score}.
                                                                                                                    function
                                               444 +
TCT-motif
                       Arabidopsis thaliana
                                                                                                       part of a light responsive element
 + Unnamed_1
 +
         Unnamed__11
         Unnamed 3
 +
 + Unname
          Unnamed 4
    box S
 + circadian
                            Organism Position Strand \frac{Matrix}{score}, sequence
                                                                                                                   function
                      Lycopersicon esculentum 62 +
Lycopersicon esculentum 407 +
Lycopersicon esculentum 179 +
                                                                6 CAANNNATC
6 CAANNNATC
6 CAANNNATC
                                                                                           cis-acting regulatory element involved in circadian control
circadian
                                                                                            cis-acting regulatory element involved in circadian control cis-acting regulatory element involved in circadian control
```

+ AATAGTGAGT CTGAGAGAGA TCGAGTAGTT GAAGAGAGGG GGGGTGGGGG GCAGTGGAAG TCAAATCCAT - TTATCACTCA GACTCTCTT AGCTCATCAA CTTCTCTCCC CCCCACCCC CGTCACCTTC AGTTTAGGTA

Lycopersicon esculentum

circadian

B-type (upstream of *PpDAM2=ProDAM2*):

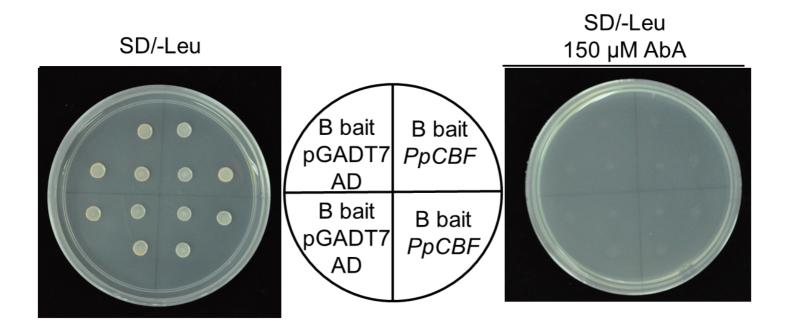


Figure S6