

HD2C interacts with HDA6 and is involved in ABA and salt stress response in Arabidopsis

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Supplementary Material

Supplementary Tables:

Table S1. Primers used for RT-PCR analysis

Genes	Primers
<i>HD2A</i>	5' – AATTGAGCTCTTAGCTCTGGTTTGCAGCCTTTC – 3' and 5' – GCTCTAGACATGGGCTATTCTGAGGAAGAAGAG – 3'
<i>HD2B</i>	5' – CAGCTTCTACTCCTCAGAAG – 3' and 5' – AATTAGATCTCTACCCTTTCCCTTGCCC – 3'
<i>HD2C</i>	5' – TGACGCTGACGGTAGTGAAG – 3' and 5' – AATTAGATCTGCACTGTGTTTGGCCTTTG – 3'
<i>HD2D</i>	5' – TGATCTCTACTTAGGGCACG – 3' and 5' – AATTGAGCTCCTACTTTTTGCAAGAGGGAC – 3'
<i>ABI1</i>	5' – GCCATGTCGAGATCCATTGG – 3' and 5' – AACGATGCATCCCCAGCCAC – 3'
<i>ABI2</i>	5' – CAAGATCCATTGGCGATAGATACC – 3' and 5' – CCTCTTTTCTCCGCCGGAAG – 3'
<i>AtERF4</i>	5' – CGGCTACTACTAACCAGACCC – 3' and 5' – TCGCTGAAGGCACAATAA – 3'
<i>MYB2</i>	5' – AGCCGGGTTTCGTTCAATTC – 3' and 5' – CGAATACGATGTCGTATCGG – 3'
<i>MYC2</i>	5' – CTAAACCAAAGATTCTACGCG – 3' and 5' – GGTTCCTTGATTTGGAGTTTCTC – 3'
<i>UBQ</i>	5' – GATCTTTGCCGAAAACAATTGGAGGATGGT – 3' and 5' – CGACTTGTCATTAGAAAAGAAAGAGATAACAGG – 3'

Table S2. Primers used for Chip assay

Genes	Primers
<i>Actin2/7</i>	5' – CGTTTCGCTTTCCTTAGTGTTAGCT– 3' and 5' – AGCGAACGGATCTAGAGACTCACCTTG – 3'
<i>Ta3</i>	5' – GATTCCTACTGTAAAGAACATGGCATTGAGAGA – 3' and 5' – TCCAAATTCCTGAGGTGCTTGTAACC – 3'
<i>ABI1 P</i>	5' – GATATTTTACCGGTGGTC – 3' and 5' – GACGTGTCGTAGTCCGAGTT – 3'
<i>ABI1 E</i>	5' – CTTGTCTTCCTAGCTTCTTC – 3' and 5' – CCTTTACCCAATCTGATCCC – 3'
<i>ABI2 P</i>	5' – CTAGTGTGGTCAGTGTAGATG – 3' and 5' – GTGTAACATGCCATATGTCAC – 3'
<i>ABI2 E</i>	5' – CTCTCCTTTCTCTTCCCAAC – 3' and 5' – GAGGGTCAGTGAATGGTCTG – 3'
<i>AtERF4 P</i>	5' – GTGTACTCACTAGAGCAAGC – 3' and 5' – CCGTACAGTATTTGACGC – 3'
<i>AtERF4 E</i>	5' – ACCATTTTCAGGCTTTGAC – 3' and 5' – AACGCCTCTGTAACGAA– 3'

Legends of supplementary figures

Figure S1. Negative controls in BiFC assays

As negative controls, HD2C and HDA6 fused with YN or YC and the empty vector (YN and YC) were co-transfected into protoplasts and visualized using confocal microscope as negative controls. No YFP signals were detected.

Figure S2. Expression of *MYB2* and *MYC2* determined by real-time RT-PCR.

Total RNA was isolated from leaf tissues. Asterisks mark values that are significantly different from the wild type (t- test, **P <0.01, *P<0.05). The experiment was repeated three times with similar results.

Figure S3. Two additional independent analyses of histone acetylation and methylation of *ABI1* and *ABI2* in *hda6* plants. Relative levels of H3K9K14Ac (A) and H3K9Me2 (B) in *ABI1* and *ABI2* promoter and first exon regions were determined. The amounts of DNA after ChIP were quantified and normalized to an internal control *ACTIN2* for H3K9K14Ac or *Ta3* for H3K9Me2. Error bars represent standard errors. Asterisks mark values that are significantly different from the wild type (t- test, *P<0.05).

Supplementary Figures:

Figure S1

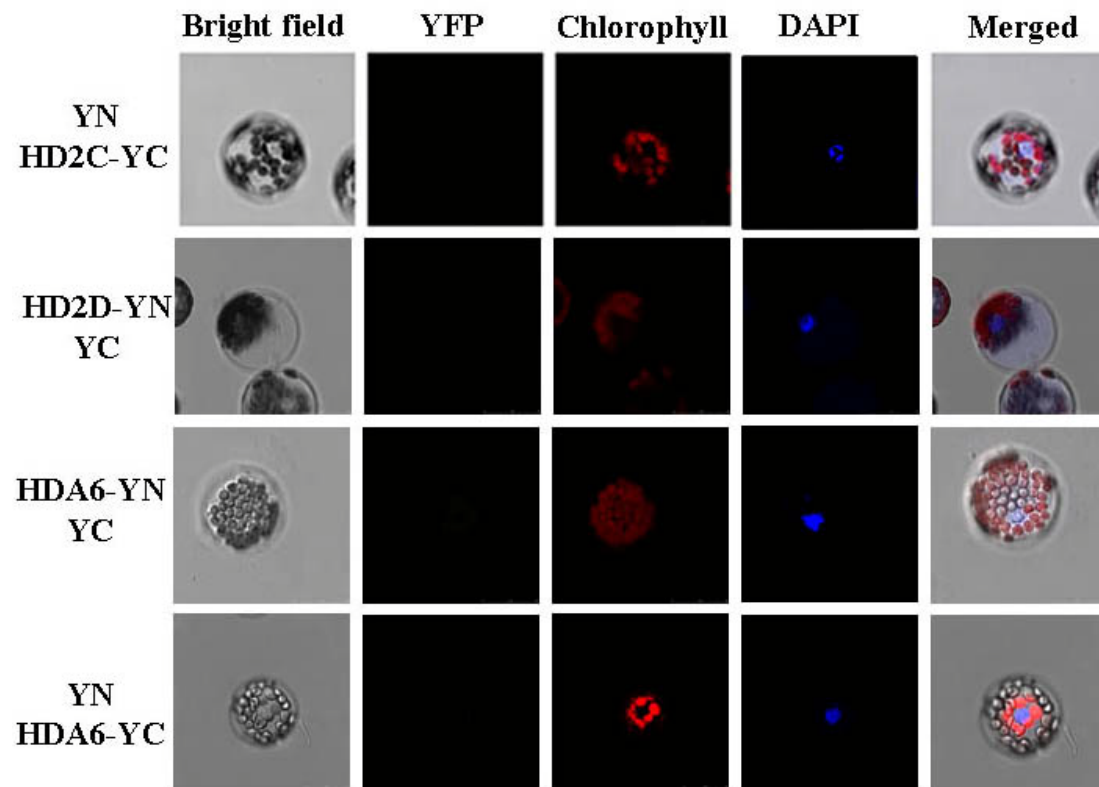


Figure S2

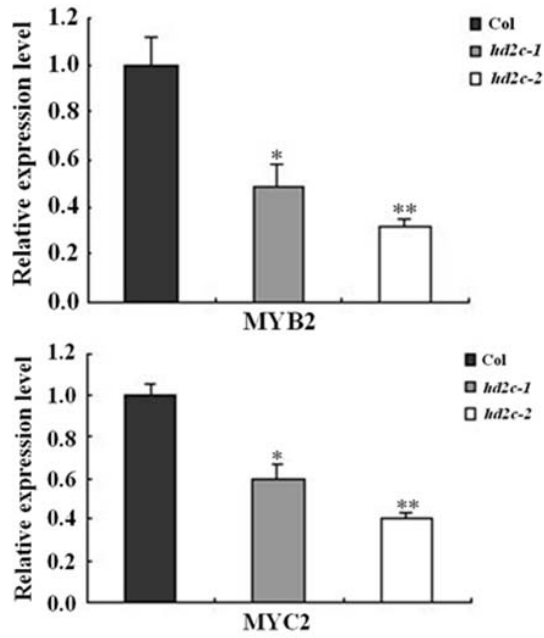


Figure S3

