



Supplementary Material

The Single-Stranded DNA-Binding Gene *Whirly* (*Why1*) with a Strong Pathogen Induced Promoter from *Vitis pseudoreticulata* Enhances Resistance to *Phytophthora capsici*

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Supplementary Table

Table S1: Sequence information of *Why1* and promoter

Genes/ promoters	GenBank ID	Sequence length/bp	ORF length/bp	Number of exon/intron	Amino acid number	Molecular weight/kD	Theoretical pI	Hydro- pathicity	Subcellular localization
<i>VvWhy1</i>	MN395403	862	807	7/6	268	68.33	5.1	0.762	Chloroplast
<i>VpWhy1</i>	MN395402	862	807	7/6	268	68.28	5.1	0.755	Chloroplast
<i>VvWhy1 promoter</i>	MN397251	1127							
<i>VpWhy1 promoter</i>	MN397250	1136							

Table S2: Primer sequence

Primer name	Primer sequence	Vector or purpose
<i>Why1-F</i>	GGTTAAGAAGAATAGAGGCCGAAG	Gene cloning of <i>VvWhy1</i> and
<i>Why1-R</i>	CAGTCCATGCTTAGAAGTACCAAG	<i>VpWhy1</i>
<i>p-F</i>	TGAGTTATGACTTTATTGTTCTCCTT	Promoter cloning of <i>pVv</i> and
<i>p-R</i>	GATGCATTCTCCGCCTCTATTCTTCT	<i>pVp</i>
<i>Why1-gfp-F</i>	CGGGATCCATGCATCACCTGCACCTGCTTTC	pBI121- <i>Why1-GFP</i>
<i>Why1-gfp-R</i>	GGGTACCCCTCTGCTCCATTCAAAGTCCCCAG	
<i>p-GUS-F</i>	CCCAAGCTTGAGTTATGACTTTATTGTTCTC	pBI121- <i>pWhy1::GUS</i>
<i>p-GUS-R</i>	TCCCCCCGGTCTTCCGCCTCTATTCTTCTTAAC	
<i>p-Why1-gfp-F</i>	ATGCCCTGCAGGTGAGTTATGACTTTATTGTTCTC	pBI121- <i>pWhy1::Why1-GFP</i>
<i>p-Why1-gfp-R</i>	CGGGATCCTCTTCCGCCTCTATTCTTCTTAAC	
<i>Why1-q-F</i>	AGCAGAACATCACTCGCA	
<i>Why1-q-F</i>	TTGGGCTCTACCGTAAGA	
<i>NbPRI-q-F</i>	CCGTTGAGATGTGGGTCAAT	q-RT-PCR
<i>NbPRI-q-R</i>	CGCCAAACCACCTGAGTATAG	

<i>NbPR2-q-F</i>	CAACCCGCCAAAGATAGTA
<i>NbPR2-q-R</i>	TGGCTAAGAGTGGAAAGGTTATG
<i>NbPR4-q-F</i>	GGATGATGTTGACAGCAGAGA
<i>NbPR4-q-R</i>	GTAGGACACGAGGTAGGTATCA
<i>NbPR5-q-F</i>	GCTCGATTACGTCTTGTCTCTC
<i>NbPR5-q-R</i>	CTCTAGCATGGTGGATTGACTT
<i>NbPR10-q-F</i>	GAAGAAGAACACAATGAAGGCA
<i>NbPR10-q-R</i>	CAGTAGGATTGGCAAGAAGGTA
<i>V-EF1α-F</i>	GAACGGGTGCTTGATAGGC
<i>V-EF1α-R</i>	ACCAAAATATCCGGAGTAAAAGA
<i>Nb-EF1α-F</i>	AGAGGCCCTCAGACAAAC
<i>Nb-EF1α-R</i>	TAGGTCCAAGGTCACAA

Red letters indicate restriction enzyme sites, GGATCC:*Bam*HI; GGTACC: *Kpn*I; AAGCTT: *Hind*III; CCCGGG: *Sma*I; *Sbf*I:CCTGCAGG.

Why1-gfp-F and *Why1-gfp-R* were used to construct pBI121-*VvWhy1-GFP* and pBI121-*VpWhy1-GFP* for subcellular localization verification.

p-GUS-F and *p-GUS-R* were used to construct pBI121-*pVv::GUS* and pBI121-*pVp::GUS* for promoter GUS activity assay.

Based on the constructs of pBI121-*VvWhy1-GFP* and pBI121-*VpWhy1-GFP*, *p-Why1-gfp-F* and *p-Why1-gfp-R* were used to construct pBI121-*pVv::VvWhy1-GFP*, pBI121-*pVv::VpWhy1-GFP*, pBI121-*pVp::VvWhy1-GFP* and pBI121-*pVp::VpWhy1-GFP* for *Ph. Capsici* resistance experiments in *N. benthamiana*.

Supplementary Sequence

Sequence S1: *VvWhy1*

>*VvWhy1* ORF 26-832 bp

GGTTAAGAAGAATAGAGGCCGAAGAACATGCATCACCTGCACCTGCTTCTCCTCATTCAACCACATCCAAAACCC
 TAGATTATGCCCTAATCACTCCCTTCGCTCCACTCCTCCTCACCCCTGAGCTTCACTCACGAACCTCTCT
 TCTGCTTCCACAACCAGGCTTCCGTAAGCGCTATTGCAATGTCGCCAGTCGGATTACTTCAACAG
 CAGAACATCACTCGCAGACAACCTCAAATGACTCTTCTTGGAGGAGCTTGCAGCCTAGGGTTTTGTTG
 GTCATTGATATAAAAGGGAAAGCTGCTTACGGTAGAGCCCCAAAGCCCCGGAGTTACACCTTAGATT
 CAGGGGCATTAAAGTCCAAAGAAGGTTTGCTGCTCCAGTTGCTCCTGCTGCAGGTGTTGACAATA
 TGATTGGGGCAGAAAGCAGGTGTTCTCATTATCTGTGACTGAAATTGGAAGTCTTATTAGCCTGGTCAAGA
 GAGTCATGCGAATTTCATGATCCTTAAAGGAAGAAGTGGAGGAAGGTAAGGTCCCGAAGGTGTTGAAG
 GTAGAGCCGCTCCAGATGGCTCTGGTCATTCTCAATCTCAGTGTCAAAACAAGCTTGAATATGGATG
 AGAACATTATATCCCTGTCACCAGAGCAGAATTGCTGCTCATCTCAGCTTAAATTCAATTGCTGCCATAT
 CTTCAGGCTGGCATGCCTATGCGAACTCCATCAAGCCAGATGATAACAAGTCGTGTAATAATGCTAATCCGA
 GATCTGGGACTTGAATGGAGCAGATAGTAAACACTGGTACTTCTAAAGCATGGACTG

Sequence S2: *VpWhy1*

>*VpWhy1* ORF 26-832 bp

GGTTAAGAAGAATAGAGGCCGAAGAACATGCATCACCTGCACCTGCTTCTCCTCATTCAACCACATCCAAAACCC
 TAGATTATGCCCTAATCACTCCCTTCGCTCCACTCCTCCTCACCCCTGAGCTTCACTCACGAACCTCTCT
 TCTGCTTCCACAACCAGGCACTCCTAAAGCGCTATTGCAATGTCGCCAGTCGGATTACTTCAACAG

CAGAACATCACTCGCAGACAACCTCAAATGACTCTTCTTGAGGGAGCTTGCAGCCTAGGGTTTGTG
 GTCATTGATATAAAAGGGAAAGCTGCTTACGGTAGAGCCAAAGCCCCGGAGTTACGCCTTAGATT
 AGGGGCATTAAAGTGTCAAAGAAGGTTGTGCTGCCAGTTGCTCCTGCTGCAGGTGTTGACAATAT
 GATTGGGGCAGAAAGCAGGTGTTCTCATTATCTGTGACTGAAATTGGAAGTCTTATTAGCCTGGTGCAAGAG
 AGTCATGCGAATTTCATGATCCTTAAAGGAAGAAGTGAGGAAGGTAAGGTCCGGAAGGTGTTGAAGG
 TAGAGCCGCTCCCAGATGGCTCTGGTCATTCTCAATCTCAGTGTCAAAACAAGCTTTGAATATGGATGA
 GAACATTATATCCCTGTACCAGACAGAATTGCTGCTCATCTCAGCTTTAATTGATTGTGCCATATCT
 TTAGGCTGGCATGCCTATGCGAATCCATCAAGCCAGATGATAACAAGTCGTGAAATAATGCTAATCCGAG
 ATCTGGGACTTGAATGGAGCAGATAGTAAACACTGGTACTTCTAAAGCATGGACTG

Sequence S3: *VvWhy1* promoter

>*VvWhy1* promoter (pVv) 1127 bp

TGAGTTATGACTTTATTGTTCTCCTTATACACAATTATAGTTGAATTAGATAAAATAGATTATAAATTAA
 GTAAGAAATAATTAAATAATTAAATTCTTAAATAATTAAATAAAATTAAAGAATAATTAGAATAAAAAAAA
 CCAATTAAATTAAATTGTAATTAAATTAAAAACATTAAATTAAATTATATTGTAATTCAACTAAATGTGTAC
 ACGAAATAAAAAACTGACTTATTACGTATTAAAAAGTTAAAATTATTATTATTATTCAATTAAATAAAAT
 ATTCTAAAACCTATTCTAAATTGTTATTATTCAATTAAAGAAAAAAATTACGAAGATGTTAATATCCTAT
 GGTTTAATATATACTAAAATAATTTTTAATAAAAAGATATAATTGATTGTTATTATAATTATTATT
 AATGTTTATTAAAAACTATATATTTTTAATTATTGTAATTACAAAACATTGTTATTGTTAATAAGACTTG
 AATTAAATTAAATTAAATTATAATTGAATTAGTAATGTTTATAAAATAAAAAAAATCAATTAA
 AAACAGTTATCCAAGAAAATCAAATCATTATACAAACAAGTTGTTTATTGTTAATAAGACTTG
 AAAACAAATTAGCAAACACTCTTATATTATAAAATACAAGAAAACGTGTTATTCTTAATTAAAC
 AATTCTAAAACAAATTAAAGAAAATGACCAAATAGATCCTAATTGTTGAAATAATTCTAA
 AATTCTTATTCAATAATATATTGCCCACCTTAATTGTTATGAAAATATCTCATAGTGAAATT
 CGAAAATAACATATTCTAAAACAAATTCTCAAAACAAATTGTTAAGAAAATTGTTATTCTTAATTAA
 TCCTAAGATGCCAACAGACTTTATTCTCACGAAATTACTTGTAAATAGTAATAAGAAACGGCAGCGT
 GCCTCAGGGCATCGTCAGCAGCCTCATGCTCAAAGTAAAAGCTCATCTCCGAAGCGAAATTGGGGTT
 AAGAAGAATAGAGGCCGAAGAATGCATC

Sequence S4: *VpWhy1* promoter

>*VpWhy1* promoter (pVp) 1136 bp

TGAGTTATGACTTTATTGTTCTCCTTATACACAATTATAGTTGAATTAGATAAAATAGATTATAAATTAA
 GTAAGAAATAATTAAATAATTAAATTCTTAAATAATTAAATAAAATTAAAGAATAATTAGAATTCAAAAAA
 AACCAATTAAATTGTAATTAAATTAAAAACATTAAATTAAATTATATTGTAATTCAACTAAATGTGT
 ACACGATATAAAAAACTGACTTATTACATATTAAAAAGTTAAAATTATTATTATTCAATTAAATAAAC
 TATTCTAAAATTATTCTAAATTGTTATTATTCAATTAAAGAAAAAAATTGAAGATGTTAATATCCTTA
 TGGTTTAATATATAATAAAATAATTGTTAATAAAAAAAATATAATTGATTGTTATTATAATTATT
 TTAATGTTTATTAAAAACTATATATTGTTATTGTAATTAAAAAAACTATCTTATTGTTAATAAGA
 CTTGAATTAAATTAAATTATATTAAATTGAATTAGTAATGGTTATAAAATCAAAAAAAGAAAAAAATC
 ATTGTTAGAAACAGTTATTCAAGAAAATCAAATCATTATACAAACAAGTTGTTATTGTTATTGAAATT
 TTTTAAACAAATTCTCAAACACTCTTATATTATAAAACACAAGAAAACGTGTTATTGTTATT
 AAAACAAATTGTTAAAAACAAATTAAAGATATGACCAAATAGACCCTAATTGTTGAAAATAATTCTT

AAAAATAATTTTATTTCAATAATATTTTTGTCCCACTTAATTTATGAAAATATCTCATATAGTGT
AAATTCAAAAAATAACATATTCTAAAACGAATTTCAAAAAAATTGTATTAAAAGTTAAAGGGGCTT
GAGATTCCTAAGATGCCAGCAGACTTTATTCTCACGAAATTAACCTTGTAATTAGTAATAAGAAACGG
CAGCGTGCCTCAGGGCATCGTCAGCAGCCTCATTGCTCAAAGTTAAAAGCTCTATCTTCCGAAGCGAAATTG
GGGGTTAAGAAGAATAGAGGCGGAAGAATGCATC