

Supplementary Materials: Identification and Functional Divergence Analysis of WOX Gene Family in Paper Mulberry

Feng Tang, Naizhi Chen, Meiling Zhao, Yucheng Wang, Ruiping He, Xianjun Peng and Shihua Shen

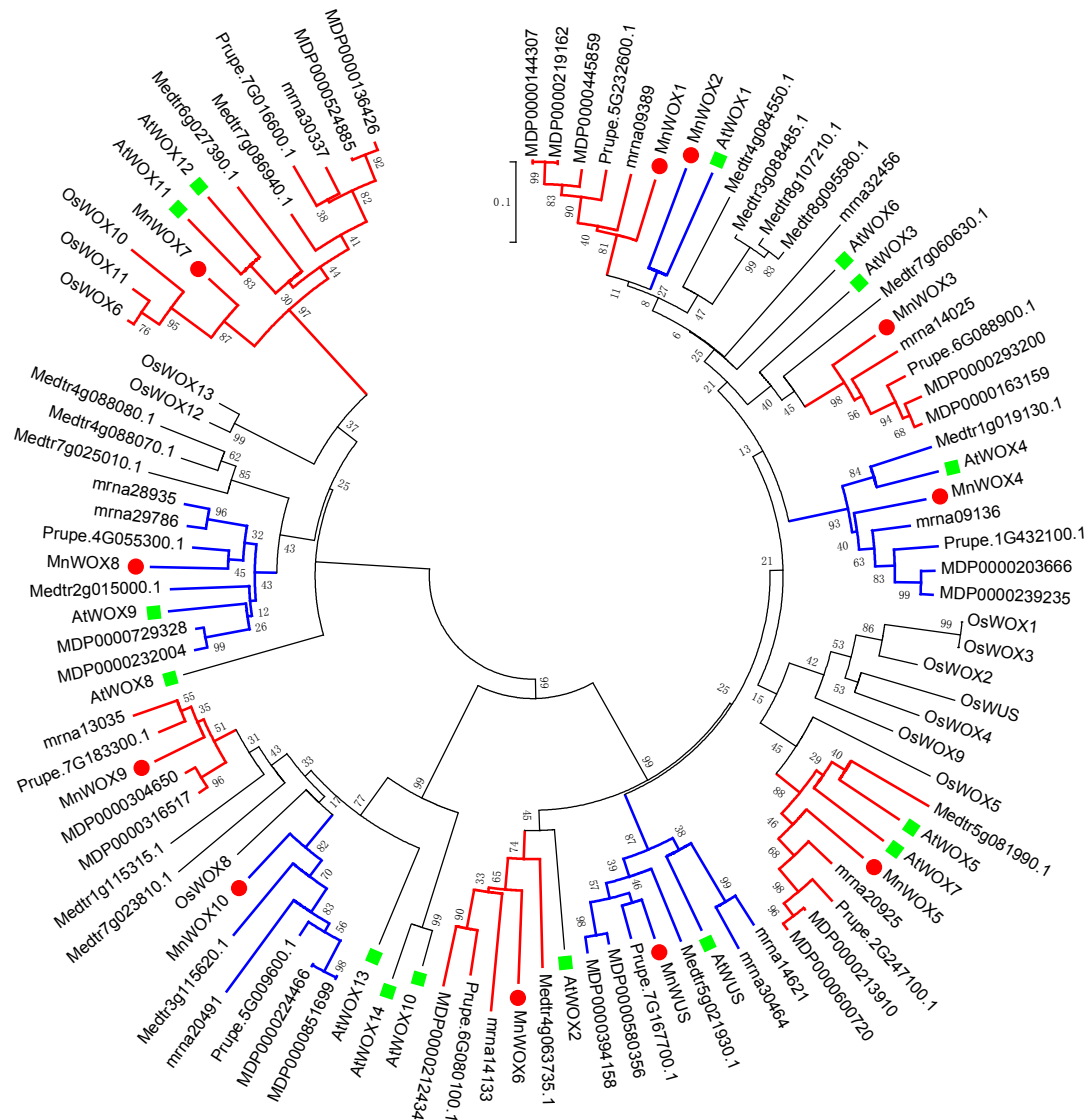


Figure S1. Phylogenetic analysis of BpWOX genes in *Morus notabilis*, *Fragaria vesca*, *Prunus persica*, *Malus domestica*, *Medicago truncatula*, *Arabidopsis thaliana* and *Oryza sativa*. There were eleven WOX genes in mulberry. Thus, we inferred that there might be eleven WOX genes in paper mulberry. Each of subclade WOX gene from seven species were aligned to find the conserved region and design the degenerate primers. The selected sequences for alignment were labeled with red or blue line. The MnWOX proteins were labeled by red dots, and the AtWOX proteins were labeled by green square. *M. notabilis*, Mn; *F. vesca*, mrna; *P. persica*, Prupe; *M. domestica*, MDP; *M. truncatula*, Medtr; *A. thaliana*, At; *O. sativa*, Os.

Figure S2. The sequences of *WUS* subclade were aligned to find the conserved regions and design the degenerate primers. The red line was the location of degenerate primers for *BpWUS*.

Figure S3. The sequences of *WOX1* subclade were aligned to find the conserved regions and design the degenerate primers. The red line was the location of degenerate primers for *BpWOX1*.

[illegible]

Figure S4. The sequences of *WOX2* subclade were aligned to find the conserved regions and design the degenerate primers. The red line was the location of degenerate primers for *BpWOX2*.

[illegible]

Figure S9. The sequences of *WOX8* subclade were aligned to find the conserved regions and design the degenerate primers. The red line was the location of degenerate primers for *BpWOX8*.

Medtr3g115620.1			0
MDP0000224466	ATGGGGGAGTGGCGGCAGCAGCAGCAAAATAATCAACAGAAATTCAGAAACCCAGAAGAAGATGATGGGGTTGGAAATGGAAATGGAGGGTTGTTCTGTAAGGTTATGACCGACGAG	120	
MDP0000851699		0	
mrna20491		0	
Prupe. 5G009600.1		0	
MniWOX10		0	
Medtr3g115620.1		0	
MDP0000224466	CAAAATGGAGCAGCTGAGGCAACAGATCGCCGCTCTACACCAACCATTTTGACCAAGCTTTGAGCTTGCACAAAGCAACTCTCTGCCAACAGGATCTTACTTGTCGGTCTGCCCTCTCTCT	240	
MDP0000851699		0	
mrna20491		0	
Prupe. 5G009600.1		0	
MniWOX10		0	
Medtr3g115620.1		0	
MDP0000224466	CTCTTCCAATCTGAATGGGCTTTTGTAATGAATTTATTCACCTTGCAAAATATTGATTTCATGGGTGGGAGCATGAANACCCGTTCTCAAGTTGGAGAAGCTTACCTCAGCTTGGTGG	360	
MDP0000851699		0	
mrna20491		0	
Prupe. 5G009600.1		18	
MniWOX10	ATGGTTATGGAGTGAAGAGTGAACAACATCAACATCAACAACAACAGATCTCATCATCAT	63	
Medtr3g115620.1		75	
MDP0000224466	ATATATCTCCTCTCAACTCTTTCCAGGAGTTCAGGTGCAGGACCTTGNCAAATTCAGCCCCGGT...GCCCAACCTTAACCTCGRAATAAAGCAGATAGCGANACACAAA	477	
MDP0000851699		0	
mrna20491	ATGGAGGGGCGAGAGAAGAGTGTGCAGTGAATTTGACGGCTGAGGACTCAGAACAGATGGGTTTGCTCATGACGGACCAAGTC	93	
Prupe. 5G009600.1	CAGCAGAGCAGAGATTCAGATCAGAATCAACAGCAATAATCCACCATTTGCAGAACCCAGAGATGCGTTTGGAAATGGAATGAGGGTTGTCCTGAAGTGTATCCGACAGCAAAATG	138	
MniWOX10	CATCTATAGAATATCAGAGACTGCAGAAAGAGGAGAGAGAAATGATTGTTAATGGGTTTCTCCAAAGAGAGATTAGGGTTAATGTCCTGAGAGTGTATGCTGACGACAAATG	183	
Medtr3g115620.1	GAACTTTTAAACAACAATATCTGCTACACCAACATTTCTGACCAACTTGTGAGATGCATTAATCAATTAACATCAACATGACCTCGCTGGACGAGCTTCGAAATTCGACTGT	195	
MDP0000224466	ATGTGGATTAGTAATTTAAGTTTCTTTGTGGAAGAAGATTTTATTATTGGTGTGGCCTTCATTTCCTATTAATCATCAGTCTTCTGTTTCAGGATGAGACTCGCGAATTTTGTGTGT	597	
MDP0000851699		24	
mrna20491	GAGCAATCTCCAGAAACAGATCGCGCTCTACGCGCTCATCTGTGAGCAGCTTGTGATATGCACAAGGCTTCACTGCCCAACAGGATCTTCTGGATGAGAGTGGAAATCTGAGGC	213	
Prupe. 5G009600.1	GAGCTGCTGAGGGCAGAGATCGCTGCTCTACGCCACCATTTTGTGAGCAGCTTGTGAGATGCACAAGGCGCTCACTGCTCAACAGGATCTTACTGGATGAGACTGAGGAAATTTGTGT	258	
MniWOX10	GAGCTTCTGAGGGCAGATCTCTGCTACGCCACCATCTGCGACAGCTTGTGAGATGTTCAACTCTGCTCACTGCCCAACAGGATCTTCTGAGCTGAGACTGAGGAAATCTGAGTAT	303	
Medtr3g115620.1	GATCGACTGAGATGCTTCTCTCTGAGACAGATTAAGCTCAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	315	
MDP0000224466	GATCGATTTAGCCTATCTG...TGSCACAGATTAAGCTCAGGCGAGCTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	714	
MDP0000851699	GATCGATTTAGCCTATCTG...TGSCACAGATTAAGCTCAGGCGAGCTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	141	
mrna20491	GAT...TTGATGCTATTTG...TGSCACAGATTAAGCTCAGGCGAGCTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	327	
Prupe. 5G009600.1	GATCGATTTAGCCTATCTG...TGSCACAGATTAAGCTCAGGCGAGCTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	375	
MniWOX10	GATCGATTTAGCCTATCTG...TGSCACAGATTAAGCTCAGGCGAGCTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	420	
Medtr3g115620.1	AAGCAGAGATCAAAAGATTAACCATTAAGCTAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	435	
MDP0000224466	AAGCAGAGATCAAAAGATTAACCATTAAGCTAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	831	
MDP0000851699	AAGCAGAGATCAAAAGATTAACCATTAAGCTAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	258	
mrna20491	AAGCAGAGATCAAAAGATTAACCATTAAGCTAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	444	
Prupe. 5G009600.1	AAGCAGAGATCAAAAGATTAACCATTAAGCTAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	492	
MniWOX10	AAGCAGAGATCAAAAGATTAACCATTAAGCTAGGCGACTGCGACCTTATCTGCTTAATCTGCTTAATCTGAGCTGCTTCTGCTGAGAGCTATGCTGCTGAGCT	537	
Medtr3g115620.1	CTCTGGGATTCATGCGATCTCAGGCGAGGCAAGAGT...GAGCTCTGAGAGAGGAGCAAGCTGTGAGAGAGGCTTCTGAGCTTCTGAGCTTCTGAGCT	552	
MDP0000224466	CTCTGGGATTCATGCGATCTCAGGCGAGGCAAGAGT...GAGCTCTGAGAGAGGAGCAAGCTGTGAGAGAGGCTTCTGAGCTTCTGAGCTTCTGAGCT	939	
MDP0000851699	CTCTGGGATTCATGCGATCTCAGGCGAGGCAAGAGT...GAGCTCTGAGAGAGGAGCAAGCTGTGAGAGAGGCTTCTGAGCTTCTGAGCTTCTGAGCT	366	
mrna20491	CTCTGGGATTCATGCGATCTCAGGCGAGGCAAGAGT...GAGCTCTGAGAGAGGAGCAAGCTGTGAGAGAGGCTTCTGAGCTTCTGAGCTTCTGAGCT	549	
Prupe. 5G009600.1	CTCTGGGATTCATGCGATCTCAGGCGAGGCAAGAGT...GAGCTCTGAGAGAGGAGCAAGCTGTGAGAGAGGCTTCTGAGCTTCTGAGCTTCTGAGCT	600	
MniWOX10	CTCTGGGATTCATGCGATCTCAGGCGAGGCAAGAGT...GAGCTCTGAGAGAGGAGCAAGCTGTGAGAGAGGCTTCTGAGCTTCTGAGCTTCTGAGCT	648	
Medtr3g115620.1	GTCAAGGATTCGACCTTCAGAGTCTTGACATAGGATTTGAGCACTTGTGGGAAATTTGAAGTTGCAGGCTGCTATATCTTACTTCTTTGA	648	
MDP0000224466	GCTCAAGATTCGACCTTCAGAGTCTTGACATAGGATTTGAGCACTTGTGGGAAATTTGAAGTTGCAGGCTGCTATATCTTACTTCTTTGA	1058	
MDP0000851699	GCTCAAGATTCGACCTTCAGAGTCTTGACATAGGATTTGAGCACTTGTGGGAAATTTGAAGTTGCAGGCTGCTATATCTTACTTCTTTGA	423	
mrna20491	ATTCAAAATTCGCTTGGCTTG	570	
Prupe. 5G009600.1	GCGCAATATTCGACCTTCAGAGTCTTGACATAGGATTTGAGCACTTGTGGGAAATTTGAAGTTGCAGGCTGCTATATCTTACTTCTTTGA	657	
MniWOX10	GAGCTTAATTCGAGTTTCAGAGTCTTGAG...ATAATCAATG...ATTGTTGGCACTTTTCACCCTTCAGGAAGCTCTTGGCTCCGCTCACCCACCCTGGTGTGCTAGTTGCCTTTAGG	765	
Medtr3g115620.1			
MDP0000224466	AAAAAGTGGTGTGAATGATTTACAAAAACAGTTTCATGACCGAGTCTTGA	1107	
MDP0000851699			
mrna20491			
Prupe. 5G009600.1			
MniWOX10	GTGTGAGTGGTCCCGAGTTGATGAATTTGTCGGCTAGTGGAAAGTGCATTGAGCTACAAGCTTCGCTTCCTGGGAGGATGGATGTGGATGAGCTGTA	866	

Figure S11. The sequences of *WOX10* subclade were aligned to find the conserved regions and design the degenerate primers. The red line was the location of degenerate primers for *BpWOX10*.

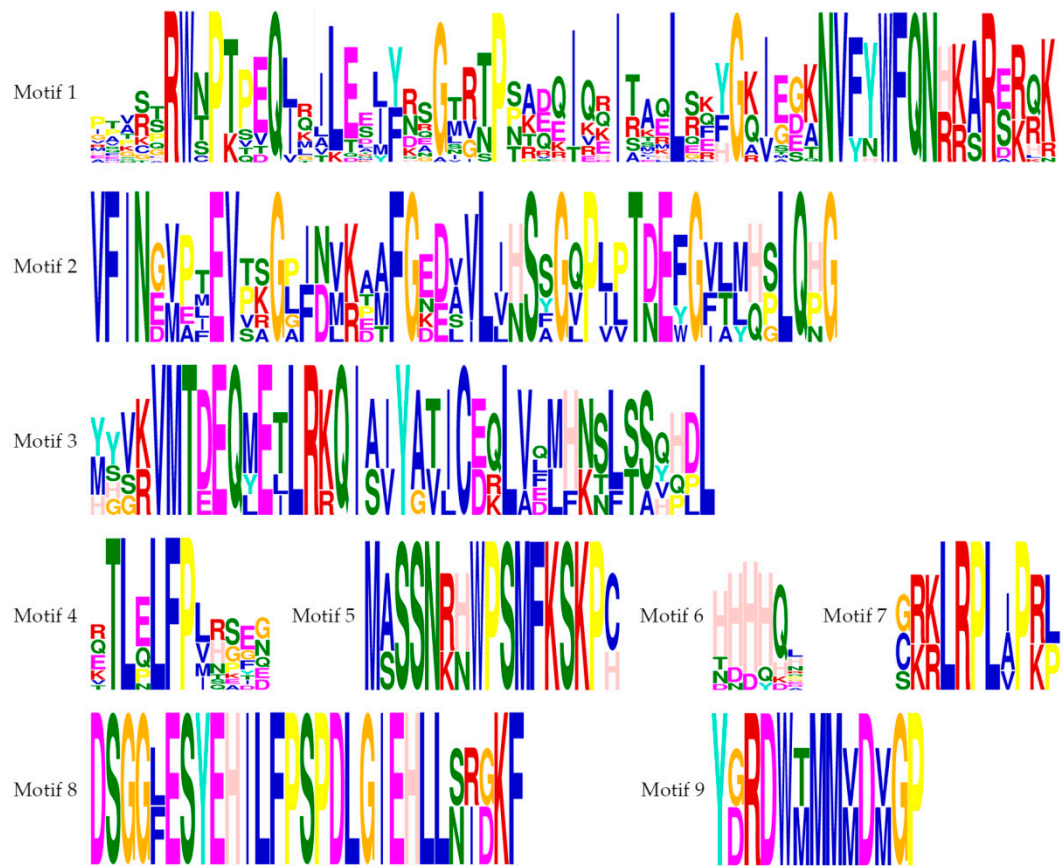


Figure S12. The conserved amino acid sequences of WOX protein in *A. thaliana* and *B. papyrifera*.

Table S1. The degenerate primer sequences of *BpWOXs* used to get fragments from paper mulberry.

Gene Name	Forward primer (5'-3')	Reverse primer (5'-3')
<i>BpWUS</i>	TGCMGGCAAAGCAGYACAAGRTGG ACDCC	TGAACCTTTTCTTYGCCTCTCNCGAGC YTT
<i>BpWOX1</i>	TCAAGATGGAACCCNACDGAGCAG YTGAGG	GYTTCTGCCTYTCTCTGGCCTTRTGRTTT TG
<i>BpWOX2</i>	AATATTGCACAAGTAGTGCCATTAC AAAGCAC	CGCTGCTTCTTTTGTCCCTTGCTTTGTG GTT
<i>BpWOX3</i>	GATGGTGCCCAACVCCCGAGCARCT GATGMTCT	TTCTGYCTRICYCTDGCCTTGTGRTTYTG
<i>BpWOX4</i>	GTGAGACGCACCCRGGAGGGACR MGNTGGAA	GGCTGAGGCCRAGGCTGYTGCGYTTYTG CTT
<i>BpWOX5</i>	GGAACVAAGTGYGGDCGTTGGAATC CNAC	GARACYTTGCGGCGCTTCTGYCTYTCYC TG
<i>BpWOX7</i>	GAACCGNGCCGTNAGGTCAAGDTG GAYWCCAAAG	GCCTGCADCTGCCTBTGDCGGCGDCGR GABC
<i>BpWOX8</i>	ATGGCTTCWTCAAACAGACACTGG CCDAGCAT	GCTTGTGYTTGCTTCTBGATTNCKGTTY TG
<i>BpWOX9</i>	TGGGAAATCTRTACTGCGAYACGYT GATGACATC	GGATCMAAGAAATGCAGRTCAGARCTY ATCTC
<i>BpWOX10</i>	CTGGAMTGARRCTDGGDARTBTTCA CTATGAT	GATTGCTTTCTYTTYGAACGAGCTCTYCT GTT

Table S2. The sequences of PCR products amplified with the degenerate primers in paper mulberry. The fragment sequences are used to design the RACE-PCR primers. The red backgrounds are the 3'-RACE primers, and the green backgrounds are the 5'-RACE primers.

Gene Name	Sequence (5'-3')
<i>BpWUS</i>	TGCAGGCAAAGCAGCACAAAGATGGACACCAACAAGTGACCAGATAAGGATTCTCAA GGACCTTTACTACAACAATGGCGTGAGGTCCCGAAGTGCAGAGCAGATTCAGAGGAT CTCTGCCCCGCTGAGGCAGTACGGCAAGATTGAAGGCAAGAATGTCTTTATTGGTTT CAGAACCATAAAGCTCGCGAGAGGCAAAACAAAAGGTTCA
<i>BpWOX1</i>	GGTGAACCCGACGCCGAGCAGCTGAGGGCCCTCGAAGAGCTCTACCGCCGCGGG ACCCGAACGCCATCGGCCGAGCAAATTCAGCACATCACGGCTCAGCTTCGGAGATAT GGAAAGATTGAAGGGAAGAACGTGTTCTATTGGTTCCAAAATCACAAGGCCAGAGA GAGGCAGAAGC
<i>BpWOX2</i>	AATATTGCACAAGTGGTGCCATTTCAAAGCACGCGGTGGAGTCCGACGCCGGAGCAG TTGATGGCGCTGCAAGAGCTGTACCAGCGTGGACGCGGACCCCAACGGCGCAGCA GATCCAGGAAATCACCGCGAAGCTCCGGCGGTTTCGGGAAGATTGAAGGCAAAAACG TGTTCTACTGGTTTCAGAACCACAAAGCCAGGGACAAAAAGAAGCAGCG
<i>BpWOX3</i>	TCAAGATGGTGCCCAACACCCGAGCAGCTGATGATCCTGGAAGAGATGTACAGGGCT GGAACAAGAATCCCAACGCGTCTCAAATCCAACAGATCACAGCCCATCTCTCGTTT TACGGAAAGATTGAAGGC AAGAATGTGTTCTACTGGTTTCAGAACCACAAGGCCAGG GACAGGCAGAAGC
<i>BpWOX4</i>	GTTGAGACGCACCCGGGAGGGACGCGGTGGAATCCGACGCAAGAACAGATAGGGAT ACTGGAGACGCTGTACAGAGGAGGAATGCGCACTCCGAACGCGCAACAAATAGAGC AAATCACGGCTCAGCTGGGCAAGTTCGGCAAGATCGAAGGCAAGAACGTGTTCTACT GGTTCAGAACCAAGGCCCGGAGCGGCAGAAAGCAGAAAGCGCAACAGCCTCGG CCTCAGCC
<i>BpWOX5</i>	GGAACAAAGTGCGGCCGTTGGAATCCCACGAGCGAGCAGGTTAAAGTCTGACCGA TCTCTTCCGGTCCGGTCTCCGAACCCGAGCACCGACCAGATCCAGAAGATCTCCGC TCAGCTCAGCTTTTACGGCAAGATCGAGAGCAAGAACGTCTTCTACTGGTTTCAGAA CCACAAGGCCAGAGAGAGACAGAAGCGCCGCAAGGTCTC
<i>BpWOX7</i>	GAACCCGCCGTGAGGTCAAGGTGGAATCCAAAGCCAGAACAAATTCATCCTTGA GTCCATCTTCAATAGTGGAATGGTAAACCCCTCCCAAGGACGAGACCGTGAGGATACG GAAGCTACTCGAGAAATTCGGGGCCGTTGGCGACGCCAACGTATTCTACTGGTTCCA AAACCGCCGCTCGAGGTCCCGCCGTCGCCAGAGGCAGATGCAGGC
<i>BpWOX8</i>	ATGGCTTCATCAAACAGACACTGGCCAGCATGTTCAAGTCCAAGCCTTGCAACACC CACCCCCACCAATGGCAACACGACATCAACCAGTCTCTCATCTCAACTGGCTGCCAC AGAATCCCTACACTTCAGTTGGCGGGTGCGAGGAGCGGAGCCCGGAGCCGAAGCC GAGATGGAATCCGAAGCCGGAGCA GATTCGGATCCTGGAAGCGATCTCAACTCGGG GATGGTAAACCCGCCGAGAGACGAGATAAGAAAGATCAGGGCACAATTGCAGGAGT ACGGCCAAGTGGGTGACGCCAACGTGTTTACTGGTTCCAGAACAGGAAATCCAGA AGCAAACACAAGC
<i>BpWOX9</i>	TGGGAAATCTATACTGCGACACGTTGATGACATCTGCGGGCCACAAGATCACGGCGA GACAGCGTTGGACTCCGACTCTGTGCGAGCTACAGATTCTGGAGAGCATGTTTGACC AGGGGAATGGAATCCAAGCAAGCAAAAAGATCAAAGAGATAACCTCTGAAGTGAAGC

	CAGCATGGCCAGATTTGGGAGACTAATGTCTACAATTGGTTCCAGAACAGGCGTGCT CGGTCGAAAAGGAAACAGCAGAATAATGCAGCACCCAACAATACTGAATCTGAAGT GGAGACTGAAGTTGAGTCACCTAAGGACAAGAAGACAAAGCCGGAGGAGTTTCTGT CCCAACAGAACTCGGCTTCGAGGGCTGAAGATATCTGCTTCCAGAGCTCCGAGATCA GCTCTGACCTGCATTTCTTGATCC
<i>BpWOX10</i>	CTGGAATGAAGCTGGGAAGTCTTCACTATGATCCACTAATGACATATGGTGGCCACAA GATCACTGCAAGGCAAGGTGGTCACCAACGCCTGTGCAGCTTCAAACCTAGAGA AAATTTTCGATGAAGGGAATGGGACTCCGAGCAAGCAGAAGATCAAAGAGATAACC GTAGAGCTGAGTCAACATGGCCAAATTCGGAAACCAATGTCTACAATTGGTTCCAG AACAGGAGAGCTCGTTTCAAGAGAAAGCAATC

Table S3. The primer sequences of *BpWOXs* used for full-length cDNA amplifications.

Gene Name	Forward primer (5'-3')	Reverse primer (5'-3')
<i>BpWUS</i>	ACATGGGGACCATCCACATAACAT TTT	GTGTCAAGACCATCGTCATATATAATAATC A
<i>BpWOX1</i>	GATAATAGTACTACACTACAATGCCT ACCTCC	CCATTTTAAGGCAAACCTCCTTTTCA
<i>BpWOX2</i>	ATGTCCATTTGAAGCCTAGATATCAA ACTA	AAATCTTCTATATTTTACTCACGTCCTTTAT TC
<i>BpWOX3</i>	ACCCCAAGAGACAAAGACAAAAGT CGAATC	AGTCTTTTGAAGGTAAACAAGGCATTTT
<i>BpWOX4</i>	CCTCACCTCATCATTTGGTCTTGGTT ATC	TGATCTGCTGCTTATTGGAACCAAAGAGC G
<i>BpWOX5</i>	ATACATGCACACGGGTTTCACATTG	TCATAAACATAAACTACCGATGTCCTACC GTAG
<i>BpWOX7</i>	CAATGGAAGATCATCACCATCATCA AGGC	GTAAAGACGTAATATTTATGGGAGACTG GC
<i>BpWOX8</i>	ATCTCATCAGTACCAAAAACAAGAG CAAC	CCACGAAAAGCATTCATTTTTACTACAC
<i>BpWOX9</i>	GTTTGTCTCTTTCTTTCCAACCTTTTT C	GGTTCATCCATTCATATCCAATCATAAAAT G
<i>BpWOX10</i>	AGTGTCACTACTAAACACTTGCTTG CAATCTAAG	GGTTTTTCAATAGACCCAGTTCATCATAAT CTG

	CTACTAATCTCAAAGAAGAGTGTACGACGTCAAAAACATGTCCCTCATTCCACCCCAACGATGA CCATCTAATGACCAGTCTAAATTCCTATCTGCATGGCTCTCTTTCTCTCTCTCTCTCTCTCTCT CTCTCTCTCTCTAAGTCTATGTCTAGATGCTACAAGGGTAGGCTTGAAAGTGTCTGTGTGTTGTT TATTTTCTTGGTTTCTGTTATTTGTCTAGAGCTCTAGGAAGAGAAAGATCGCAATTGTCTAGCGA GTAATACGTTACTGGCTCTATTGTCTTACTACTACCTACTCAATGAAATGCCTTGTTTACCTTC AAAAGACT
<i>BpWOX4</i>	CCTCACCTCATCATTTGGTCTTGGTTATCCCAAGCACACCTCATTTCCCACTTTGGTGACTTTA ATCGTACCAGTACATTACTTTGCTATATAACCAGGGGTTTGTGTAGTGATTAGTCTCTCCACCAT GGGAAGAAGTAGCTGCAGCATGAAGGTGCATCAGTTCGCACGTGGACTCTGGGAGGCTCACG AACCTCCCTCACGCTTGGCTGCAAACGCTTACGCCCTCTTGCTCCCAAGCTCCCTAACTCTCC TCCTTCTACTACCGACACTACTCACATTACTACCCTGCTTCTTCCCCCACTCCATCGCTCCTT TCGATCTCAAGAGCTTCATCAGACCCGAAAGCGGGCCCCGAAAGCTCGGTTCTCCGACGAC AAGAAAGACTCGCTCCGTTGAGACGCACCCGGGAGGGACGCGGTGGAATCCGACGCAAG AACAGATAGGGATACTGGAGACGCTGTACAGAGGAGGAATGCGCACTCCGAACGCGCAACA AATAGAGCAAATCACGGCTCAGCTGGGCAAGTTCGGCAAGATCGAAGGCAAGAACGTGTTCT ACTGGTTCCAGAACCACAAGGCCCGGAGCGGCAGAAAGCAGAAGCGCAACAGCCTCGGCCT CAGCCACAGCCCGGAACCCCGACCCCAATCGCCATAGCCACCTGCGGTTTGGACACTAGGG GAGACTTGGACAGAGACCCGGAAGATAGTACTAGTCCGTACAAGAGGAAGTGCCGGAGCTG GGGATTTGAATGCTTGGTAGAAGAGCATAGTAGTAGAATATTGTATAACAATTATAGAGATCAG GATCAGGGAGATAGGACTCTGGAGCTTTCCATTACACCCGGAAGGCATGAGATGATGGGG GGTTTTGGGGAGAGAGAAAATACTAGTAATAATTCATCCTTCTTTCACCATCATATCATTTTTCA GCTTTAGTTTATGTGGCCGGTATATGAATATTATCAGCTTAATATGTGCTTTTCTTTCCTTTTCTT CTTTTTTCTTTCGCTTCGGGAAAAGAAGAAAAAAGAAGGTACTTGTCTGATCTCTTTTGTAGT ACTGATCTGCTGCTTATTGGAACCAAAGAGCG
<i>BpWOX5</i>	GCACACGGGTTTCACATTGAAAACCTGATCAGAAAACCTGCTATATCACTAGGAACTTCAGTGAC AGAGATCAGAAGATTTTGGAGCTCAAAAAATGGAAGATCATGGCATGTCAGGCTTTTGCAATTA AAGCGGGACATAACATGGCTAACAAACGTTGTCGTTTCGCGGTACTACTGGAACCAACGGCAGC ACTGGAACAAAGTGCGGCCGTTGGAATCCCACGAGCGAGCAGGTTAAAGTCCTGACCGATCT CTTCCGGTCCGGTCTCCGAACCCGAGCACCGACCAGATCCAGAAGATCTCCGCTCAGCTCA GCTTTTACGGCAAGATCGAGAGCAAGAACGTCTTCTACTGGTTTCAGAACCACAAGGCCAGA GAGAGACAGAAGCGCCGCAAGGTCTCCGTTGATGATCATGACAAGGATTTCTGTTCTAAGAGA GGACAAGATTCCCTCTCCCAAACGTAATTTTGGGAGATATATCACAGTCGCCATCATCATCAT CATCATCATCATCATGATCACGAGGAATATCAGGTTTCTGAGCCGGAAGAGTGATTGAGACG CTCCAACCTCTTCCGTTGAATTCGTTTCGGAGAGCCGGATTTCGGAGAAGCTGAGATTCATGAT CAGTGCAGAGAAACAGCGGCTTTTCCGTACACAATTGAGTCAGCAATAGAACATCCTCCAGT ACTGGATCTGCGCTTAAGCTTTGTCTAATTAAGCATGTTTAGTTTACATCTTAGCCAAAAAAA AAGAAAGAAAGAAAAAGAGAGAGATAGTTTTTGTATCACGTATTAATTAGG
<i>BpWOX7</i>	CAATGGAAGATCATCACCATCATCAAGGCCAAGACGCTAACAGTCCAAGCAACGGCTCCGAG AAAAGTAGTAGTGCTAGTACCGAACCCGCCGTGAGGTCAAGGTGGACTCCAAGCCAGAAC AAATTCTCATCCTTGAGTCCATCTTCAATAGTGAATGGTAAACCCTCCCAAGGACGAGACCG TGAGGATACGGAAGCTACTCGAGAAATTCCGGGGCCGTGGCGACGCCAACGTATTCTACTGGT TCCAAAACCGCCGCTCGAGGTCCCGCCGTCGCCAGAGGCAGATGCAGGCCAGCCTCGAAGT CCAGCAGCGGAACCAGGCCGACAAGGTGTTGGTGGTGCATTCAGTATGAAAGCAATAATT ATAACAGCAATAGTACTACTCAGGCCATGGCCGGTTATGTAGCTGGTTCGGCTCTTTTGTGTTG

	CTCTCCTTCTTATAGTAATCTTGTGGGTTCTTCTTCTCTCTTCATCATCTTCATGCGGAGTTTGT GCAGATCATCATGATCACCATCATGGTTTGGACTACAATACCAATTCTTTCAAGTCTCTGGTC AAATGGGTTTTCCGCAAATTGAGCAAAGCTCTCCTGTAAACGTCTGTTTTGTATCCTCCAGATAG CTCAAACATGCACCTCGAATCTGGATTCATCACAGTGTTTCATCAATGGGGTTCCAACAGAAGT TCCTAAGGGCCTAATTGACTTGAAACCAATGTTTGAAAAAGATGTGATTTTGGTGCATTCTCC GGAGTTCCACTTCCAATAATGAATTTGGCATTCTAATGCAGGGCTTGACGCTGGTGAAAGC TATTTCTGGTGCCGAGAACACGTTAAGTAATAACTGAAGCAGCATCTAGCTAGATTGCTAC CAATGGAAGCTTCCCTACATATTCCAATCTTGATTTCTTCATTCTTTTTTCTCTCCTTATACATT TATTATAGATTTCTATTTTGTGGTTTTAAGGATTTAAATTGAGAGTGCTATTATATATATGTGTGTG ATTATTAGAGACAATTAAGATCTATACATAAGAGTTGCCAATATCATGCATATGATTGCTTATTT GTAATGCCAGTCTCCCATAAATATTACGTCTTTAAC
<i>BpWOX8</i>	ATCTCATCAGTACCAAAAAACAAGAGCAACAAAACGGCTACTTCATCTCATTGTTTCTCTTTTGT TTATCTACTTTTAAATTACTTCTCAGATCATCAGTTTCATCATCATCATCATCATGCGTTTCATCAAA CAGACACTGGCCAGCATGTTCAAGTCCAAGCCTTGCAACACCCACCCCAACCAATGGCAAC ACGACATCAACCAGTCTCTCATCTCAACTGGCTGCCACAGAACTCCCTACACTTCAGTTGGCG GGTGCGAGGAGCGGAGCCCGGAGCCGAAGCCGAGATGGAATCCGAAGCCGGAGCAGATTCC GATCCTGGAAGCGATCTTCAACTCGGGGATGGTAAACCCGCCGAGAGACGAGATAAGAAAG ATCAGGGCACAATTGCAGGAGTACGGCCAAGTGGGTGACGCCAACGTGTTTTACTGGTTCCA GAACAGGAAATCCAGAAGCAAAACAAAGCTCCGACACCTCCAAAACCTCAAACAACAACA ACAACAGCAAACTCAGAACAATCACCAAACTCTCCGAACCTCAAATTTACCACCACTACAA CCACTAGTAGTACTACTCTCTACCACCAGCACAAAGAGCTCCTTCTCATCGTCTCTGCTCTC CTCGGAGAAATCGTCTCCGAAAGCCCCGAACAGATCGCCGATCTTCTCGATGGGCTTCACCG AAGCCCTAAACTCTCCGACCGGCTCGGTCAACCAGAGCACCTTCTTTCAGACCAACGGTAGT ACCATCACTAGTACTGGTCACGACCCGTTCTTGCCGAACCTTTTTCTTCCGGTGCAACAGT CCAGTACTAGTACTACTCAAGGGTTTTGCTTCCCTCATGATCAGCTCACGCCGACTGCGGTCCA TCAGATTATGCCCCGAGATCATTATCAACAGATTAACAACATCGGACCGTGCACTAGCCTCTTG CTCAGCGAGATCATGAACCATCATGGGGCTTCAACGAAAGGCCATGATCATGATCATCATAAG CTGCTACCGCTTAATTACACTCCGTGAGGCGCGCGGACGACGGTGCCAAGGACTACTTCT CCGGCCAGTACTCTCTCACTACGGCTGCTACTAGTACTATTGAATCTTCAACATCACAAAGGA ATCAAATCCAAGCAGGTGTTGGAGAACAGGGAACCTTCTGGAGTGGGGCCAATCGGAGGGGG CGGTAAACGACGGTGTTATAAACGACGTGGCGTTTGAGGTGCCGGCGGGGCCGTTCAATG TGAGGGAGGCGTTTGGGGAGGAGGCTGTGCTCATCCACTCCTACGGTCAGCCAGTTCTCACC AACGAGTGGGGGGTCACTCTTCACTCCCTCCAGCCTGGTGCTTTTATTATCTAATTTAGGAGC TTGGGAGATATATCTAGCTAGATATGAGAAAAACACACGCTGGCTGTGGTTCAATTAATCACT ACATATATTTCTGCTCTTATCTCTTAAAGTTTGGATTAGGGCTTTCAATGACTTGTATATTAGCTC TGTTAATTAGTCCGTTTAGTGACAACTTGAAAATTTGGTGACTCATATTATATATATGATTCTC TTTCAGTTGTTAAGTAGAGAATTATATCAAAGATGTGTAGTAAAAATGAATGCTTTTCGTGG
<i>BpWOX9</i>	GTTTGTCTCTTTCTTCCAACCTTTTTCTGTCTCTCTCCTCTCTCCGACTAAAGACTGGTTTT TTAGACAAACATACTCCTCCGTTTCCAGACTGTGAGCCTGAGAAAGAGGAGAGAGAAAGA AGGAGGATGATGGGGTGGGAGAAGCAGG AGGAGAGAGAAGGGCATAATAATAGTAACAGCAGTAATCTGTATCATCACCATGTGAAGGTG ATGACTGACGAGCAATTGGAGACGCTTCGCAAGCAAATCTCCATCTACGCCACCATTGCGAG CAGCTCGTTCAGATGCACAAGAACCTCACTTCCCATCACGATCTCCCTGGCGTCCGATTGGGA AATCTATACTGCGACACGTTGATGACATCTGCGGGCCACAAGATCACGGCGAGACAGCGTTG

	<p> GACTCCGACTCCTGTGCAGCTACAGATTCTGGAGAGCATGTTTGACCAGGGGAATGGAATC CAAGCAAGCAAAAGATCAAAGAGATAACCTCTGAACTGAGCCAGCATGGCCAGATTTCTGGA GACTAATGTCTACAATTGGTTCCAGAACAGGCGTGCTCGGTGCAAAAGGAAACAGCAGAATA ATGCAGCACCCAACAATACTGAATCTGAAGTGAGACTGAAGTTGAGTCACCTAAGGACAAG AAGACAAAGCCGGAGGAGTTTCTGTCCCAACAGAACTCGGCTTCGAGGGCTGAAGATATCTG CTTCCAGAGCTCCGAGATCAGCTCTGACCTGCATTTCTTTGATCCGCATTCCAACAAAGGAGA AGCCATGTTCTCATCAAACAATTCTTTACGGCCTGCGAGGAATTTGACTCAAATGCCATTCTAC GACGGTCTGCTCTCGAACTCAAGAAATGAGCATCTGAGTGGAATAATGGAAGTGCCAGGGA GTTATGGTCTTTACGAGCATGCGGAAAACCTACAGCATGACGGGATGAGGCTATCGTCCGAGGG AGAAGTTAATATTGTGAATAGAAAGAGTACAGATTCCAGATGCAGCTTGGTGATGTGTTATTA GCAATCATTGCAATTGGGTGCCGTACAAACAGTCTTGGATATGGAATGTACCTGAACCCACC TACTTAGTAAAGAGTACAAAGCTAACTTAGCAGAGTTTAATGTTTCTCTTTTTGTCTTCATTTT ATCTGGTCCAGTTTATCAGGGAGTAGAATGTCATGGTTTTGATTGGTGAACAAACAACTTGA GTCTCTTTAAGTTCCTGTTAACTTTTGATAAATGTGAGGACTTTTTGCCCTTTTTGTGTTTTGGC TGATAGTTGATCCGTGAGTTCCTTTTGCTTTTGTCATACAATGAACCCATTTATGATTGGATAT GAATGGATGAACC </p>
<i>BpWOX10</i>	<p> AGTGTCAGTACTAAACACTTGCTTGCAATCTAAGGCTGACTCTCTCTCTTTGTTTCTCTTTCTCT GTCTAAGTCAGATTCTCTGACAGAGGTTGAAGATCCAGGGCTTTTGGACAAATGGTTAACTA ACTCTGGACTGCTCTACTTTTGTGTTGTGTCAGAAACAGGAGTGAGAAAGAGAGCAATTATAG TTAAGGAAGATGATGGAGTGGAAGATTCAACAACAACAAGATCATCATGATCATGATCA TGATAAACATGATCATCAGATGGAGATGGGAATATTGCAGAAAAAGGAAGATCAAGAGAATT CAACTATTGTTGGGGTTGTTCTCTCTCCAAATCCAAAAGATGGGTAGGGTTAATGTACGTGAA GGTTATGACTGACGAGCAAATGGAGCTTCTCAGGCGCCAGATCTCCGTCTACGCCACCATCTG CGAGAAGCTTGCTGACATGTTCAACTCCTTCTCTTCCCAACATGATCTCGCTGGAATGAAGCT GGGAAGCTTCACTATGATCCACTAATGACATATGGTGGCCACAAGATCACTGCAAGGCAAA GGTGGTCACCAACGCCTGTGCAGCTTCAAACCTAGAGAAAAATTTTCGATGAAGGGAATGGG ACTCCGAGCAAGCAGAAGATCAAAGAGATAACCGTAGAGCTGAGTCAACATGGCCAAATTTCT GGAAACCAATGTCTACAATTGGTTCCAGAACAGGAGAGCTCGTTCCAAGAGAAAGCAATCCG TTCCAGCTCCAACTGCGTGGAAGTAGAAGTAGAGACCGAACTCAAGAGCATTTGAAGGAC AGTGCAAAACCAGAAGATACTGAATTCTGTGACAACCTCGGCTGCAGGGGAGGATCATGTGTA TTTTCCAAGTCCTGAGGAGTTGATCAACTGGTCCGCGAATGGAAGTGCCATTGAGCTGCATGC CTTGCCCTTCATGGGGAGGATTTGATGTGGATGAGCTCTAGTTGGGATTCTTATATGGTCTCTGTTG TTGCTCTTCAGCTGATCTTATTATAAATTTCTGTTCAATGGATTAGAATTTTACATTAGAGAGAA CTTCATTATTAAAATATTATAGTCATAGGTAATTTAGCAATCACTGACAATCTGACATGTGGACT TATAAAATCGTATGTGTCAGACTTGGCTAAAAAACTAACTGATAGCGAATGTCTTTGATGTTA GCATGTATATGTAATCTTTACAATCAGATTATGATGAACTGGGTCTATTGAAAAACC </p>

Table S5. The species information used to reconstruct the phylogenetic tree. Letter A represents the ancient clade, I represents the intermediate clade, and W represents the WUS clade.

Species	The numbers of WOX gene	A	I	W
<i>Ostreococcus lucimarinus</i>	1	1	0	0
<i>Physcomitrella patens</i>	3	3	0	0
<i>Selaginella moellendorffii</i>	8	7	1	0
<i>Picea abies</i>	18	4	8	6
<i>Amborella trichopoda</i>	9	1	2	6
<i>Oryza sativa. ssp. Japonica</i>	14	1	6	7
<i>Brachypodium distachyon</i>	13	2	5	6
<i>Populus trichocarpa</i>	18	3	4	11
<i>Medicago truncatula</i>	19	3	7	9
<i>Morus notabilis</i>	11	2	2	7
<i>Broussonetia papyrifera</i>	10	2	2	6
<i>Prunus persica</i>	10	2	2	6
<i>Cucumis melo</i>	10	2	2	6
<i>Vitis vinifera</i>	11	3	2	6
<i>Gossypium raimondii</i>	20	4	4	12
<i>Arabidopsis thaliana</i>	15	3	4	8
<i>Solanum lycopersicum</i>	10	1	2	7
<i>Theobroma cacao</i>	11	2	2	7
<i>Utricularia gibba</i>	17	1	4	12
Total number	228	47	59	59

Table S8. The primer sequences of *BpWOXs* used in qRT-PCR.

Gene Name	Forward primer (5'-3')	Reverse primer (5'-3')
<i>BpWUS</i>	TGAGCTAACCTCAACTCCTACG	TCACACAATCATCATAAATAGAGCATG
<i>BpWOX1</i>	GCCGGCCTACAGATCAGAAAATG	CTTCAATGGTAGGAACTCAAAGAACTG
<i>BpWOX2</i>	ACTGCTACAACTCTTGAGCTATTTCC	ACTCACGTCCTTTATTCTTACTTTGC
<i>BpWOX3</i>	GAAGGAGAACTCCATGATGAGAATG	GAGAGCCATGCAGATAGGAATTTAG
<i>BpWOX4</i>	CTAGTCCGTACAAGAGGAAGTGCC	CATATACCGGCCACATAAACTAAAG
<i>BpWOX5</i>	GCATTAAAGCGGGACATAACATG	GTCCTCTCTTAGAACGAAATCCTTGT
<i>BpWOX7</i>	CGTTAAGTAATAACTGAAGCAGCATCT	ATGGGAGACTGGCATTACAAATAAG
<i>BpWOX8</i>	GACGACGGTGGCAAGGACTAC	AGGCTGGAGGGAGTGAAGAGTG
<i>BpWOX9</i>	ATGCGGAAAACTACAGCATGAC	CAAAACCATGACATTCTACTCCCTG
<i>BpWOX10</i>	CTAGAAAGTAGAGACCGAAACTCAAGAGC	CTGAAGAGCAACAACAGGACCAT
<i>BpGAPDH</i>	TCAACATCATTCCTAGCAGTACCG	AGTCAGTGGAACCACGTCATC

Table S9. The restriction enzymes and primers used to construct the recombinant expression vectors for subcellular localization (SL) and transactivation activity assay (TA).

Gene Name		Forward primer (5'-3')		Reverse primer (5'-3')
<i>BpWUS</i> -SL	<i>Xba</i> I	GCTCTAGAATGGAACCTCAAC	<i>Sal</i> II	ACGCGTCGACAAGCGAATCCGG
		AACAAAACCAATC		TGACCCGCTAG
<i>BpWOX1</i> -SL	<i>Xba</i> I	GCTCTAGAATGTGGATGATGG	<i>Sal</i> II	ACGCGTCGACATTCTCAATGGT
		GCTATAACGACG		AGGAACTC
<i>BpWOX2</i> -SL	<i>Xba</i> I	GCTCTAGAATGTGGACGATGG	<i>Pst</i> I	AACTGCAGGAGGAAGAAACGCA
		GTCGTAGTACTA		ACGAGAGG
<i>BpWOX3</i> -SL	<i>Xba</i> I	GCTCTAGAATGTCTCCAGCAG	<i>Sal</i> II	ACGCGTCGACGATGGTCATCGTT
		CCAATTCATCAAG		GGGGTGGAAATG
<i>BpWOX4</i> -SL	<i>Xba</i> I	GCTCTAGAATGGGAAGAAGTA	<i>Sal</i> II	ACGCGTCGACTCTCATGCCTTCC
		GCTGCAGCATGA		GGGTGTAA
<i>BpWOX5</i> -SL	<i>Xba</i> I	GCTCTAGAATGGAAGATCATG	<i>Sal</i> II	ACGCGTCGACGACAAAGCTTAA
		GCATGTCAGGCT		GCGCAGAT
<i>BpWOX7</i> -SL	<i>Xba</i> I	GCTCTAGAATGGAAGATCATC	<i>Sal</i> II	ACGCGTCGACGTTATTACTTAAC
		ACCATCATCAAGG		GTGTCT
<i>BpWOX8</i> -SL	<i>Xba</i> I	GCTCTAGAATGGCTTCATCA	<i>Sal</i> II	ACGCGTCGACAAATTAGATAATA
		ACAGACACTGGC		AAAAGCACCAGG
<i>BpWOX9</i> -SL	<i>Xba</i> I	GCTCTAGAATGATGGGGTGGG	<i>Sal</i> II	ACGCGTCGACTCCCGTCATGCTG
		AGAAGCAGGAG		TAGTTTTTC
<i>BpWOX10</i> -SL	<i>Sma</i> I	TCCCCCGGGATGATGGAGTGG	<i>Sal</i> II	ACGCGTCGACGAGCTCATCCACA
		AAGATTCAACAAC		TCAAATCCTC
<i>BpWUS</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGGAACCTCA	<i>Sal</i> II	ACGCGTCGACGGTTAAAGCGAAT
		ACAACAAAACCAATC		CCGGTGACCCGCTAG
<i>BpWOX1</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGTGGATGAT	<i>Sal</i> II	ACGCGTCGACGGTCAATTCTTCA
		GGGCTATAACGACG		ATGGTAGGAACTC
<i>BpWOX2</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGTGGACGAT	<i>Pst</i> I	AACTGCAGCCTCAGAGGAAGAA
		GGGTCGTAGTACTA		ACGCAACGAGAGG
<i>BpWOX3</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGTCTCCAGC	<i>Sal</i> II	ACGCGTCGACGGTTAGATGGTCA
		AGCCAATTCATCAAG		TCGTTGGGGTGGAAATG
<i>BpWOX4</i> -TA	<i>Eco</i> RI	CCGGAATTCCGGATGGGAAGA	<i>Sal</i> II	ACGCGTCGACGGTCATCTCATGC
		AGTAGCTGC		CTTCCGGGTGTAA
<i>BpWOX5</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGGAAGATCA	<i>Sal</i> II	ACGCGTCGACGGTTAGACAAAG
		TGGCATGTC		CTTAAGCGCAGAT
<i>BpWOX7</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGGAAGATCA	<i>Sal</i> II	ACGCGTCGACGGTCAGTTATTAC
		TCACCATCATCAAGG		TTAACGTTGTCT
<i>BpWOX8</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGGCTTCATC	<i>Sal</i> II	ACGCGTCGACGGCTAAATTAGAT
		AAACAGACACTGGC		AATAAAAGCACCAGG
<i>BpWOX9</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGATGGGGTG	<i>Sal</i> II	ACGCGTCGACGGTCATCCGTC
		GGAGAAGCAGGAG		TGCTGTAGTTTTTC
<i>BpWOX10</i> -TA	<i>Sma</i> I	TCCCCCGGGGATGATGGAGTG	<i>Sal</i> II	ACGCGTCGACGGCTAGAGCTCAT
		GAAGATTCAACAAC		CCACATCAAATCCTC

