

Figure S1: Multiple sequence alignment with the amino acid sequences of HD in different classes of *Phyllostachys edulis* (Pe), *Arabidopsis thaliana* (At), and *Oryza sativa* (Os).

| | | | | | |
|------------|--------------------|-----------------------------|---------------------------------|--------------------------------|---------|
| PeHB011 | WRPQRGIPERAVTILR | IAWLFEEHFLHPYPSD | DVKHILARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB014 | WRPQRGIPERAVAILR | IAWLFEEHFLHPYPSD | DVKHILARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB015 | WRPQRGIPERAVAILR | IAWLFEEHFLHPYPSD | DVKHILARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB018 | WRPQRGIPERAVTILR | IAWLFEEHFLHPYPSD | DVKHILARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB023 | WRPQRGIPERAVAILR | IAWLFEEHFLHPYPSD | DVKHILARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB024 | WRPQRGIPERAVASVL | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB031 | WRPQRGIPERAVASVL | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB032 | WRPQRGIPERAVAILR | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB037 | WRPQRGIPESAVSVL | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB039 | WRPQRGIPESAVSVL | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB042 | WRPQRGIPERAVSVL | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB047 | WRPQRGIPERAVSVL | IAWLFEEHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| AT1G19700 | WRPQRGIPENSVLSVL | IAWFEHHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| OS124g3950 | WRPQRGIPENSVLSVL | IAWFEHHFLHPYPTDGD | KQMLARQFGLSRSQVS | NWFLNARVREWK | |
| PeHB093 | KRTIIMNDQVNEIEKALI | YEPMEMHNKATLQAAWEAKL | SQGSQEITSSQLQNWLNRRKAKL | A | BEL |
| PeHB070 | KRTIIMNDQVNEIEKALI | YEPMEMHNKATLQAAWEAKL | SQGSQEITSSQLQNWLNRRKAKL | A | BEL |
| PeHB017 | KRKRAKGKLPGDTTTL | LKQWOQOHPWPYTEDDKAKVEETG | GLOKQIENNWFINQRKRWNH | | NDX |
| PeHB057 | KRKRAKGKLPGDTTTL | LKQWOQOHPWPYTEDDKAKVEETG | GLOKQIENNWFINQRKRWNH | | NDX |
| PeHB058 | KRKRAKGKLPKDARS | MDDWNTNHYWPWPYPTEDDKAKVEETG | VRLAAMGLDPKQIENNWFINQRKRWNH | | NDX |
| PeHB074 | KRKKGKLPKEAROKL | LHWWELHYWPWPYSTEKKIALAE | STGLDQKQIENNWFINQRKRWNH | | NDX |
| PeHB104 | KKKKKGKLPKPRDARQKL | LHWOLHYWPWPYSTEKKIALAE | STGLDQKQIENNWFINQRKRWNH | | NDX |
| PeHB103 | KRKRAKGKLPGDTTTS | TKSWSOSHARWPWPYPTPSTES | FHFHRHVTVLARSRSRSILLRAERA | | KNOX |
| PeHB104 | KKKKKKSLPDRARQKL | LHWOLHYWPWPYSTEKKIALAE | STGLDQKQIENNWFINQRKRWNH | | KNOX |
| PeHB104 | KKKKKKSLPDRARQKL | LHWOLHYWPWPYSTEKKIALAE | STGLDQKQIENNWFINQRKRWNH | | KNOX |
| AT1G22380 | QMSPLQLETLKIVS | EKAQPEATRAELSERK | TDQKQIENNWFINQRKRWNH | | KNOX |
| PeHB009 | VMKPYLEVILEKTNT | EDPYPDETLRAELSVKL | GTDQKQIENNWFINQRKRWNH | | KNOX |
| PeHB034 | TKKSPQIQLQMEVCP | EVQVPKPDEMDAYEATSV | GQASQVRIWEKEERRRER | | DDT |
| PeHB036 | VMKPYLEVILEKTNT | EDPYPNETLRLSVKL | GTDQKQIENNWFINQRKRWNH | | DDT |
| PeHB041 | TKKSPQIQLQMEVCP | EVQVPKPDEMDAYEATSV | GQASQVRIWEKEERRRER | | DDT |
| PeHB082 | TKKSPQIQLQMEVCP | EVQVPKPDEMDAYEATSV | GQASQVRIWEKEERRRER | | DDT |
| PeHB093 | TKKSPQIQLQMEVCP | EVQVPKPDEMDAYEATSV | GQASQVRIWEKEERRRER | | DDT |
| AT1G28420 | QMSPLQLETLKIVS | EKAQPEATRAELSERK | TDQKQIENNWFINQRKRWNH | | DDT |
| PeHB021 | QDMVLPVSSSRQVER | LDYKRYDEAYGKRESDSSDDEEWS | GNSTPQKGNEQESTDSDLAE | | PHD |
| PeHB022 | QDMVLPVSSSRQVER | LDYKRYDEAYGKRESDSSDDEEWS | GNSTPQKGNEQESTDSDFAE | | PHD |
| PeHB035 | QDLVLPVSSGRREAQ | LYYYKKYDYLRLH... | EHFKIDQYPSR...AVKESLAQELGL | | PHD |
| PeHB079 | SVEGFPAIDLDEKVQ | PQRKQFRIPP... | AAVQVLRKVFAENELPAR... | | PHD |
| PeHB101 | SVKGFDPDVISDEKVQ | PQRKQFRIPP... | AAVQVLRKVFAENELPAR... | | PHD |
| PeHB102 | LDDDGAGVSERRRNRV | ERLDYKKYDDEEYDNVP | TSQDDDWDTKARMGK... | | PHD |
| AT3G19510 | DCWSHWRVYLQDGMLH | PGQCEGAGHOSCI | RDVLIVGCAEGITVGDKEVMS | NSDRIMSNVNI | PINTOX |
| PeHB060 | DCWSHWRVYLQDGMLH | PGQCEGAGHOSCI | RDVLIVGCAEGITVGDKEVMS | NSDRIMSNVNI | PINTOX |
| PeHB114 | DCWSHWRVYLQDGMLH | PGQCEGAGHOSCI | RDVLIVGCAEGITVGDKEVMS | NSDRIMSNVNI | PINTOX |
| PeHB000 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB008 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB012 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB016 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB020 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB022 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB028 | GERKERRLAVFOVRALE | RSFETDNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB030 | TERKERRLAVFOVRALE | XNNELGKNGLEPERKQL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB046 | GERKERRLAVFOVRALE | XNNELGKNGLEPERKQL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB050 | GERKERRLAVFOVRALE | XNNELGKNGLEPERKQL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB061 | GERKERRLAVFOVRALE | XNNELGKNGLEPERKQL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB099 | GENRURAAFOVRALE | RSFSEADNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| Os08g37580 | GENRURAAFOVRALE | RSFSEADNKLDPDKRARI | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP I |
| PeHB001 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB025 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB029 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB040 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB059 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB065 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB067 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB081 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB087 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP II |
| PeHB025 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | ZIP III |
| PeHB007 | KVVRYTPQEVALERV | YAHCPKPTTSRQOLRE | PLIQLANIEQOIQKVWEQNRRCRD | KORKE | ZIP III |
| PeHB038 | KVVRYTPQEVALERV | YAHCPKPTTSRQOLRE | PLIQLANIEQOIQKVWEQNRRCRD | KORKE | ZIP III |
| PeHB060 | KVVRYTPQEVALERV | YAHCPKPTTSRQOLRE | PLIQLANIEQOIQKVWEQNRRCRD | KORKE | ZIP III |
| PeHB088 | KVVRYTPQEVALERV | YAHCPKPTTSRQOLRE | PLIQLANIEQOIQKVWEQNRRCRD | KORKE | ZIP III |
| PeHB095 | KVVRYTPQEVALERV | YAHCPKPTTSRQOLRE | PLIQLANIEQOIQKVWEQNRRCRD | KORKE | ZIP III |
| Os10g33960 | KVVRYTPQEVALERV | YAHCPKPTTSRQOLRE | PLIQLANIEQOIQKVWEQNRRCRD | KORKE | ZIP III |
| PeHB000 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB043 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB044 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB045 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB046 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB047 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB048 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB049 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB050 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB051 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB052 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB053 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB054 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB055 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB056 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB057 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB058 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| PeHB059 | RRKRYHHTHTPROJQ | QKELKECPHIDPMQARQLS | REL...GIEQQLKFWFONRRT | KOL | ZIP IV |
| AT1G05230 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | PLINC |
| Os10G42400 | KKLRILITQEQALIEDR | FEKEHSLNRRKAKLQNL | ARDIGLPROVAVWFONRRAWRWT | KOL | PLINC |
| PeHB052 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| PeHB053 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| PeHB065 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| PeHB075 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| PeHB089 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| PeHB096 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| AT1G14440 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| Os04g35500 | RHKTFTEAQKARMGL | PEEVGQALQKLE | ...DAVVGQRFQEVVKVWEQNRRCRD | KORKE | SAWADEE |
| PeHB084 | DYQEYLNRLNRLYRQ | QKQVLSCKVGSNLTYE | EVLSFHAAEKAQQLPRELVVPVLMQVOYS | | WOX |
| AT3G1830 | NGGPAAFRFLP | EVTEMEAI | LLQHNTAMPGRHLLEALADKFESPERGKVVV | OF KOLWNW | WOX |
| PeHB003 | RWPTTANQLOI | LENIFD | QGNGTPSKQKIEI | T. AELQSQHQISETNVNVWFQNRKARSKR | WOX |
| PeHB075 | RWPTTANQLOI | LENIFD | QGNGTPSKQKIEI | T. AELQSQHQISETNVNVWFQNRKARSKR | WOX |
| PeHB010 | RWPTTANQLOI | LENIFD | QGNGTPSKQKIEI | T. VELSQHQISETNVNVWFQNRKARSKR | WOX |
| PeHB019 | RWPTTANQLOI | LENIFD | QGNGTPSKQKIEI | T. VELSQHQISETNVNVWFQNRKARSKR | WOX |
| PeHB066 | OLLGTAAAG | AGAPPPLCTT | PPAPHSCLDQQAAA | AHUTTAPAAATYSSYYPFGAAPASRCT | WOX |
| PeHB101 | RWNPSPKQKIEI | KVLEM | YRGGMRTPNALQIERI | T. EELGKYGRIEGNNVYWFQNRKARSKR | WOX |
| PeHB070 | RWNPSPKQKIEI | KVLEM | YRGGMRTPNALQIERI | T. EELGKYGRIEGNNVYWFQNRKARSKR | WOX |
| Os01g02070 | RWTPTPMQLOI | LENIFD | QGNGTPSKQKIEI | T. AEQSQHQISETNVNVWFQNRKARSKR | WOX |

Figure S2: Phylogenetic tree based on the amino acid sequences of HB proteins of *Phyllostachys edulis* (Pe), *Arabidopsis thaliana* (At), and *Oryza sativa* (Os).

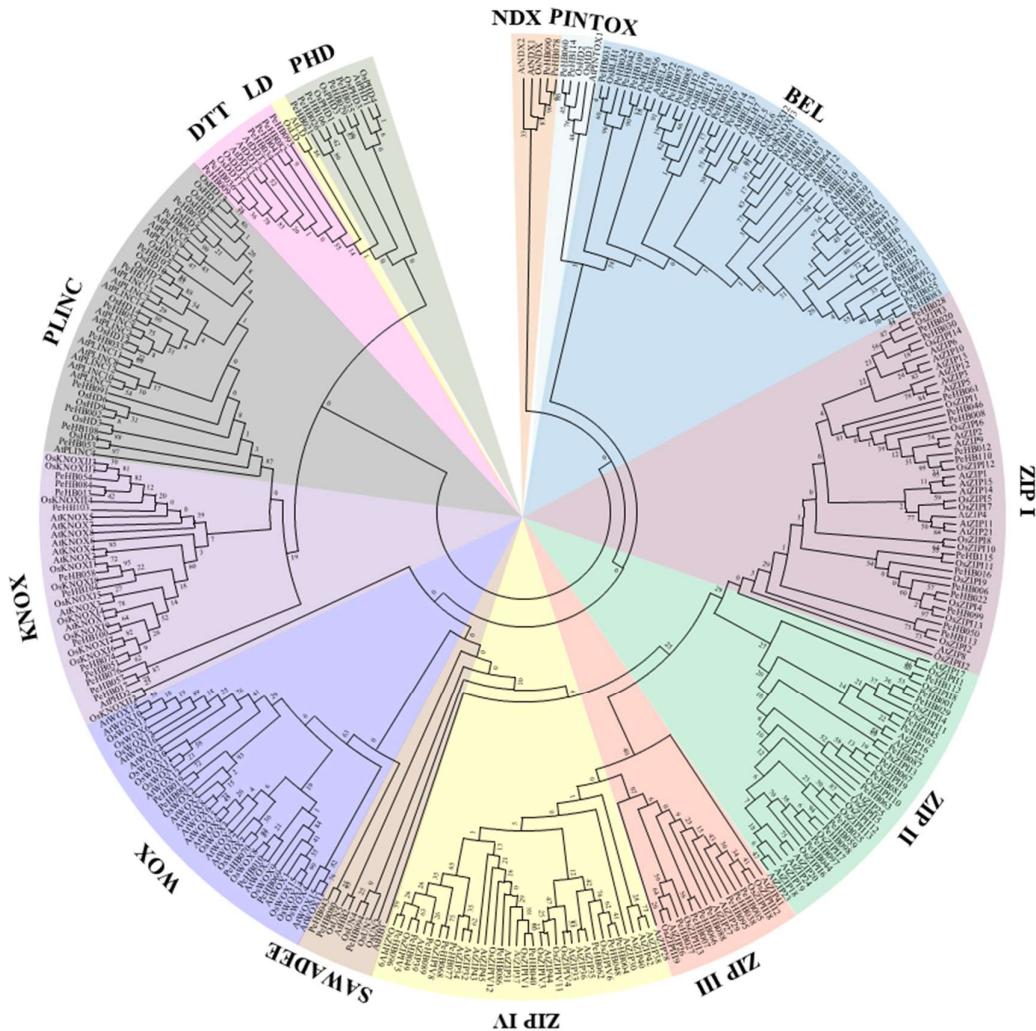


Figure S3: Gene structures of *PeHBs* in moso bamboo.

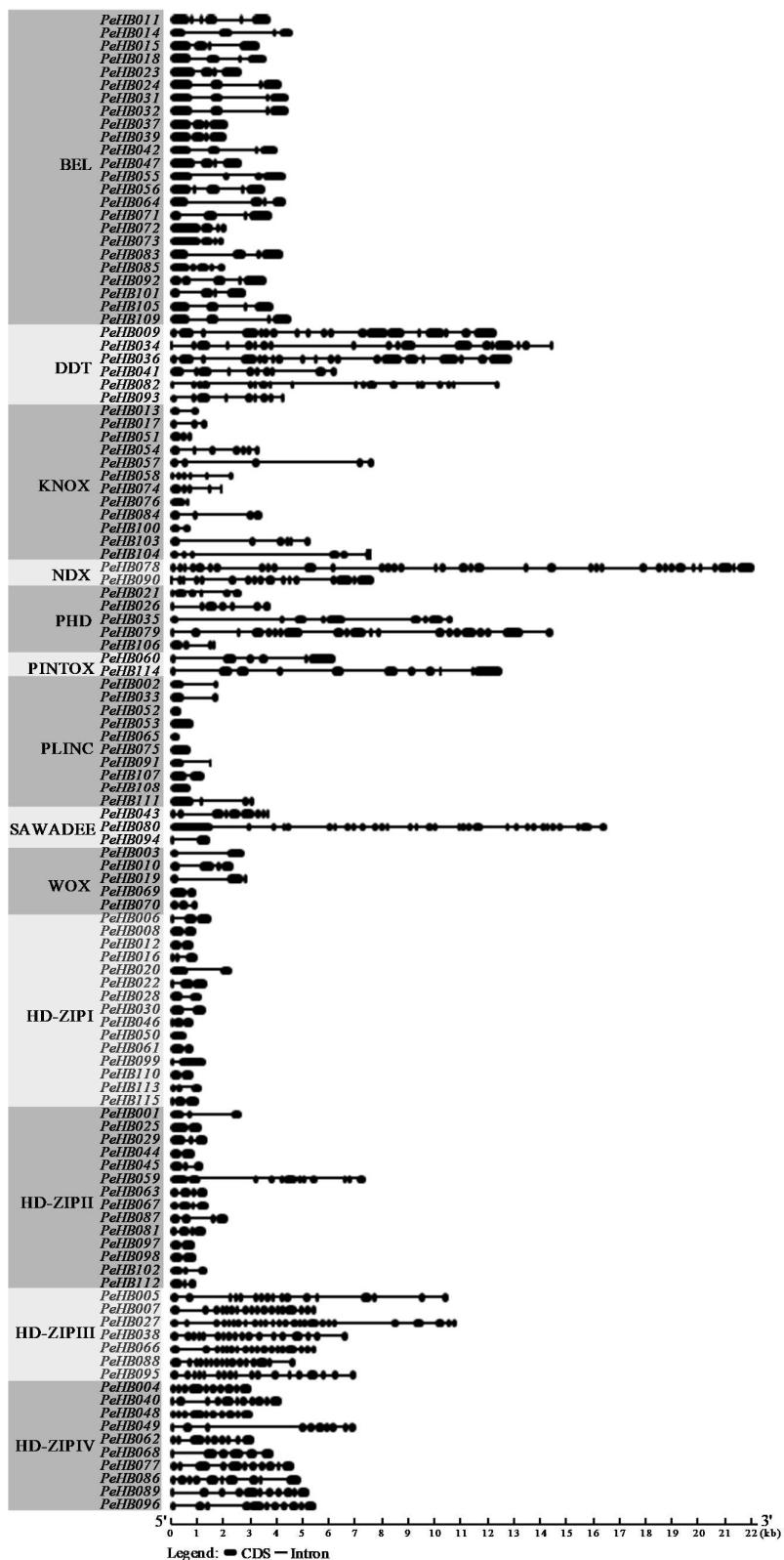


Figure S4: Information of the 20 motifs in PeHBs.

| | E-value | Sites | Width | | E-value | Sites | Width |
|----------|-----------|-------|-------|----------|----------|-------|-------|
| motif 1 | 3.4e-1062 | 95 | 21 | motif 11 | 6.1e-171 | 15 | 36 |
| motif 2 | 6.8e-578 | 29 | 29 | motif 12 | 1.8e-169 | 8 | 50 |
| motif 3 | 5.4e-595 | 21 | 50 | motif 13 | 9.1e-168 | 18 | 29 |
| motif 4 | 3.3e-569 | 46 | 29 | motif 14 | 2.5e-166 | 7 | 50 |
| motif 5 | 1.7e-239 | 14 | 29 | motif 15 | 8.1e-152 | 23 | 21 |
| motif 6 | 2.3e-288 | 15 | 50 | motif 16 | 3.0e-173 | 12 | 47 |
| motif 7 | 6.3e-224 | 10 | 50 | motif 17 | 2.7e-143 | 13 | 35 |
| motif 8 | 5.4e-215 | 30 | 29 | motif 18 | 3.4e-153 | 7 | 50 |
| motif 9 | 5.0e-203 | 16 | 29 | motif 19 | 1.7e-136 | 15 | 25 |
| motif 10 | 1.4e-201 | 7 | 50 | motif 20 | 1.4e-135 | 8 | 50 |

Figure S5: GO analysis of *PeHBs*.

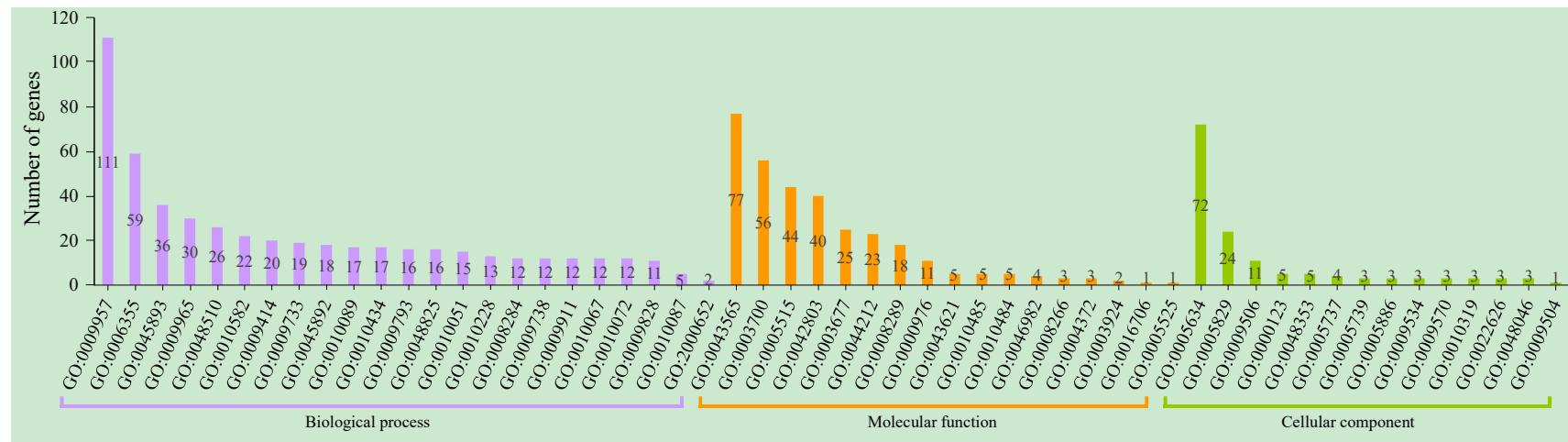


Figure S6: Transactive analysis of four members belonging to KNOX class in yeast. The control vectors and fusion constructs of four KNOX genes were transformed into AH109 yeast cells respectively, and inoculated onto SD/-Trp and SD/-Ade /-Leu/-Trp/X-a-GAL plates for further selection.

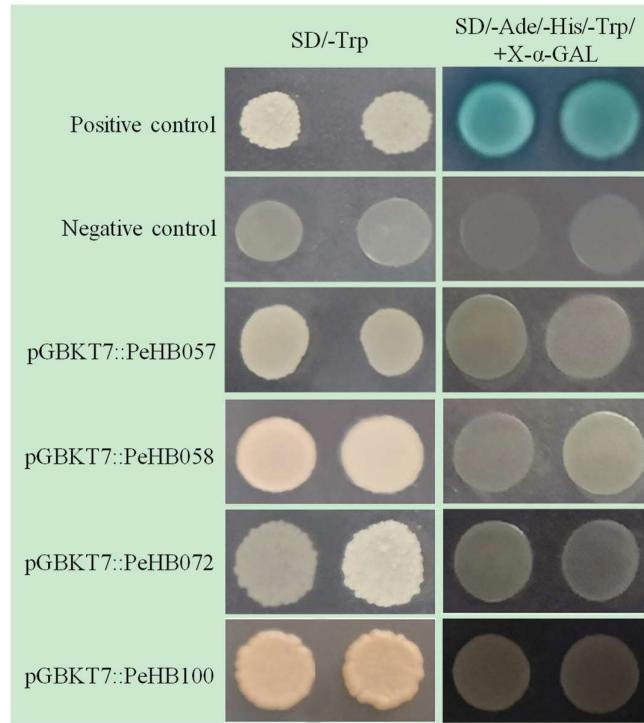


Figure S7: The content changes of cellulose (a) and hemicellulose (b) in winter bamboo shoots during storage. Asterisks indicated a significant difference between the storage shoots and the fresh shoots (* $p < 0.05$).

