

Table S1. List of primers.

| Primer name | Primer sequences (5'-3') |
|--------------------------|---|
| <i>CpBBX19-F</i> | GAAGGGAAATGAAAGGTGG |
| <i>CpBBX19-R</i> | CTTCTTTAAGCCGCCAATAGTC |
| <i>CpBBX19-F1</i> | ATGCGAACCCTCTGCGACGTTTGC |
| <i>CpBBX19-R2</i> | TTACTTCTCACACACACCTTTG |
| <i>pGWB551-CpBBX19-F</i> | <u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> ¹ |
| | GAAGGGAAATGAAAGGTGG |
| <i>pGWB551-CpBBX19-R</i> | <u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> ² |
| | CTTCTTTAAGCCGCCAATAGTC |
| RT- <i>CpActin-F</i> | AGGCTAAGATTCAAGACAAGG |
| RT- <i>CpActin-R</i> | TTGGTCGCAGCTGATTGCTGTG |
| RT- <i>AtActin-F</i> | CTTCGTCCTCCACTTCAG |
| RT- <i>AtActin-R</i> | ATCATACCAGTCTCAACAC |
| RT- <i>CpBBX19-F</i> | AACCTAGTTCTTGTGAAGACTTGGC |
| RT- <i>CpBBX19-R</i> | CATCCATGTTAGCATCATTAGTTGG |
| SP1 | CTCAACTGGGTCCATTGGTTGCAAG |
| SP2 | TAAAAGAAAGCAGGTGCATTTTCACATATG |

¹ stands for *attB1*, ² stands for *attB2*

Table S2. Analysis of cis-acting elements of promoters. The amounts are the sum of Cis-elements in sense and antisense strand.

| No. | Name of elements | Amount | Sequence | function |
|-----|------------------|--------|-------------------------|---|
| 1 | TATA-box | 1 | TATA | core promoter element common cis-acting element |
| 2 | CAAT-box | 1 | CAAAT/CAAT/CCAAT | in promoter and enhancer regions |
| 3 | ABRE | 3 | GCAACGTGTC/ACGTG/CACGTG | cis-acting element involved in the abscisic acid responsiveness |
| 4 | ARE | 1 | AAACCA | cis-acting regulatory element essential for the anaerobic induction |
| 5 | CGTCA-motif | 2 | CGTCA | cis-acting regulatory element involved in the MeJA-responsiveness |
| 6 | MBS | 1 | CAACTG | MYB binding site involved in drought-inducibility |
| 7 | MYB | 2 | CAACCA | |
| 8 | MYC | 4 | CATTTG | |
| 9 | MYb | 1 | CAACTG | |
| 10 | MYc | 1 | TCTCTTA | |
| 11 | TC-rich | 1 | GTTTTCTTAC | cis-acting element involved in defense and stress |

| | | | | |
|----|-------------|---|------------|--|
| 12 | TCA-element | 1 | CCATCTTTTT | responsiveness cis-acting element involved in salicylic acid responsiveness |
| 13 | TGACG-motif | 2 | TGACG | cis-acting regulatory element involved in the MeJA-responsiveness |

Table S3. Names and accession numbers used for phylogenetic analysis

| Name | Species | Accession number |
|---------|-----------------------------|------------------|
| AtBBX1 | <i>Arabidopsis thaliana</i> | AT5G15840 |
| AtBBX2 | <i>Arabidopsis thaliana</i> | AT5G15850 |
| AtBBX3 | <i>Arabidopsis thaliana</i> | AT3G02380 |
| AtBBX4 | <i>Arabidopsis thaliana</i> | AT2G24790 |
| AtBBX5 | <i>Arabidopsis thaliana</i> | AT5G24930 |
| AtBBX6 | <i>Arabidopsis thaliana</i> | AT5G57660 |
| AtBBX7 | <i>Arabidopsis thaliana</i> | AT3G07650 |
| AtBBX8 | <i>Arabidopsis thaliana</i> | AT5G48250 |
| AtBBX9 | <i>Arabidopsis thaliana</i> | AT4G15250 |
| AtBBX10 | <i>Arabidopsis thaliana</i> | AT3G21880 |
| AtBBX11 | <i>Arabidopsis thaliana</i> | AT2G47890 |
| AtBBX12 | <i>Arabidopsis thaliana</i> | AT2G33500 |
| AtBBX13 | <i>Arabidopsis thaliana</i> | AT1G28050 |
| AtBBX14 | <i>Arabidopsis thaliana</i> | AT1G68520 |
| AtBBX15 | <i>Arabidopsis thaliana</i> | AT1G25440 |
| AtBBX16 | <i>Arabidopsis thaliana</i> | AT1G73870 |
| AtBBX17 | <i>Arabidopsis thaliana</i> | AT1G49130 |
| AtBBX18 | <i>Arabidopsis thaliana</i> | AT2G21320 |
| AtBBX19 | <i>Arabidopsis thaliana</i> | AT4G38960 |
| AtBBX20 | <i>Arabidopsis thaliana</i> | AT4G39070 |
| AtBBX21 | <i>Arabidopsis thaliana</i> | AT1G75540 |
| AtBBX22 | <i>Arabidopsis thaliana</i> | AT1G78600 |
| AtBBX23 | <i>Arabidopsis thaliana</i> | AT4G10240 |
| AtBBX24 | <i>Arabidopsis thaliana</i> | AT1G06040 |
| AtBBX25 | <i>Arabidopsis thaliana</i> | AT2G31380 |
| AtBBX26 | <i>Arabidopsis thaliana</i> | AT1G60250 |
| AtBBX27 | <i>Arabidopsis thaliana</i> | AT1G68190 |
| AtBBX28 | <i>Arabidopsis thaliana</i> | AT4G27310 |
| AtBBX29 | <i>Arabidopsis thaliana</i> | AT5G54470 |
| AtBBX30 | <i>Arabidopsis thaliana</i> | AT4G15248 |
| AtBBX31 | <i>Arabidopsis thaliana</i> | AT3G21890 |
| AtBBX32 | <i>Arabidopsis thaliana</i> | AT3G21150 |

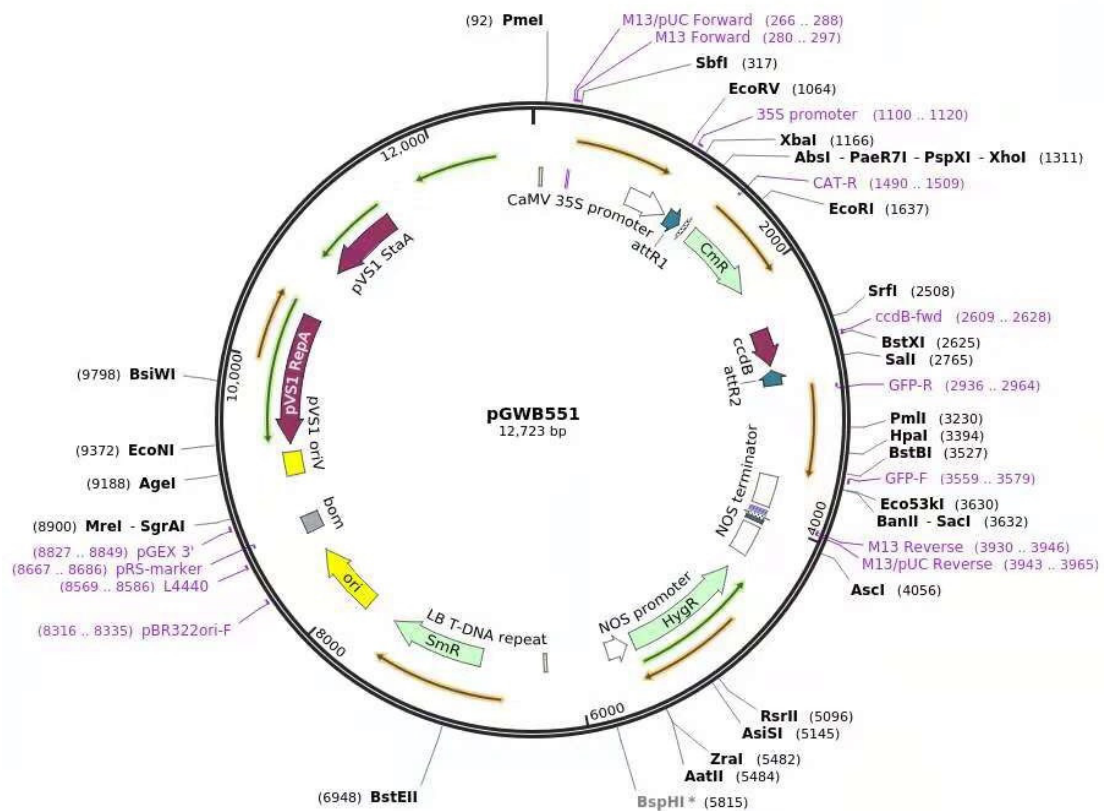


Figure S1. The simple map of the pGWB551 vector.

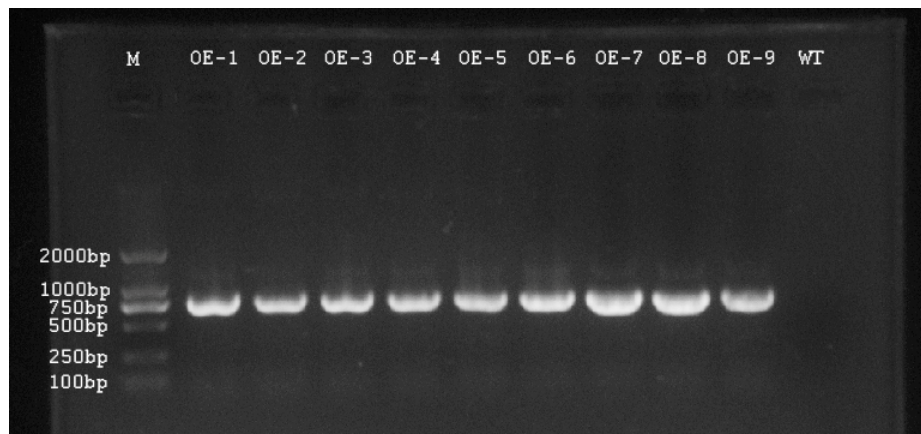


Figure S2. PCR Detection of *CpBBX19* overexpression *Arabidopsis*. OE-1-9: *CpBBX19* transgenic *Arabidopsis*; WT: wild-type *Arabidopsis*. M: DNA Maker DL2000.