

1 TCCACTATATTAGAATCAGAAGATTCACTGAAAGAATTAGTTTATTTGGTAGCTTGTAGGGTTCTTCACCTGTATATTGTAACACG
 91 CCTTATATGATTCAATTCTTGTGTTTATCTGGTCTCCCTAATCCCTAACCTACAATTATAAAAACATTGATTATGTACTT
 1 M S N P P F G S P L Y N E E D D F S F P F Q L P N N D
 181 TTTGGTTTATGCTAACCCCTCTTGGAAAGTCCCCCTATAATGAGGAAGATGACTCTCTTCCCTTTCAACTACCAATAATGAT
 28 D P I N F P P L I E D H N S L P S T S S L P M S E Y Q D L S
 271 GATCCCACATCAATTCCACCTCTATTGAAGATCATAATTCTTGCCCTACATCTTCACTACCCATGTCAAGAGTACCAAGATTAAAGC
 58 M F I D D H H E E D H T L I G Q Y N P N I Q E T I N W A C P
 361 ATGTTTATTGATGATCATGATGAAGAAGATCATACACTAATAGGGCAATAATCCCAACATACAAGAGACAATAATTGGGCATGTCCA
 88 I E Q V G L N Q V H Q G A I E G S S I G Y H E M Q M Q Y Y C
 451 ATTGAACAGGGTTGGCTTAATCAGGTGCACCGGGTCAATAGAAGGCTCATCTATAGGATATCATGAGATGCAAATGCAAACTACTGT
 118 P T T T T T T T T I A V A A S A A T M M I N D Q A A P P P G
 541 CCTACCACCAACCAACCCACCAACCCATCGCTGCTGCCCGCCACCATGATGATAATGATCAAGCTGCTCCACCACTGGT
 148 V C R G G G G E G S S K K L D H N A K E K V R R R M K L N
 631 GTTGTCTGYGTGGCGGGGTGGTAAGGATGCTAGCAAGAACCTGATCATATGCTAAGGAGAAAGTAGGGAGATGAAGCTTAAT
 178 E T Y L A F R S L L P D S K R A K K R W S A P Y I I D R A L
 721 GAAACATATTGGCTTTCGCTCCCTGCTACCAAGATTCTAAAGAGCTAACAGAAAGATGGAGTCACCATATAATTGATAGAGCTTG
 208 D Y I P Q L Q A E I E K L T L E K K N M L T M L E K K Q Q L
 811 GATTACATCCCTCAGCTCAAGCTGAGATAGAGAACGTAACACTCGAAAGAACATGCTAACATGCTAGAGAAAAAGCAACACTC
 238 V E R S K D H N N A T E D K K T L T V S M N E V K I G E V I
 901 GTTGAACGCTAGCAAGGATCATAATAATGCAACTGAGATAAGAAGACGCTAACAGTATCGATGATGAAGTAAATTGGTGAAGTGTATA
 268 V Q I C E Q N N K I G M L P T L I E K L E G E S M Q I V G A
 991 GTCCAGATTGTGAACAGAATAACAAAATTGGTATGTTACCTACGGTATCGAAAAACTCGAAGGAGAAAGCATGCAAATCGTAGGTGCC
 298 S S Q R A C E D R S C F H I H V Q M G E N P V E A D Y V A I
 1081 TCTTCACAAACGTGATGTGAAGATAGATGCTGCTCCATATTGTTCAAATGGTGAAAACCCAGTGGAGCTGATTATGTTGCAATT
 328 L H K K I I S W L S *
 1171 TTACATAAGAAGATAATCTCTGGCTGCTTAGTACGGAGAACATGAGTCAGTGTACTCCGACACCTTCATTACATTGTTCAACATGGGG
 1261 TGTCGTGCTGCTGCCGACATGACACTTACTGAAATCTTCACTAAACTAAACTGACTTTTCTACAAAATAGTCGAGTCGGAC
 1351 ACTTGGACACCCGTATTCGAGTAAACACTCAATCCGTCTAAAGTAACATAGGGAGAAAATTGTTCAAGGAAAAACTAATATGAGCTCT
 1441 TTGAATATGAAATTGAAATCTCTCTTACTTTCTGTAGAGAAGTAGAAATACAGCGTGCCTCAATTTCATGGA

 Basic  helix  loop

Figure S1. Sequence analysis of a cDNA encoding a *BvbHLH93* isolated from the sugar beet. Nucleotide and deduced amino acid sequence of the *BvbHLH93*.

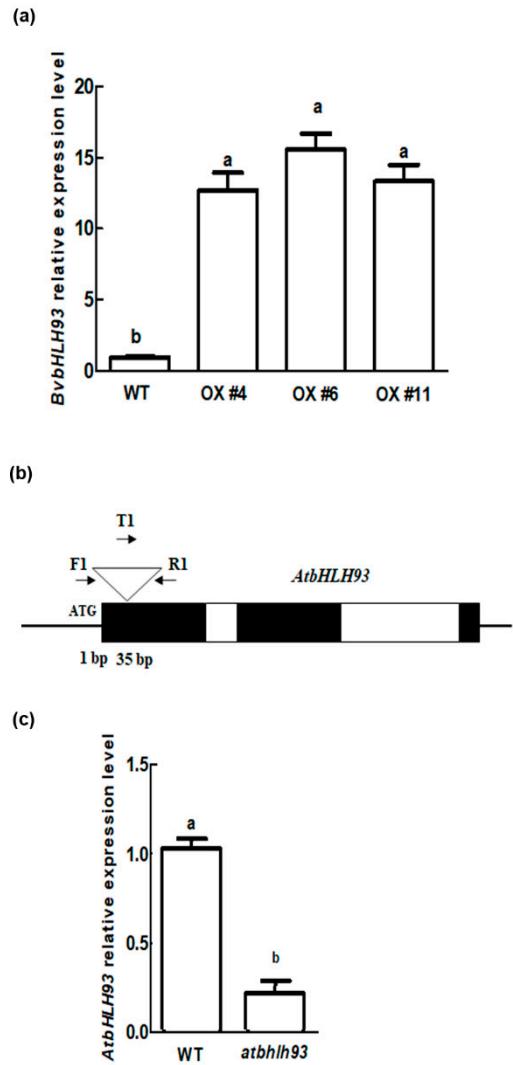


Figure S2. Identification of *atbhlh93* mutant and overexpression of *BvbHLH93* in *Arabidopsis*. (a) QRT-PCR analysis of the expression levels of the overexpressed *BvbHLH93* (OX4, OX6 and OX11) in *Arabidopsis*; (b) Structure of the *AtbHLH93* gene. **The T-DNA insertion site was at 35 bp downstream of the start codon. The primers used to identify the T-DNA insertion were marked with arrows (T1, R1 and F1); (c) Real-time PCR analysis of the expression levels of *AtbHLH93* in the *atbhlh93* mutant. Different letters indicate significant difference at $p < 0.05$. Three biological replicates were performed. Please refer to supplementary materials for the primers used.**

Table S1. The list of primers used in this article

Primer name	Primer sequence
<i>BvbHLH93-F</i> (Full length cDNA)	5-TCCACTATATTAGAATCAGAAGATT-3
<i>BvbHLH93-R</i> (Full length cDNA)	5-TCCATGAAAATTGAAGGCACCGCTGT-3
<i>BvbHLH93-F</i> (GFP vector construct)	5-CCGAAATTCATGTCTAACCCCTCCTTTGG-3
<i>BvbHLH93-R</i> (GFP vector construct)	5-ACGCGTCGACAGCCAAGAGATTATCTTC-3
<i>BvbHLH93-F</i> (Yeast vector construct)	5-GAATTCCCGGGATCATGTCTAACCCCTTTGG-3
<i>BvbHLH93-R</i> (Yeast vector construct)	5-GCAGGTGCGACGGATCCTAACAGACAGCCAAGAGAGATTA-3
<i>BvbHLH93-F</i> (Overexpression vector construct)	5-CTAGTCTAGAATGTCTAACCCCTCCTTTGG-3
<i>BvbHLH93-R</i> (Overexpression vector construct)	5-ACGCGTCGACAGACAGCCAAGAGACTGAC-3
<i>BvbHLH93-F</i> (QRT-PCR)	5-AAGCAGCAACTCGTTGAACG-3
<i>BvbHLH93-R</i> (QRT-PCR)	5-CATGCACGTTGTGAAGAGGC-3
<i>Bv18S rRNA-F</i> (QRT-PCR)	5-CCCCAATGGATCCTCGTTA-3
<i>Bv18S rRNA-R</i> (QRT-PCR)	5-TGACGGAGAATTAGGGTTCG-3
<i>AtRbohD (At5g47910)-F</i> (QRT-PCR)	5-TCAGGGACGACTCGGTGG-3
<i>AtRbohD (At5g47910)-R</i> (QRT-PCR)	5-GTTTATCGAAACGTTGGTC-3
<i>AtRbohF (At1g64060)-F</i> (QRT-PCR)	5-GTTCGATGCATTGAGTAG-3
<i>AtRbohF (At1g64060)-R</i> (QRT-PCR)	5-TTTAATCTTGATACTGTTATT-3
<i>AtbHLH93 (At1g71200)-F</i> (QRT-PCR)	5-TGGCTCAAGTAGTGGTGCAG-3
<i>AtbHLH93 (At1g71200)-R</i> (QRT-PCR)	5-CCCCAATTCAAGAACCGAGA-3
<i>β-Actin (At3g53750)-F</i> (QRT-PCR)	5-GAGGCTCCTCTTAACCCAA-3
<i>β-Actin (At3g53750)-R</i> (QRT-PCR)	5-TACAATTCCCGCTCTGC-3
<i>SOD (At5g18100)-F</i> (QRT-PCR)	5-AACGAGGAAGAGCGTCATG-3
<i>SOD (At5g18100)-R</i> (QRT-PCR)	5-GCGTTCCAGTTGATTG-3
<i>SOD (At4g25100)-F</i> (QRT-PCR)	5-ATGAGAAGTTCTATGAAGAG-3
<i>SOD (At4g25100)-R</i> (QRT-PCR)	5-GTCTTATGTAATCTGGT-3
<i>POD (At1g14550)-F</i> (QRT-PCR)	5-CCATAGGACAATCTCAATGC-3
<i>POD (At1g14550)-R</i> (QRT-PCR)	5-TGATCGGTTACTAATAG-3

POD (At5g66390)-F (QRT-PCR) 5-CTCACTAAGTTCAAGCGTC-3

POD (At5g66390)-R (QRT-PCR) 5-GAATAGGGTCTGGTCACC-3
