Supplemental Material

Table S1 Strains and plasmids used

Strain or plasmid	Relevant characteristic(s)	Reference
Strains		
Pst DC3000	*Pseudomonas syringae* pv. *tomato* DC3000, wild type	(1)
Pst DC3000 Δ*algU*	Δ*algU*	(2)
Pst DC3000 Δ*algUmucAB*	Δ*algUmucAB*	(2)
Pst DC3000 D36E	*Pst* DC3000 with all 36 known virulence effector genes deleted	(3)
Pst DC3000 D36E Δ*fliC*	Effector polymutant with *fliC* deleted	(4)
Pst DC3000 D36E Δ*algUmucAB*	Effector polymutant with *algUmucAB* deleted	This work
Pst DC3000 D36E Δ*algUmucAB* Δ*fliC*	Effector polymutant with *algUmucAB and fliC* deleted	This work
Agrobacterium tumefaciens GV2260	VIGS vector delivery strain	(5)
Pst DC3000 Δ*hopQ1-1*	*hopQ1-1* effector gene deletion renders *Pst* DC3000 pathogenic on *N. benthamiana*	(6)
Pst DC3000 Δ*hopQ1-1* Δ*fliC*	Δ*hopQ1-1* Δ*fliC*	(6)
Pst DC3000 Δ*hopQ1-1* Δ*algUmucAB*	Δ*hopQ1-1* Δ*algUmucAB*	This work

Pst DC3000 Δ*hopQ1-1* Δ*algUmucAB* Δ*fliC*	Δ*hopQ1-1* Δ*algUmucAB* Δ*fliC*	This work
Pst DC3000 Δ*hrcQ-U*	Non-pathogenic, Type three secretion system deficient *Pst* DC3000	(7)
Pst DC3000 Δ*hrcQ-U* Δ*fliC*	Δ*hrcQ-U* Δ*fliC*	(8)
Pst DC3000 Δ*hrcQ-U* Δ*algUmucAB*	Δ*hrcQ-U* Δ*algUmucAB*	This work
Pst DC3000 Δ*hrcQ-U* Δ*algUmucAB* Δ*fliC*	Δ*hrcQ-U* Δ*algUmucAB* Δ*fliC*	This work
Pst DC3000 Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB*	Δ*avrPto* Δ*avrPtoB* mutants are unable to suppress flagellin activated PTI	(6)
Pst DC3000 Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB* Δ*fliC*	Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB* Δ*fliC*	(6)
Pst DC3000 Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB* Δ*algUmucAB*	Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB* Δ*algUmucAB*	This work
Pst DC3000 Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB* Δ*algUmucAB* Δ*fliC*	Δ*hopQ1-1* Δ*avrPto* Δ*avrPtoB* Δ*algUmucAB* Δ*fliC*	This work
Plasmids		
pJN105	P_{BAD}::empty; Gm^r	(9)
pEM53	P_{BAD}::*algU*; Gm^r	(9)
pTRV1	VIGS destination vector	(5)
pQ11-EC1	VIGS control vector	(5)

pTRV2-FLS2	FLS2 silencing construct	(5)
pZB28	pK18mobsacB/Δ*algUmucAB*	(2)
pME6010-fliC	*fliC* expressed from its native promoter; Tc^r	(10)

Table S2. Oligonucleotide primers used in qRT-PCR

No.	Sequence	Description
381	CCGCAAGGTGATTATCTCAGC	*gap-1* Fwd
382	TGGAGATGATCTGGTGCGACT	*gap-1* Rev
6868	CCAGACAATGGCGATCAA	PSPTO_1949 (*fliC*) Fwd
6869	TACGCTGCAGAATGTTGG	PSPTO_1949 (*fliC*) Rev
6980	GAACATGCCGAACATGGA	PSPTO_915 Fwd
6981	CGATTCCTGAGACTCGGT	PSPTO_915 Rev
6982	AAATCACGGACAAGCTGAC	PSPTO_1925 Fwd
6983	ACACGGTTGCTGTCTACTA	PSPTO_1925 Rev
6984	AACAGCCGCCATATCGAA	PSPTO_1933 Fwd
6985	AGTCAACGGTGTTCTGGT	PSPTO_1933 Rev
6986	CGCTAAAGATCACGAGGTTG	PSPTO_1979 Fwd
6987	AGGTCGTCGAAACTGAAGA	PSPTO_1979 Rev
6988	CTTCCTGTACTGATGGTTACTG	PSPTO_1980 Fwd
6989	CCGTGAATGGCTTGACTAC	PSPTO_1980 Rev
6990	TATCGAAACGCCGGAAGA	PSPTO_1982 Fwd
6991	GGAAATCGACAGCAGGATATG	PSPTO_1982 Rev
6992	ATCCGAACGAGATCAAATGG	PSPTO_1987 Fwd
6993	CTCTGCAACATCCAGCAA	PSPTO_1987 Rev

6994	GGTTTACTTGCGTCAGTCC	PSPTO_1988 Fwd
6995	CAGTTCGCCGTTCTTGTT	PSPTO_1988 Rev

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