

Development of a real-time PCR method for the identification of cucurbit seed-borne *Pseudomonas syringae* species

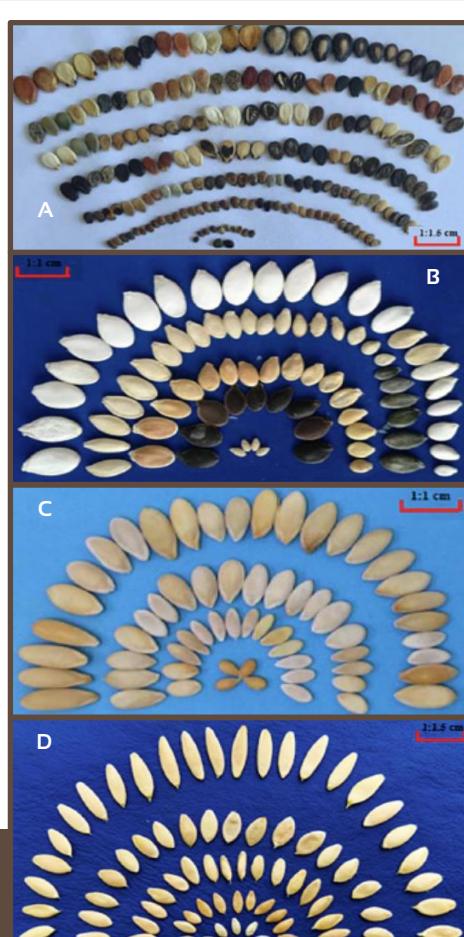
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Pseudomonas syringae-induced diseases in cucurbits

Cucurbits:

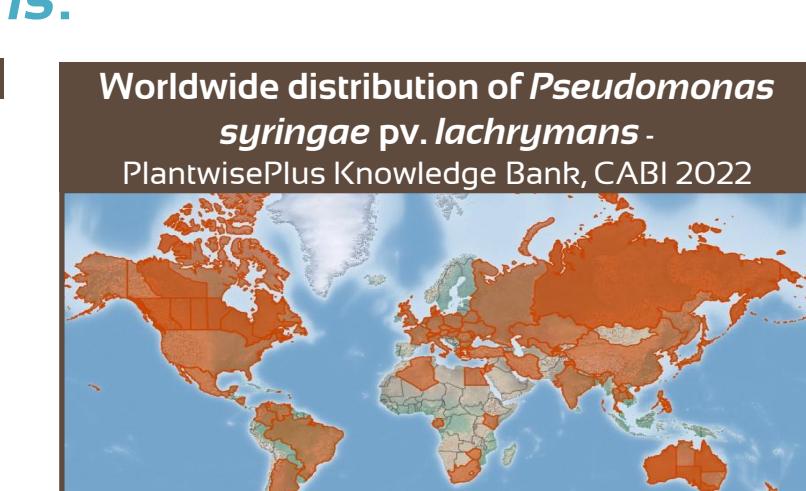
- Among the most cultivated crops
- Most economically important species: watermelon, melon, cucumber...
- Remarkable diversity in seeds and in fruit size, shape and colour



Examples of seed-borne diseases:

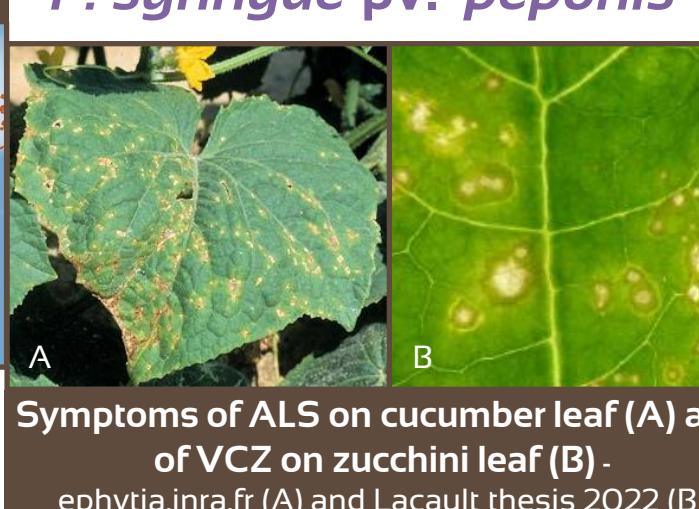
Angular leaf spot (ALS) caused by *P. syringae* pv. *lachrymans*:

- Most widespread bacterial disease of all cucurbit crops
- Significant yield reduction in the number of fruits and fruit weight
- Symptoms: Angular spots with yellow halo on leaves and water-soaked tan, small circular spots on fruits



Vein Clearing on Zucchini (VCZ) caused by lineages of *P. syringae*:

- First observation: 2004
- Causal agent was provisionally named *P. syringae* pv. *peponis*'

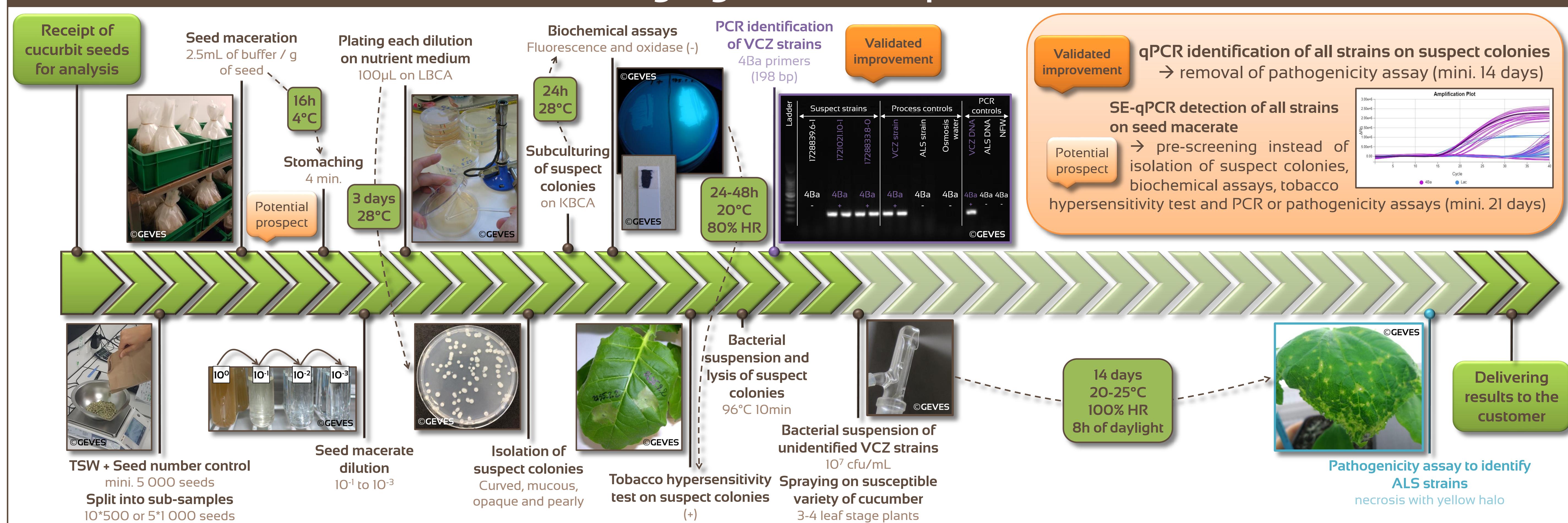


- Symptoms: cotyledon and leaf necroses, vein clearing, and stunting

Diseases management:

- Resistant crop cultivars
 - Cultural practices
 - Chemical and biological control
 - Seed health analysis
- Bacteriological analysis of cucurbit seeds at GEVES (detection and identification of ALS and VCZ strains)
- Around 450 analyses per year (approx. 20%)
 - Analysis performed on 5 000 seeds and in 36 days

Cucurbit seeds detection of *Pseudomonas syringae* strains responsible of diseases



Development of a qPCR identification method for *Pseudomonas syringae*-induced diseases in cucurbits

1. Characterisation of the strain collection:

- Host and morphological criteria
- Biochemical and tobacco hypersensitivity tests
- Identification PCR
 - PCR 4Ba for VCZ strains (Manceau et al., 2011)
 - other PCRs for non-target strains
- Pathogenicity assays (*result from provider)
 - on zucchini or cucumber for target
 - according to bacterial species for non-target
- Phylogenetic analyses
 - Multilocus sequence analysis based on concatenated partial sequences of *cts*, *gapA*, *gyrB* and *rpoD* (Berge et al., 2014)
 - PCRs and qPCRs to specifically identify 2ba and 2a phylogenetic clusters and to confirm 2b and 2d phylogroups (Lacault et al., 2023)

5 ALS strains

+ 24 VCZ strains (15 alongside)

 + 12 *P. syringae* pathogenic strains (10 alongside)

41 target strains

33 non-target strains (20 alongside)

 + 3 *P. syringae* non-pathogenic strains

36 non-target strains

Strains	Name	Host	Country	Date	Provider	PCR 4Ba	Pathogenicity assay	Phylo-group	qPCR 4Ba Lac	
ALS target strains	CFBP2104	cucumber	USA	1935	CFBP	-	+ on cucumber	1a	- +	
	CFBP8534	cucumber	USA	1935	CFBP	-	+ on cucumber	1a	- +	
	CFBP6463	cucumber	Hungary	1958	CFBP	-	+ on cucumber	3	- +	
	psLO-1	na	na	na	NSHS	-	+ on cucumber	3	- +	
	PsI7	na	na	na	Syngenta	-	na	na	- +	
VCZ target strains	Psp pep 99	zucchini	na	na	HM.Clause	+	+ on zucchini	2ba-A	+ -	
	PI2855	zucchini	Chile	2007	HM.Clause	+	+ on zucchini	2ba-B	+ -	
	PI37	zucchini	France	2014	HM.Clause	+	+ on zucchini*	2ba-A	+ -	
	Peponis9	na	na	na	Syngenta	+	na	2ba-A	+ -	
	PI	zucchini	France	2005	HM.Clause	+	+ on zucchini*	2ba-B	+ -	
	P69	zucchini	USA	2010	HM.Clause	+	+ on zucchini	2ba-B	+ -	
	P88	zucchini	China	2011	HM.Clause	+	+ on zucchini	2ba-C	+ -	
	PIOO	zucchini	na	2011	HM.Clause	+	+ on zucchini*	2ba-C	+ -	
	ZUM3584	na	na	2023	Syngenta	+	na	2ba-C	+ -	
	P77	zucchini	France	2010	HM.Clause	+	+ on zucchini	2a-D	+ -	
	P70	zucchini	Thailand	2010	HM.Clause	+	+ on zucchini*	2a-D	+ -	
	ZUM3889	na	na	2018	Syngenta	+	na	2a-D	+ -	
	P89	zucchini	France	2011	HM.Clause	+	+ on zucchini	2a-E	+ -	
	P90	zucchini	USA	2011	HM.Clause	+	+ on zucchini*	2a-E	+ -	
	Psp pep 66	zucchini	na	HM.Clause	+	na	2a-E	+ -		
Other <i>Pseudomonas syringae</i> strains isolated from cucurbit seeds	1007009	cucumber	na	2010	VMK	-	+ on cucumber*	2b	+ -	
	1008020	cucumber	na	2010	VMK	-	+ on cucumber*	2b	+ -	
	PAS52384	zucchini	na	2021	GEVES	-	+ on cucumber	2b	+ -	
	PI08	zucchini	Chile	2011	HM.Clause	-	+ on zucchini	2b	+ -	
	PI35	zucchini	India	2014	HM.Clause	-	+ on zucchini	2d	+ -	
	CFBP6576	melon	France	1992	CFBP	-	+ on zucchini	2b	+ -	
	PAS3754	pumpkin	na	2023	GEVES	-	+ on cucumber	2b	- -	
	PAS3795	squash	na	2023	GEVES	-	+ on cucumber	2b	- -	
	CI	cucumber	na	na	VMK	-	+ on cucumber*	2b	- -	
	CFBP6452	melon	New-Zealand	1977	CFBP	-	+ on zucchini	2b	- -	
	PAS3618	pumpkin	na	2022	GEVES	-	+ on cucumber	2b	- -	
	PAS3619	pumpkin	na	2022	GEVES	-	+ on cucumber	2b	- -	
	PAS3620	pumpkin	na	2022	GEVES	-	+ on cucumber	2b	- -	
Xanthomonas cucurbitae	O2-19-01	zucchini	na	na	HM.Clause	-	+ X. cucurbitae	na	- -	
Acidovorax citrulli	Aac 06-90	na	na	na	HM.Clause	-	+ A. citrulli	na	- -	
	PAS156	cabbage	na	na	GEVES	-	+ maculicola	1a	+ -	
	PAS3794	radish	na	2023	GEVES	-	+ maculicola	1a	- -	
	Psm20314	radish	na	2014	VMK	-	+ maculicola	na	- -	
	CFBP1617	beet	USA	1959	CFBP	-	+ apata	2b	+ -	
	PAS3781	beet	na	2023	GEVES	-	+ apata	2b	+ -	
	P. syringae pv. syringae	bean	na	2020	GEVES	-	+ Pss	2d	+ -	
	CFBP4888	bean	USA	1999	CFBP	-	+ Pss	2d	+ -	
	Pseudomonas corrugata	CFBP5321	tomato	France	1986	CFBP	-	+ corrugata	na	- -
	P. syringae pv. apii	CFBP6476	pea	France	2023	GEVES	-	+ apii	na	- -
	P. syringae pv. coriandricola	PAS3574	coriander	na	2022	GEVES	-	+ coriandricola	na	- -
	P. syringae pv. helianthi	CFBP2067	sunflower	Mexico	CFBP	-	+ helianthi	6	- -	
	P. savastanoi pv. phaseolicola	CFBP6471	bean	France	1999	GEVES	-	+ Psp	na	- -
	Pseudomonas viridis	CFBP2107	bean	Switzerland	1927	CFBP	-	+ viridis	na	- -
	P. syringae pv. pisi	CFBP6472	pea	France	1997	CFBP	-	+ pisi	na	- -
	P. syringae pv. glycinea	CFBP6470	soybean	France	1994	CFBP	-	+ glycinea	na	- -
	P. syringae pv. tomato	CFBP1427	tomato	France	1972	CFBP	-	+ tomato	na	- -
	P. syringae pv. cicori	CFBP2101	chicory	na	1929	CFBP	-	na	na	- -
	P. syringae pv. tabaci	CFBP8589	tobacco	USA	1905	CFBP	-	na	na	- -

3. VCZ specificity and sensitivity:

- 4Ba primers/probe (Lacault et al., 2023)
- Amplification of 24 out of 24 VCZ strains
- Amplification of 8 strains out of 12 of *P. syringae* pathogenic
- Amplification of 3 strains out of 4 of 'apata' and 2 out of 2 *Pss* strains (previously reported in Lacault et al., 2023)
- No-amplification of the 4 remaining *P. syringae* pathogenic, the 3 *P. syringae* non-pathogenic and the 5 *ALS* strains and the 28 other non-target

Addition of *P. syringae* strains isolated from cucurbit seeds to the strain collection
 non-pathogenic = non-target
 pathogenic = target (with VCZ primers/probe)

	Expected +	Expected -	Specificity = 87.80 %	
Obtained +	24			