

Table S2 Information of primers used in this study

Primer name	Primer 5'-3'	Purpose
BeXyl1 F	CGGGATCGATATTTC C AATGTAAATC C	To amplify the DNA fragment of BeXyl1 for expression in <i>Pichia pastoris</i>
BeXyl1 R	CGGAATTC GCATAAAAGT AATTATCCGA GGTAGT	To amplify the DNA fragment of BeXyl1 for expression in <i>Pichia pastoris</i>
BeXyl1rec F	TATTGTGCGCGGTGGAGGAGCCCAGTCTGCTTCTGCAAAGGTTATCCAGCA	To amplify the DNA fragment of site-directed mutagenized BeXyl1rec for expression in <i>Pichia pastoris</i>
BeXyl1rec R	CGCGCACAATAGCGTTACCATCCCCCT TAATACTCAG CGCTGCCTCCAGACA	To amplify the DNA fragment of site-directed mutagenized BeXyl1rec for expression in <i>Pichia pastoris</i>
BeXyl1 21-329-F	CGGAATCCGTCAACAACGTCACCCCCAACA	To transient expression BeXyl1 21-329 protein (deleted the N-terminal signal peptide)
BeXyl1 21-329-R	GCTCTAGA GCATAAAAGT AATTATCCGA GGTAGT	To transient expression BeXyl1 21-329 protein (deleted the N-terminal signal peptide)
BeXyl1-T-F	CGGAATCCATGCATTCCTTCTTTATGCTAA CA	To transient expression BeXyl1 protein (with the N-terminal signal peptide)
BeXyl1-T-R	GCTCTAGA GCATAAAAGT AATTATCCGA GGTAGT	To transient expression BeXyl1 protein (with the N-terminal signal peptide)
RT-qPCR-NbEF1a-F	AGGATACAACCTGACAAGA	<i>N. benthamiana</i> EF-1a gene used as qRT-PCR reference
RT-qPCR-NbEF1a-R	AGGATACAACCTGACAAGA	<i>N. benthamiana</i> EF-1a gene used as qRT-PCR reference
RT-qPCR-Beggsdb-F	CGAGTACCAGGCTGGTATCT	<i>B. cinerea</i> Beggsdb gene used for qPCR measurement of pathogen levels and as qRT-PCR reference
RT-qPCR-Beggsdb-R	TCATTGGTGGTTAGCCATCTT	<i>B. cinerea</i> Beggsdb gene used for qPCR measurement of pathogen levels and as qRT-PCR reference
RT-qPCR-PR1a-F	GTGGGTCGATGAGAAACAGTAT	RT-qPCR for validation of <i>PR1a</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-PR1a-R	GAACCTAGCACATCCAACA	RT-qPCR for validation of <i>PR1a</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-NPR1-F	GGAGCAAGCAGAAAGAAGAGA	RT-qPCR for validation of <i>NPR1</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-NPR1-R	GTTTAGCCAGGCCAACTCTAT	RT-qPCR for validation of <i>NPR1</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-PR5-F	GGGCAATCTTGGAGCATT	RT-qPCR for validation of <i>PR5</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-PR5-R	CAGTCTCCAGTCTCACAAATTACC	RT-qPCR for validation of <i>PR5</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-COI1-F	GGCTTGACGTACTTAGGGAAATA	RT-qPCR for validation of <i>COI1</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-COI1-R	GGGACACCTTTGCAGTAAGA	RT-qPCR for validation of <i>COI1</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-PAL-F	ATTGCTGGTTTGCTCACTGG	RT-qPCR for validation of <i>PAL</i> gene transcription level in <i>N. benthamiana</i> leaves
RT-qPCR-PAL-R	TCCTTAGGCTGCAACTCGAA	RT-qPCR for validation of <i>PAL</i> gene transcription level in <i>N. benthamiana</i> leaves
NbBAK1-F	GGAATCCGTGAGGGTGGTGAGCGGGATAAT	Silencing of <i>BAK1</i> in <i>N. benthamiana</i>
NbBAK1-R	CCTCAGAGCTCATAACTGGGCAAAGGGCTT	Silencing of <i>BAK1</i> in <i>N. benthamiana</i>
NbSOBIR1-F	TGCTCTAGAGGAAGAAAAGGAAAGTCAGA	Silencing of <i>NbSOBIR1</i> in <i>N. benthamiana</i>
NbSOBIR1-R	CCGCTCGAGATCCATTGGGGGTTGTATAA	Silencing of <i>NbSOBIR1</i> in <i>N. benthamiana</i>