

Table S1. Bacterial strains and plasmids used in this study.

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| Strains or plasmids | Relevant characteristics | Resources |
|----------------------------------|---|------------|
| Strains | | |
| <i>Escherichia coli</i> | | |
| DH5 α | <i>F</i> <i>recA hsdR17</i> (<i>rk</i> ⁻ , <i>mk</i> ⁺) ϕ 80 <i>lacZ</i> Δ M15 | Clontech |
| Yeast | | |
| AH109 | MATa, <i>trp1-901</i> , <i>leu2-3, 112</i> , <i>ura3-52</i> , <i>His3-200</i> , <i>gal4</i> , <i>gal80</i> , <i>LYS2::GAL1UAS-GAL1TATA-His3</i> | Clontech |
| <i>Agrobacterium tumefaciens</i> | | |
| GV3101 | Rif ^r , with Ti plasmid pMP90 | [1] |
| Plasmids | | |
| pHB | Km ^r , a binary vector to express gene under control of a double CaMV 35S promoter | [2] |
| pHB-SDE1 | Km ^r , the 465-bp coding sequence of full length <i>SDE1</i> cloned in pHB | This study |
| pHB-SDE1mp | Km ^r , the 393-bp DNA fragment coding for SDE1mp lacking signal peptide cloned in pHB | This study |
| pGDR | Km ^r , RFP transient expression vector | [3] |
| pGDR-SDE1 | Km ^r , a 465-bp coding sequence of <i>SDE1</i> cloned into pGDR | This study |
| pGDR-SDE1mp | Km ^r , a 393-bp coding sequence of SDE1mp cloned into pGDR | This study |
| pGDG | Km ^r , GFP transient expression vector | [3] |
| pGDG-NbDDX3 | Km ^r , a 1413-bp <i>NbDDX3</i> gene cloned into pGDG | This study |
| pGBKT7 | Km ^r , GAL4(1-147) DNA-BD, TRP1, c-Myc epitope tag | Clontech |
| pGBKT7-SDE1 | A 465-bp coding sequence of <i>SDE1</i> cloned in pGBKT7 | This study |
| pGADT-7 | Amp ^r , GAL4(768-881) AD, <i>LEU2</i> , HA epitope tag | Clontech |
| pGADT7-CsDDX3 | Amp ^r , the 1413-bp <i>CsDDX3</i> genes cloned in pGADT7 | This study |
| TRV1 | Km ^r , a VIGS vector encoding the replication, movement and cysteine-rich protein proteoins of tobacco rattle virus | [4] |
| TRV2 | Km ^r , a VIGS vector harboring the coat protein and two non-structural proteins of tobacco rattle virus | [4] |

| | | |
|--------------|---|------------|
| TRV-NbPDS | Km ^r , a 369-bp DNA fragment of <i>NbPDS</i> cloned in TRV-RNA2 | [4] |
| TRV-NbDDX3 | Km ^r , a 437-bp DNA fragment of <i>NbDDX3</i> cloned in TRV-RNA2 | This study |
| 1301-YC | Km ^r , derived from pCAMBIA1301 which encodes the C-terminal portion of YFP | [5] |
| YC-SDE1mp | Km ^r , a 393-bp DNA fragment coding for SDE1mp cloned in 1301-Yc | This study |
| 1301-YN | Km ^r , derived from pCAMBIA1301 which encodes the N-terminal portion of YFP | [5] |
| YN-NbDDX3 | Km ^r , a 1413-bp <i>NbDDX3</i> gene cloned in 1301-YN | This study |
| SRC2-1-GFP | Gm ^r , a SRC2-1 GFP fusion in gateway vector pDONR207 | [6] |
| pGADT7-VemR | Amp ^r , the 384-bp full length of <i>vemR</i> genes cloned in pGADT7-AD vector at <i>NdeI</i> and <i>EcoRI</i> sites | [7] |
| pGBKT7-RpoN2 | Km ^r , the 1404-bp full length of <i>rpoN2</i> gene cloned in pGBKT7-BD vector at <i>NdeI</i> and <i>EcoRI</i> sites | [7] |

References

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Table S2. Primers for molecular cloning and qRT-PCR in this study

| Primer pair | Sequence(5'-3') | Cutting sites | Description or purpose |
|---|--|-----------------------------|---|
| Primers used for molecular cloning | | | |
| SDE1.pHB.F/SDE1.pHB.R | TTGGATCCATGCGTAAAAATTTATTAACCT TTGAGCTCTCAAGACTGCTCCAACATTT | <i>Bam</i> HI- <i>Sac</i> I | A 465-bp coding sequence of full length <i>SDE1</i> cloned into pHB |
| SDE1mp.pHB.F1 | TTGGATCCATGGGCAGTAGTTTTGGTTGTTG (Combined with SDE1.pHB.R) | <i>Bam</i> HI- <i>Sac</i> I | A 393-bp DNA fragment coding for SDE1mp cloned into pHB |
| SDE1.RFP.F/ SDE1.RFP.R | TCGTCGACATGCGTAAAAATTTATTAACCT TCGGATCCAGACTGCTCCAACATTTTC | <i>Sal</i> I- <i>Bam</i> HI | A 465-bp coding sequence of <i>SDE1</i> cloned into pGDR |
| SDE1mp.RFP.F | TCGTCGACATGGGCAGTAGTTTTGGTTGTTG (Combined with SDE1.RFP.R) | <i>Sal</i> I- <i>Bam</i> HI | A 393-bp coding sequence of <i>SDE1mp</i> cloned into pGDR |
| SDE1.BD.F/ SDE1.BD.R | TCGAATTCATGCGTAAAAATTTATTAACCT TCCTGCAGAGACTGCTCCAACATTTTC | <i>Eco</i> RI- <i>Pst</i> I | A 465-bp coding sequence of <i>SDE1</i> cloned into pGBKT7 |
| NbDDX3.Ri.F/ NbDDX3.Ri.R | GCTCTAGAGCCTGGCTGGGTAAAG GGGGTACCCAGAGCGATGAACAAAAGT | <i>Xba</i> I- <i>Kpn</i> I | A 437-bp DNA fragment of <i>NbDDX3</i> cloned in TRV-RNA2 for antisense |
| NbPSMD14.Ri.F/ NbPSMD14.Ri.R | GCTCTAGACCCTCCTACTCTGGACTCTT GGGGTACCTCTACCGTGGCTCTTGG | <i>Xba</i> I- <i>Kpn</i> I | A 467-bp DNA fragment of <i>NbPSMD14</i> cloned in TRV-RNA2 for antisense |
| Niben101Scf05290g02006.1.Ri.F / Niben101Scf05290g02006.1.Ri. R | GCTCTAGACAAAGCGGAGCAAACA GGGGTACCCACAAGATGAGGCAGAG | <i>Xba</i> I- <i>Kpn</i> I | A 438-bp DNA fragment of <i>Niben101Scf05290g02006.1</i> cloned in TRV-RNA2 for antisense |
| Niben101Scf04231g02014.1.Ri.F / Niben101Scf04231g02014.1.Ri. R | GCTCTAGATAGGGAAACGAGTTGTGAA GGGGTACCAGGGCGGAAGAAGGTC | <i>Xba</i> I- <i>Kpn</i> I | A 317-bp DNA fragment of <i>Niben101Scf04231g02014.1</i> cloned in TRV-RNA2 for antisense |
| NbDDX3.GFP.F/ NbDDX3.GFP.R | GCGTCGACATGCTCGCAGTTGGATTTG CGGGATCCATAGTCCCTTTTGTAGGGCA | <i>Sal</i> I- <i>Bam</i> HI | A 1413-bp <i>NbDDX3</i> gene cloned into pGDG |

| | | | |
|--------------------------|---|-------------------|---|
| SDE1mp.Yc.F/ SDE1mp.Yc.R | <u>GCTCTAGA</u> AATGGGCAGTAGTTTTGGTTGTTG GGGT <u>ACCAGACTGCTCCAACATTTTTC</u> | <i>XbaI-KpnI</i> | A 393-bp DNA fragment coding for SDE1mp cloned into 1301-YC |
| NbDDX3.Yn.F/ NbDDX3.Yn.R | <u>GCTCTAGA</u> AATGCTCGCAGTTGGATTTG GGGT <u>ACCATAGTCCCTTTTGTTAGGGCA</u> | <i>XbaI-KpnI</i> | A 1413-bp <i>NbDDX3</i> gene cloned into 1301-YN |
| CsDDX3.F/CsDDX3.R | <u>TCGAATTC</u> ATGCTTGCAGTTGGATTT <u>TCCTGCAGC</u> AGTTAGGACATTCAGAT | <i>EcoRI-PstI</i> | A 1488-bp <i>CsDDX3</i> gene cloned into pGADT7 |

Primers used for qRT-PCR analysis

| Gene | Sequence(5'-3') | Description |
|---------------|--|--------------------|
| <i>NbEF1a</i> | TGGTGCCTCAAGCCTGGTATGGTTG; ACGCTTGAGATCCTTAACCGCAACATTCTT | 160-bp |
| <i>SDE1</i> | CCATATGCGTCCTTTCACCAA; GAGATTGAGGAGCAGAACCCC | 125-bp |
| <i>NbDDX3</i> | GTTTTACACAGACAAAGCGAGATG; CCCTCTGATGTTGAGAAATGTCG | 103-bp |
| <i>CsCOX</i> | GTATGCCACGTCGCATTCCAGA; GCCAAAAGCTGCTAAGGGCATTG | 68-bp |
| <i>CsDDX3</i> | AAAACCTTCCACCCAAGCGACAGAG; CCCTCTGCAAGCTTTTCGTCTTGGT | 139-bp |

Underlined are restriction enzyme sites at 5' end of each primers

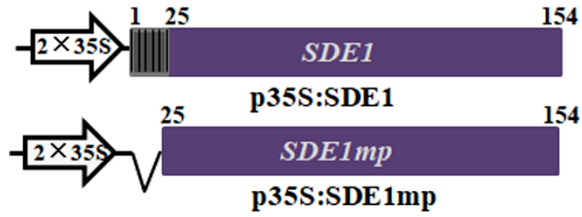


Figure S1 Schematic diagram of full-length SDE1 and mature SDE1 (SDE1mp) for transient expression. The N-terminal 24 amino acids were deleted to generate SDE1mp.

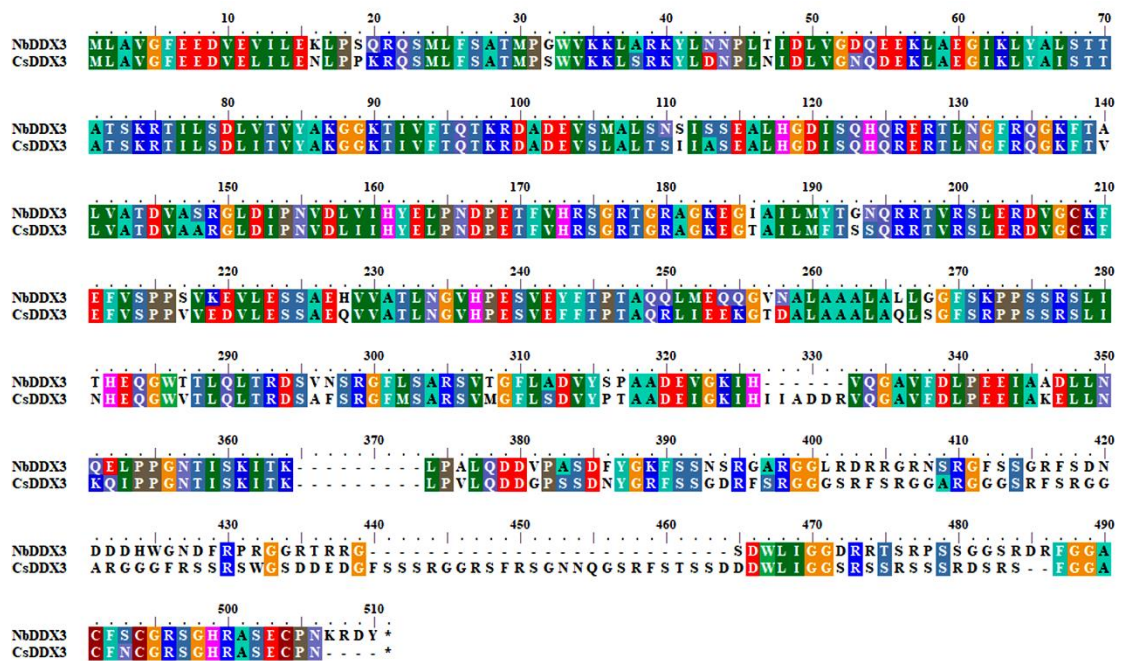
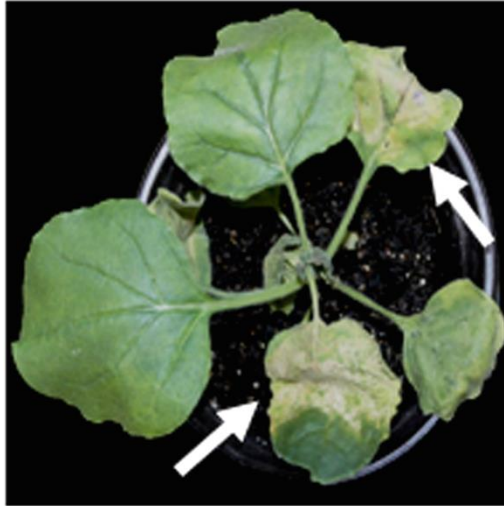


Figure S2 Alignment of amino acid sequences between NbDDX3 and CsDDX3.



TRV:NbPSMD14

Figure S3 Silencing of *NbPSMD14* gene in *N. benthamiana*. Growth phenotypes were scored at 15 dpi. White arrows indicate leaf death in silenced plants.