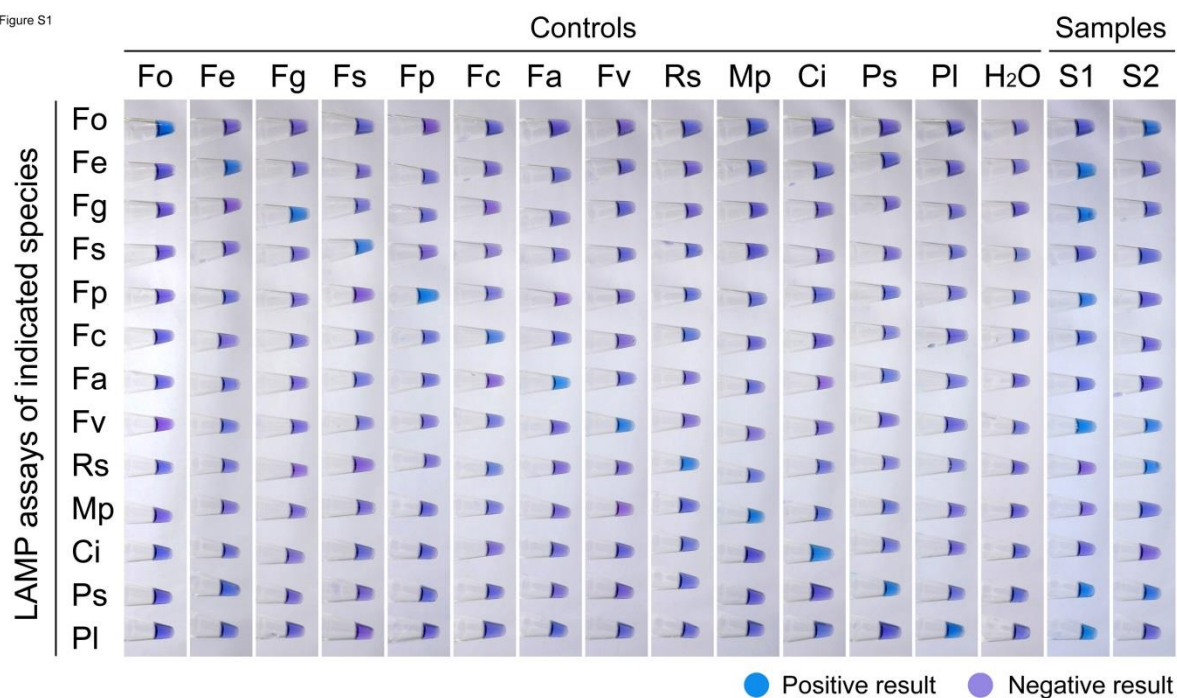


SUPPLEMENTARY INFORMATION

Figure S1



Appendix A. Results of loop-mediated isothermal amplification (LAMP) assays.

DNA was analysed using 13 LAMP assays. Results shown in the 14 columns on the left (controls) were obtained using either DNA of the indicated species or sterile ultra-pure water as templates. The 15th and 16th columns represent assay results for samples S7-07 (S1) and J7-13 (S2). Sky-blue colour indicates a positive result; purple indicates a negative result.

Appendix B. Information of the 13 LAMP assays used in this study

Target species (abbreviation)	Target gene ^a	Sensitivity	Publication
<i>Phytophthora sojae</i> (Ps)	<i>A3apro</i>	10 pg· μ l ⁻¹	(Dai et al., 2012)
<i>Fusarium oxysporum</i> (Fo)	<i>Cyp51c</i>	100 pg· μ l ⁻¹	(Zeng et al., 2016)
<i>F. equiseti</i> (Fe)	<i>Cyp51c</i>	100 pg· μ l ⁻¹	(Zeng et al., 2016)
<i>F. graminearum</i> (Fg)	<i>Cyp51c</i>	100 pg· μ l ⁻¹	(Lu et al., 2015b)
<i>F. virguliforme</i> (Fs)	<i>Tef-1a</i>	100 pg· μ l ⁻¹	(Lu, 2015)
<i>F. proliferatum</i> (Fp)	<i>Red1</i>	400 pg· μ l ⁻¹	(Rong et al., 2018)
<i>F. culmorum</i> (Fc)	<i>Cyp51c</i>	100 pg· μ l ⁻¹	(Zeng et al., 2017)
<i>F. sambucinum</i> (Fa)	<i>Tef-1a</i>	100 pg· μ l ⁻¹	(Xu et al., 2018)
<i>F. verticillioides</i> (Fv)	<i>Pgk</i>	100 pg· μ l ⁻¹	(Zeng et al., 2018)
<i>Rhizoctonia solani</i> (Rs)	<i>ITS</i>	10 pg· μ l ⁻¹	(Lu et al., 2015a)
<i>Macrophomina phaseolina</i> (Mp)	<i>ITS</i>	100 pg· μ l ⁻¹	(Lu et al., 2015a)
<i>Calonectria ilicicola</i> (Ci)	<i>β-tublin</i>	10 pg· μ l ⁻¹	(Lu, 2015)
<i>Phomopsis longicolla</i> (Pl)	<i>Tef-1a</i>	100 pg· μ l ⁻¹	(Dai et al., 2016)

^a*A3aPro* is a 300-bp transposon-like element initially identified in the promoter region of the *P. sojae* avirulence gene *Avr3a*; *Red1* encodes a reductase; *Pgk* encodes a 3-phosphoglycerate kinase; *Rpb1* encodes the large subunit of RNA polymerase II.

Reference

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Appendix C. Primer sequences of the 13 LAMP assays.

Species	Primer name	Primer sequence
Ps	F3	GCGTATTGAGGGTTGCTG
	B3	GCGTCCTATCACCTAGTGC
	FIP	ACGTGGGTTCCGGATTGGACCCTTGGGTACTGTGTACCAG
	BIP	CGCCACCGATGATTTCGACGAAATCAACCATCACTCACCG
	LF	-
	LB	GTAGGATGATTGGATGAACAC
Fo	F3	CTCGAAGGGTAATGGGGAGA
	B3	ATTGGGATCAGGCTGTTCAA
	FIP	CGCCAAAGGGTAGATATGGGCTAGTTGATTATGGCTTCGGCG
	BIP	GGACGGCATCGCTGTGTTGGTGAAGCAGTCTCACAAACGT
	LF	-
	LB	AGAGAACTACGCGTATGCC
Fe	F3	ACGATGTTCCCTAGCTTCTC
	B3	GCGGACTTCATTTCTAGA
	FIP	CTTCGACTTCTTTGACGAATTTGGAAGCTTGGTCTAACTACTGAA
	BIP	CGCACCTCACCTACTTCAACGGTGTAGATGGTGATTTCAG
	LF	-
	LB	GGGCAACACAGGAATCGTCAAC
Fg	F3	GTCCAATCCACTCCATCCTC
	B3	CGGTCTTCTCGAGAGGTTCA
	FIP	GTGTGTGGGATGGTGGCACTACAAGTCAAATCACCCATGCG
	BIP	CTTCGTCTCCCGGCACAATGGTCTTGTCCCAACGATGAGGA
	LF	-
	LB	CCCGCTCAGAAGAATTCTTCC
Fs	F3	CCTGCACACAAAAACACCA
	B3	GGGGAGTCTCGAACTTCCA
	FIP	CGAGCTCAGCGGCTTCTATTGGCGCATCATCACGTGGTT
	BIP	GTTCCCTCAAGTACGCCTGGGTAGGGCAATGTCGATGGT
	LF	CGAGCTCAGCGGCTTCTATTG
	LB	GCGCATCATCACGTGGTT
Fp	F3	ATCGGCGTTTTCAACTCTCT
	B3	TTGGATGCTGACGACATGAA
	FIP	CTATGCCAGAGCCGACGTAGACCCAAGCATTCTCCCCCT
	BIP	GACCCCAATGGCTGAGGTACCTCGGAAGTAGTCAACCATCT
	LF	CTTTGCAGCTGGGGCTG
	LB	GTTTAGTTACGCCGCCGCCAAA
Fc	F3	CTACATCCCAAAGTTCATCAA
	B3	ATGGGCTGGAACCATTG
	FIP	GGCCATGACTTCTGTAATATTGACGAGTTGAAAACACTACGTCAAACATC
	BIP	AATCACAATTTACACAGCATCTGGATAGAGAGTGGCGAATGTG
	LF	TGTCGCCCTTGAAGTATGGA
	LB	TCCCTCTAGGAAACGAAGTCC
Fa	F3	CTACAATTGGCGGGGTAGC
	B3	TGAACTCGAGCGGGGTAG
	FIP	TCGATACCACCGCACTGGTAGAAAGATACGCTTGTGCTGACA
	BIP	TTCATTTTCTCGATCGCGCGGTGTCGAGTCGTATCAGAGC
	LF	TCAAGTGACCGGTCTATGATGTA
	LB	-
Fv	F3	CCCACGCTACATGATGCTAA
	B3	GATCTTGTCCGAGACCTTGG
	FIP	GAAACCGGAGGCCCTTCTGGGCTTTCGGCACTGCTCACC
	BIP	GAAGGAGCTCGAGTACTTCGCCAGGATGGCGAGGAAAGGA
	LF	AGGTCGACACCGACCATG
	LB	AAGGCTCTCGAGGAGCCCAAG
Rs	F3	CGAAATGCGATAAGTAATGTGAA

	B3	AGAGGAGCAGGTGTGAAG
	FIP	GCTCCAAGGAATACCAAGGAGCCAGAATTCAGTGAATCATCGAATC
	BIP	TGCCTGTTTGAGTATCATGAATTCTAAAAGACCTCCAATACCAAAG
	LF	-
	LB	-
Mp	F3	GCACATTGCGCCCCTTG
	B3	G TTCAGAAGGTTTCGTCCGG
	FIP	AGGACGGTGCCCAATACCAAGCGGGGCATGCCTGTTCGA
	BIP	CTCAAAGACCTCGGCGGTGGGCTCCGAAGCGAGGTGTA
	LF	-
	LB	-
Ci	F3	CGACAGCAACAAAGCTCGA
	B3	AGATGGTCTGCCAGAAAGC
	FIP	CGGTCTGGAGGTGGACCTAGAAGACCCCAAGCACGATGTG
	BIP	TCCAGCTTCCAAAAACTGCCCTAGCACCAATTTGGTTACCCT
	LF	AGTCAGCAACCTTGTCTCCTCCGATAT
	LB	GGGGATTCACTAACATTCGCG
PI	F3	CCACCACACGTGCAGCT
	B3	CGACAGGGTTGATGTGTGAG
	FIP	AGCGTCGCAGACAGCGCATGCGCTCCATATCCCTCCA
	BIP	AGGCATTTTCACCCCTCCCTCTTGATAAGCGCACCCCGC
	LF	GCCAACGGTTTCTATGTGCA
	LB	-