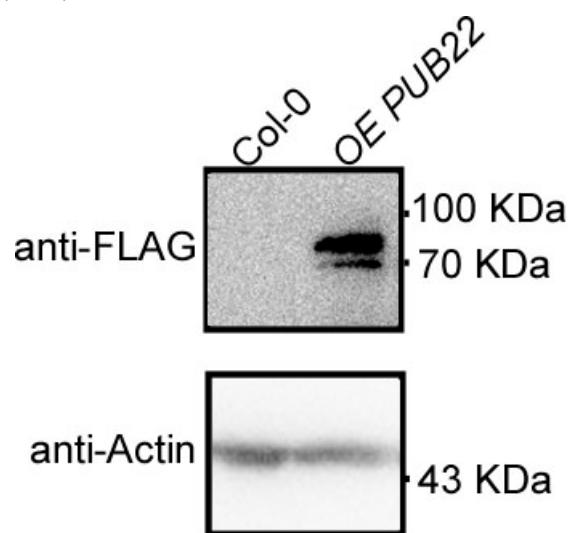
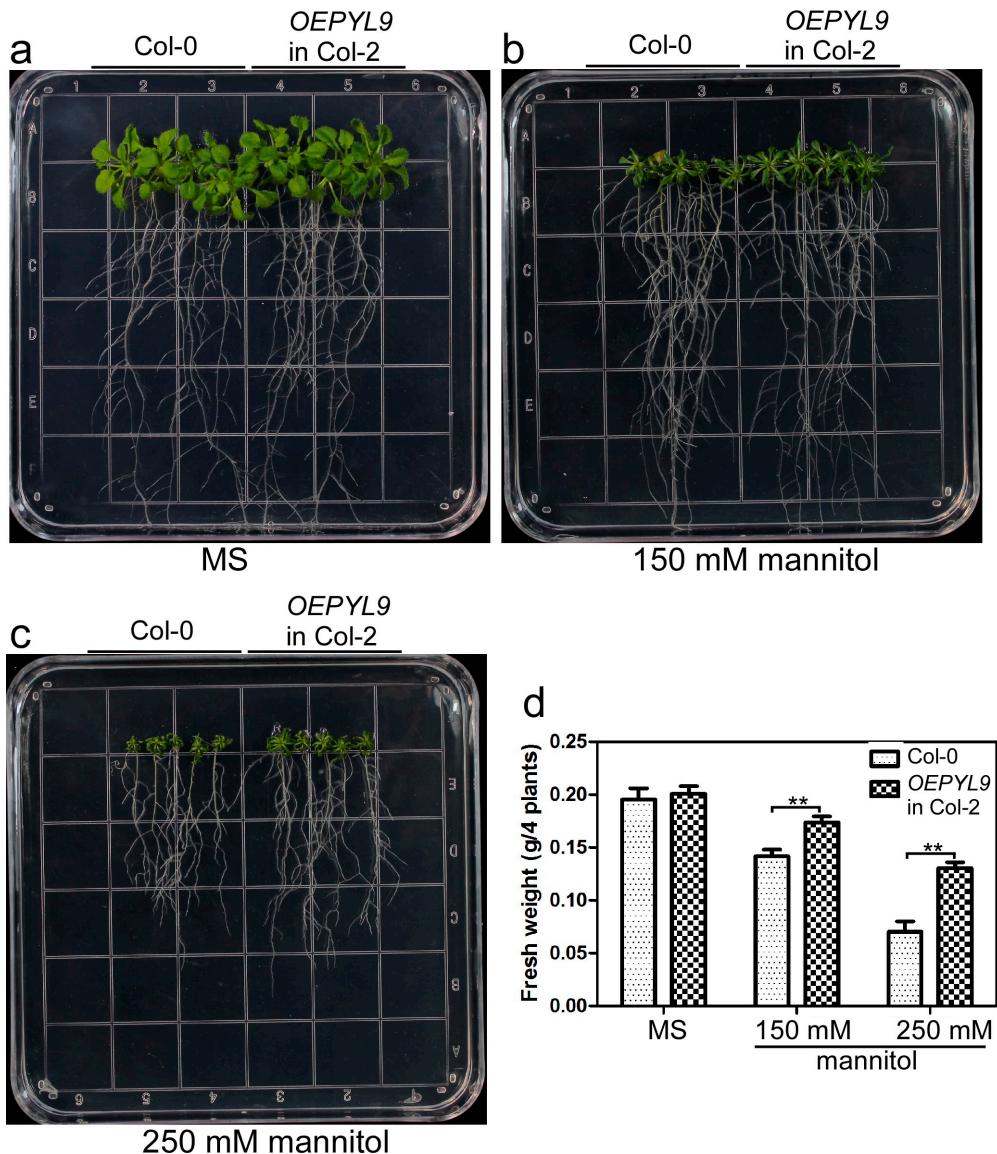


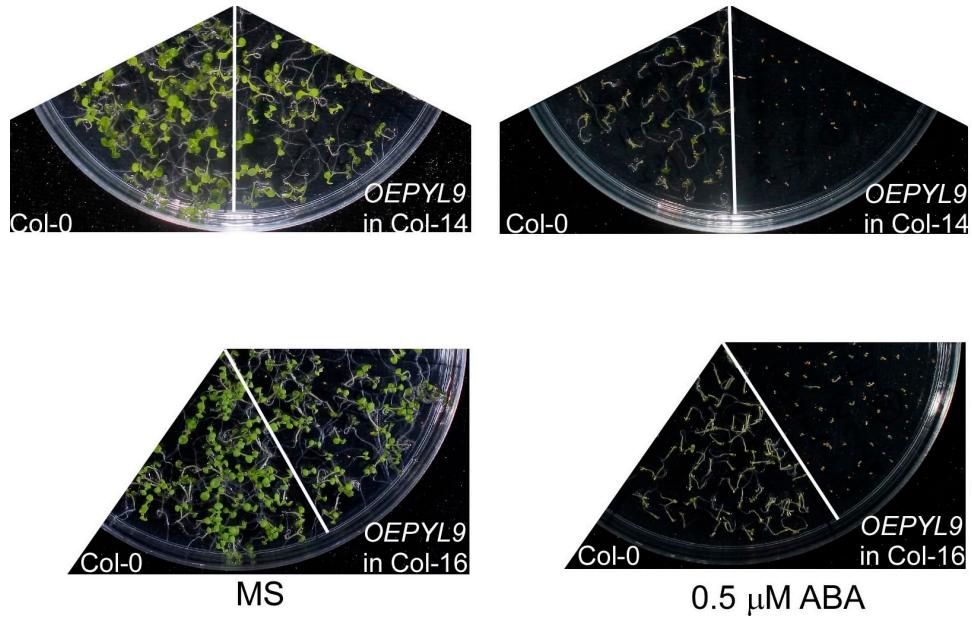
**Supplementary Figure S1.** Expression of *PUB22* and *PUB23* in the *pub22 pub23* double mutant. Quantitative real-time PCR determined the expression of *PUB22* and *PUB23* in *pub22 pub23* double mutant. The *pub22* (SALK\_072621) and *pub23* (SALK\_063470) single mutants[35]. Data are the means  $\pm$  standard errors ( $n = 3$ ).



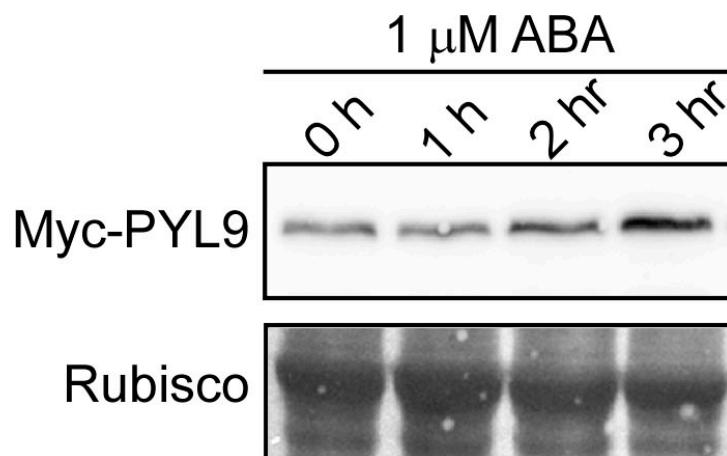
**Supplementary Figure S2.** The expression of *FLAG-PUB22* in transgenic plants. The expression of *FLAG-PUB22* in transgenic plants was analyzed by immunoblotting with anti-FLAG antibody.



**Supplementary Figure S3.** The seedling growth of the wild-type and the transgenic seedlings harboring *Myc-PYL9* on MS with or without mannitol. (a), (b) and (c). Seven-day-old seedlings grown on MS medium were transferred to the medium with different concentration of mannitol for 12 days. (d) Quantitative evaluation of the fresh weight of seedlings in (a), (b) and (c). Asterisks indicate significant differences ( $p < 0.05$ ).



**Supplementary Figure S4.** The seeds germination of the wild-type and the independent transgenic seeds with *Myc-PYL9* on MS with or without 0.5  $\mu$ M ABA. The seeds were vernalized at 4 °C for 3 days and grown on the chamber at 22 °C for 9 days.



**Supplementary Figure S5.** Accumulation of the Myc-PYL9 protein induced by ABA. The transgenic plants containing *Myc-PYL9* in Col-0 background were treated by 1  $\mu$ M ABA and the samples were harvested at the indicated time points. Coomassie brilliant blue (CBB) staining of Rubisco was used as loading control.

**Supplementary Table 1.** Primers used for plasmids construction.

Plasmids	Forward primer (5'-3')	Reverse primer (5'-3')
<i>pCAMBIA1307-Myc-PYL9</i>	ACGCGTCGACATGATGGACGGCGTTGAAGGC	GGGGTACCTCACTGAGTAATGTCCTGAG
<i>pCAMBIA1307-FLAG-PUB22</i>	ACGCGTCGACATGATGGATCAAGAGAGATAGAGATT	GGGGTACCTCAAGCAGGATAACGAATCATAC
<i>BD-PYR1</i>	CATGGAGGCCGAATTCATGCCTTCGGAGTTAACACC	GCAGGTCGACGGATCCTCACGTACCTGAGAACCAC
<i>BD-PYL1</i>	CATGGAGGCCGAATTCATGGCGAATTCAAGAGTCCTCC	GCAGGTCGACGGATCCTAACCTGAGAAGAGTTG
<i>BD-PYL2</i>	CATGGAGGCCGAATTCATGAGCTCATCCCCGGCCGTG	GCAGGTCGACGGATCCTTATTCATCATCATGCATAGGT
<i>BD-PYL3</i>	CATGGAGGCCGAATTCATGAATCTTGCTCCAATCCATG	GCAGGTCGACGGATCCTCAGGTCGGAGAACCGTG
<i>BD-PYL4</i>	CATGGAGGCCGAATTCATGCTTGCCGTTACCGTCC	GCAGGTCGACGGATCCTCACAGAGACATCTCTTCTTG
<i>BD-PYL5</i>	ACGAGGTCGGAATTCTCGCTGAGG	CGCGTCGACATAACTAATCATCAATTGCC
<i>BD-PYL6</i>	CATGGAGGCCGAATTCATGCCAACGTCGATACAGTTTC	GCAGGTCGACGGATCCTACGAGAATTAGAAGTGTCTC
<i>BD-PYL7</i>	CATGGAGGCCGAATTCATGGAGATGATCGGAGGAGAC	GCAGGTCGACGGATCCTCAAAGGTTGGTTCTGTATG
<i>BD-PYL8</i>	CATGGAGGCCGAATTCATGGAAGCTAACGGATTGAG	GCAGGTCGACGGATCCTAGACTCTGATTCTGCGTG
<i>BD-PYL9</i>	CATGGAGGCCGAATTCATGGACGGCGTTGAAGGC	GCAGGTCGACGGATCC TCACTGAGTAATGTCCTGAG
<i>BD-PYL10</i>	CATGGAGGCCGAATTCATGAACGGTGACGAAACAAAGAAG	GCAGGTCGACGGATCCTCATATCTTCTCCATAGATT
<i>BD-PYL13</i>	CATGGAGGCCGAATTCATGGAAAGTTCTAAGAAAAACG	GCAGGTCGACGGATCCTTACTTCATCATTCTTTGTGAGC
<i>AD-PUB18</i>	GGAGGCCAGTGAATTCAAGTCATAGCAGCATGATCCATACG	CGAGCTCGATGGATCCCCGAGCTAAATACAAACA

<i>AD-PUB22</i>	GGAGGCCAGTGAATTCATGGATCAAGAGATAGAGATT	CGAGCTCGATGGATCCTCAAGCAGGATAACGAATCATACT
<i>AD-PUB23</i>	GGAGGCCAGTGAATTCATGTCCGGAGGAATAATGGATG	CGAGCTCGATGGATCCTCAGCAGGGATATGCAAGAACATC
<i>pCold-GST-PYL5</i>	CGAGGGATCCGAATTCATGAGGTACCGGTGCAACTC	TAGACTGCAGGTCGACTTATTGCCGGTTGGTACTTCG
<i>pCold-GST-PYL7</i>	CGAGGGATCCGAATTCATGGAGATGATCGGAGGAGAC	TAGACTGCAGGTCGACTCAAAGGTTGGTTCTGTATG
<i>pCold-GST-PYL8</i>	CGAGGGATCCGAATTCATGGAAGCTAACGGGATTGAG	TAGACTGCAGGTCGACTTAGACTCTCGATTCTGTCGTG
<i>pCold-GST-PYL9</i>	CGAGGGATCCGAATTCATGATGGACGGCGTTGAAGGC	TAGACTGCAGGTCGACTCACTGAGTAATGTCCTGAG
<i>pCold-GST-PYL10</i>	CGAGGGATCCGAATTCATGAACGGTGACGAAACAAAGAAG	TAGACTGCAGGTCGACTCATATCTTCTCCATAGATT
<i>pCold-MBP-PUB22</i>	CGAGGGATCCGAATTCATGGATCAAGAGATAGAGATT	TAGACTGCAGGTCGACTCAAGCAGGATAACGAATCATACT
<i>pCold-MBP-PUB22C13A</i>	CTTCCTCTTCCCTGCTCCAATCTCTCTAG	CTAGAGAGATTGGAGCAAGGAAGAAGGAAG
<i>pCold-MBP-PUB23</i>	CGAGGGATCCGAATTCATGTCCGGAGGAATAATGGATG	TAGACTGCAGGTCGACTCAGCAGGGATATGCAAGAACATC
<i>pCold-MBP-PUB23C18A</i>	CCTCCGTTCTTCCCTGCTCCTATCTCTTGG	CCAAAGAGATAGGAGCAAGGAAGAACGGAGG
<i>pCAMBIA1300-PUB22C13A-NUC</i>	AGCTCGAGTAGTCGACATGGATCAAGAGATAGAG	ACGAGATCTGGTCGACAGCAGGATAACGAATCATACT
<i>pCAMBIA1300-CLUC-PYL9</i>	ACGCGTCCCAGGGCGGTACCATGATGGACGGCGTTGAAGGC	GCCCTCTAGAGGATCCTCACTGAGTAATGTCCTGAG
<i>pCAMBIA1300-CLUC-OsNAC2</i>	TCCCAGGGCGGTACCATGGACTTGCCCCCTGGC	GCTCTGCAGGTCGACTTAGTAGCCCCATAGCGC
<i>pSPYCE (M)-PUB22C13A</i>	CGCCACTAGTGGATCCATCAAGAGATAGAGATT	TACCCTCGAGGTCGACAGCAGGATAACGAATCATACT
<i>pSPYNE173-PYL9</i>	GCCTACTAGTGGATCCATGATGGACGGCGTTGAAGGC	TACCCTCGAGGTCGACTCACTGAGTAATGTCCTGAG

**Supplementary Table 2.** Primers used for the quantitative real-time PCR.

Plasmids	Forward primer (5'-3')	Reverse primer (5'-3')
<i>ACTIN</i>	ACTCTCCGCTATGTATGTC	GATGGAAGAGCTGGTCTTG
<i>PYL9</i>	AGCTCCTCTTCATCTCGTTGG	AGACTGCCGATTCAGGATCAC
<i>Myc-PYL9</i>	ATGGAGAGCTTGGCGACCTCAC	ATCACCATCGTTCTGCTCTTC
<i>PUB22</i>	TTCGGTTGGGAGGTTCTG	CAAACCCTAGCGTGAAGC
<i>PUB23</i>	GGCCGTGGAAGCTGGAGTAATC	CCCTAACCGCTCTATCGCTTGC
<i>PUB24</i>	TAGAGCCGAGATTCTTGC	ACCGTCCAACATTAACC
<i>NLUC</i>	TTCTATCCGCTGGAAGATGGAACC	TTCATAGCTCTGCCAACCGAACG
<i>CLUC</i>	TGGATGGCTACATTCTGGAGAC	GGTGTGGAGCAAGATGGATTG