

Table S1. BLAST results of *Xanthomonas* sp. strains on 16SrDNA sequence.

Strain	Bacterial strains in GenBank	Identity (%)	Accession
XL1	<i>Xanthomonas perforans</i> 91-118	99.93	CP019725.1
	<i>Xanthomonas perforans</i> LH3	99.93	CP018475.1
	<i>Xanthomonas perforans</i> GEV1001	99.93	CP116309.1
	<i>Xanthomonas perforans</i> GEV872	99.93	CP116305.1
	<i>Xanthomonas perforans</i> TC2-1	99.93	CP077785.1
XL2	<i>Xanthomonas perforans</i> 91-118	99.93	CP019725.1
	<i>Xanthomonas perforans</i> LH3	99.93	CP018475.1
	<i>Xanthomonas perforans</i> GEV1001	99.93	CP116309.1
	<i>Xanthomonas perforans</i> GEV872	99.93	CP116305.1
	<i>Xanthomonas perforans</i> TC2-1	99.93	CP077785.1
XL3	<i>Xanthomonas perforans</i> 91-118	99.87	CP019725.1
	<i>Xanthomonas perforans</i> LH3	99.87	CP018475.1
	<i>Xanthomonas perforans</i> GEV1001	99.87	CP116309.1
	<i>Xanthomonas perforans</i> GEV872	99.87	CP116305.1
	<i>Xanthomonas perforans</i> TC2-1	99.87	CP077785.1
XL4	<i>Xanthomonas perforans</i> 91-118	99.16	CP019725.1
	<i>Xanthomonas perforans</i> LH3	99.16	CP018475.1
	<i>Xanthomonas perforans</i> GEV1001	99.16	CP116309.1
	<i>Xanthomonas perforans</i> GEV872	99.16	CP116305.1
	<i>Xanthomonas perforans</i> TC2-1	99.16	CP077785.1
G1	<i>Xanthomonas perforans</i> 91-118	99.87	CP019725.1
	<i>Xanthomonas perforans</i> LH3	99.87	CP018475.1
	<i>Xanthomonas perforans</i> GEV1001	99.87	CP116309.1
	<i>Xanthomonas perforans</i> GEV872	99.87	CP116305.1
	<i>Xanthomonas perforans</i> TC2-1	99.87	CP077785.1