

Table S1. PCR primers used in this study

| Primer | Sequence (5'-3') | Relevant characteristics |
|------------|---|--|
| P1 | CGAGATGCCAGGTACGTG | PCR primers to amplify <i>BdATG1</i> upstream fragment for construction of the <i>BdATG1</i> deletion fragment |
| P2 | CAAAATAGGCATTGATGTGTTGACCTCCGTGCGACGGGTGTGGAG | |
| P3 | CTCGTCCAGGGCAAAGGAATAGACTAGCAGTTCAACCATGGCTCGTC | PCR primers to amplify <i>BdATG1</i> downstream fragment for construction of the <i>BdATG1</i> deletion fragment |
| P4 | CGATCTTCCAGCACTCCGAG | |
| P5 | GCAGAGCGCTCGTATGTG | PCR primers to amplify the <i>BdATG1</i> deletion fragment for generation of the <i>BdATG1</i> deletion mutants |
| P6 | GCTGCAGGGTAACTAACTC | |
| P7 | CCATCAACCTCGTCGCTGC | |
| P8 | CCGTACCATGAAC TGATGC | PCR primers for identification of <i>BdATG1</i> deletion mutants |
| Probe-F | GACAGGCAGACGTGCAGAC | PCR primers to amplify the <i>BdATG1</i> upstream fragment used as the probe for Southern blotting analysis |
| Probe-R | GGCAGTGATGTGGGACATG | |
| ATG1-GFP-F | ACTCACTATA GGCGAATTGGTACTCAAATTGGTCAACCGGTGATCTGAAC | PCR primers to amplify the native promoter and open reading frame of <i>BdATG1</i> to construct <i>BdATG1</i> -GFP fusion vector |
| ATG1-GFP-R | CACCA CCCC GGTAACAGCTCCTCGCCCTTGCTCACCGAGGGCGATGCGTTGGC | |
| ATG1-ID-F | CGTATGCTTGAGGCAATC | PCR primers for PYF11- <i>BdATG1</i> -GFP vector identification |
| GFP-ID-R | GTGCTGCTTCATGTGGTC | |
| H1-GFP-F | ACTCACTATA GGCGAATTGGTACTCAAATTGGTCAACACACTCGATCAAGTC | PCR primers to amplify the native promoter and open reading frame of <i>BdH1</i> to construct <i>BdH1</i> -GFP fusion vector |
| H1-GFP-R | CACCA CCCC GGTAACAGCTCCTCGCCCTTGCTCACGCTTGGCCTCGGCCTC | |
| H1-ID-F | CTGCTGTACGTTAGACG | PCR primers for PYF11- <i>BdH1</i> -GFP vector identification |
| GFP-ID-R | GGCATGGCGGACTTGAAG | |

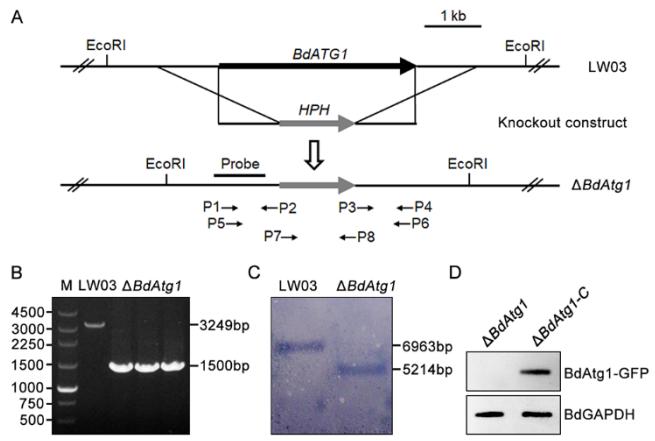


Figure S1: Target deletion of *BdATG1* in *B. dothidea*

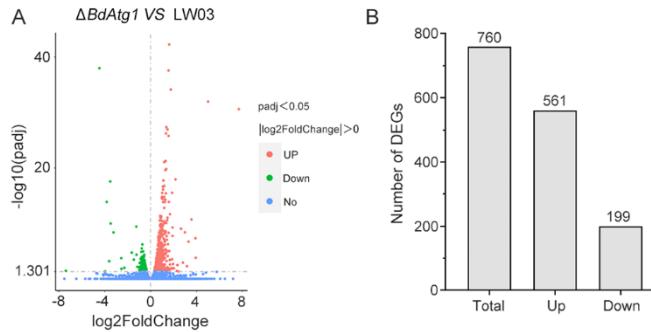


Figure S2: Differentially expressed genes (DEGs) of transcriptome sequencing