

Les traits d'histoire de vie de la bactérie phytopathogène et glaçogène *Pseudomonas syringae*: un lien entre l'agriculture et les processus atmosphériques



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Pseudomonas syringae, an important plant pathogen



cantaloupe



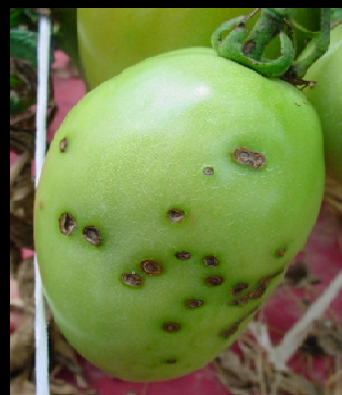
horse chestnut



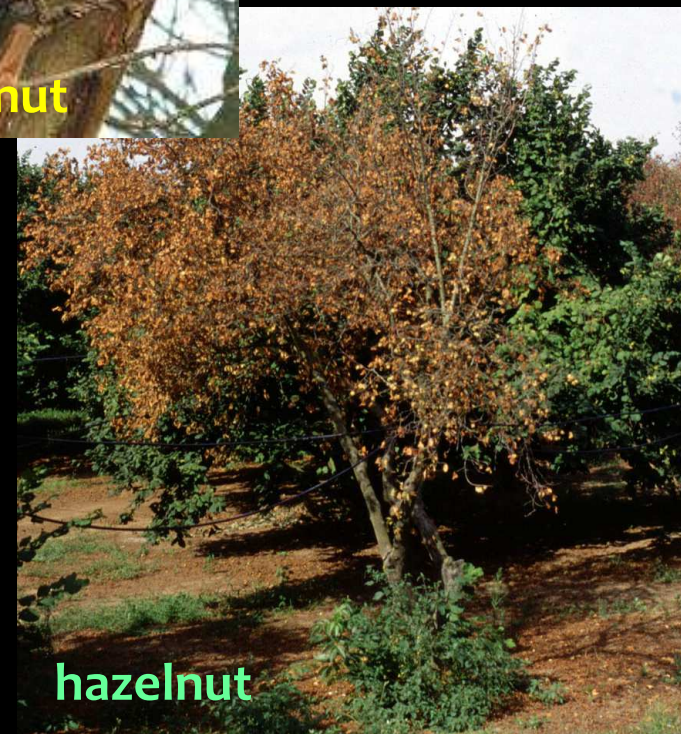
crucifers



kiwifruit

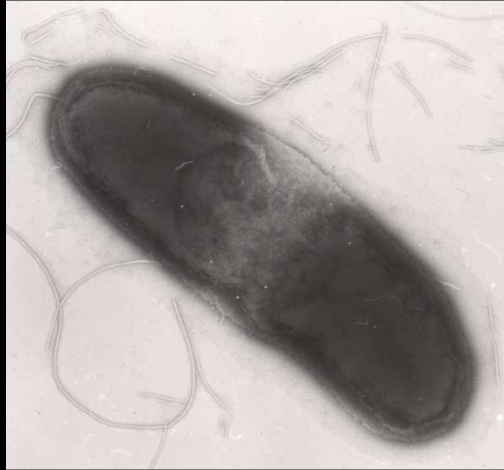


tomato



hazelnut

Aspects of the ecology of *P. syringae* suggest it is ubiquitous in the environment



- Ubiquitous epiphyte
- Cool weather opportunist
- Multiple means of dissemination including airborne
- Ice nucleation active
(watch INA movie)





Are there habitats of *P. syringae* outside of agriculture?



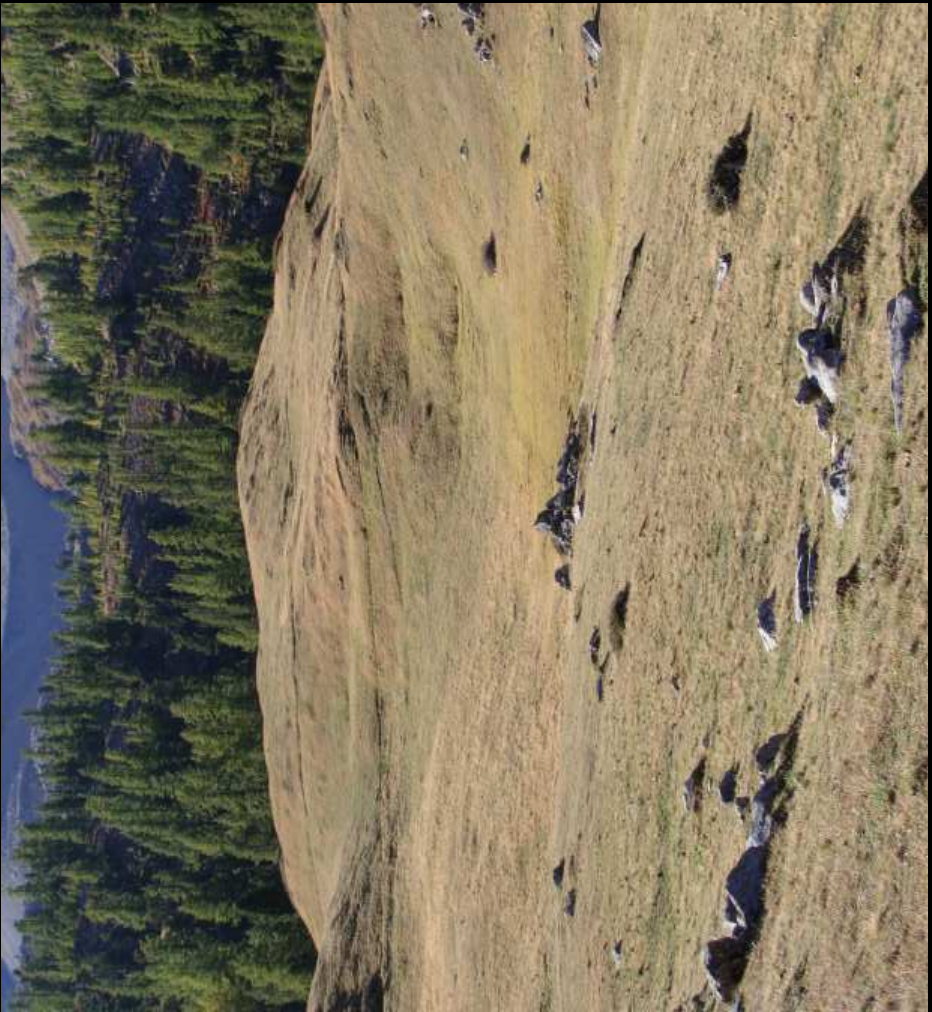
The search for *P. syringae* beyond agriculture





2800 m, Oetzal Valley, Austria





The search for *P. syringae* – agriculture and beyond

Crops (reported in the literature)	$10^2 - 10^8$ b / leaf (or /g)
Wild plants (healthy)	$500 - 10^7$ b / g
Cloud water (Sands 1982; Amato et al 2007)	ca. 10^4 b / L
Rain	$100 - 10^4$ b / L
Snow	$100 - 10^5$ b / L melt water
Leaf litter in alpine prairies	10^8 b/m ² of ground cover
Alpine lakes and rivers	$100 - 10^4$ b / L
Irrigation water basins	$12 - 70$ b / L
Epilithic biofilms in rivers	$50 - 10^4$ b / g
Aerosols (Lindemann et al 1982)	↑ 30 b / m ² / sec

The life history of *P. syringae* is linked to the water cycle

P. syringae inhabits multiple substrates associated with the water cycle and is disseminated among them.



Morris et al. 2008. The life history of the plant pathogen *Pseudomonas syringae* is linked to the water cycle. *ISME Journal* 2:321-334.

New lessons about *P. syringae* ecology from an expanded view of its life history

1. “Natural” environments and non-plant substrates can be significant reservoirs of *P. syringae*.
2. Hydrological and atmospheric circulation transport *P. syringae* over large spatial scales.
3. Pandemic populations are disseminated over long ranges and co-exist with endemic populations.
4. Genetic diversification of *P. syringae* occurs in environmental reservoirs.
5. Ice nucleation activity links pathogen dissemination and evolution – with potential impact on the water cycle

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2. Dissemination with atmospheric circulation and the water cycle

P. syringae trajectory:

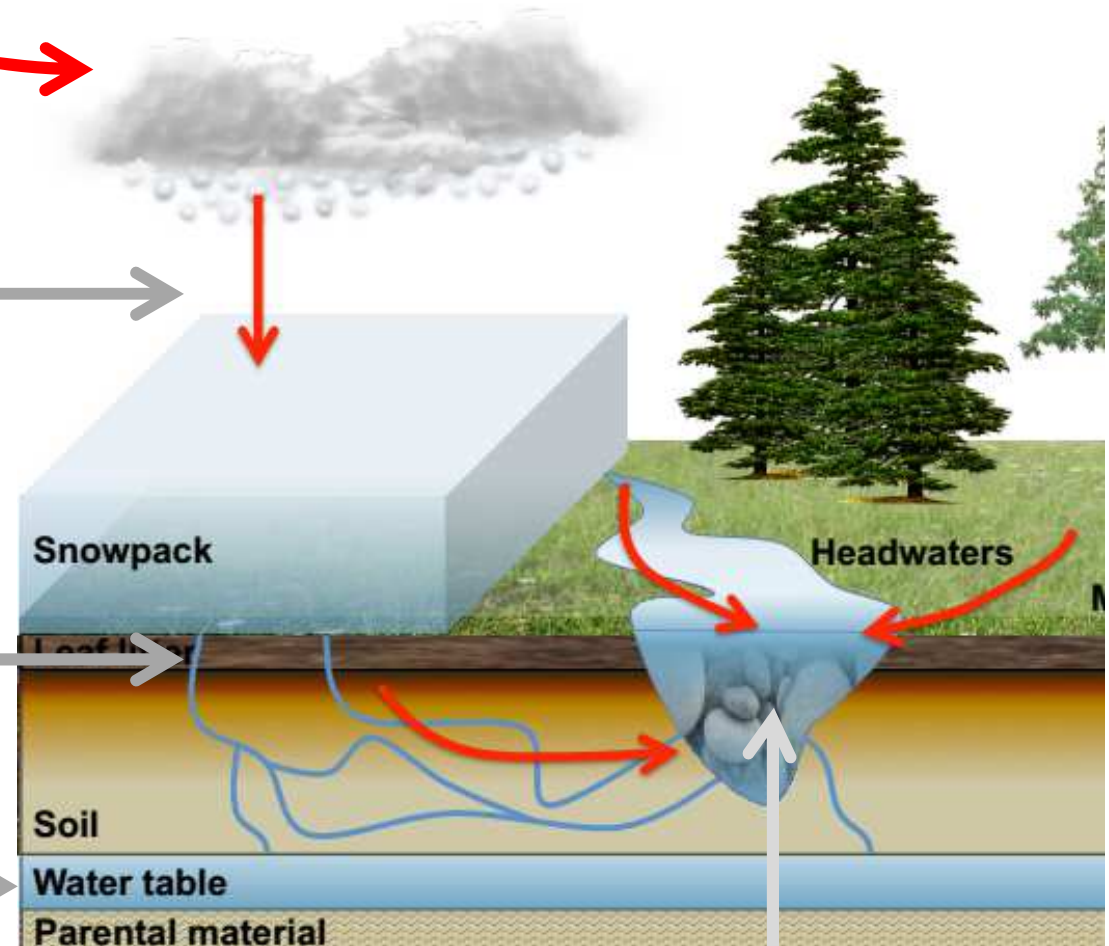
Clouds →

Precipitation →

Percolation water →

Ground water →

Rivers →

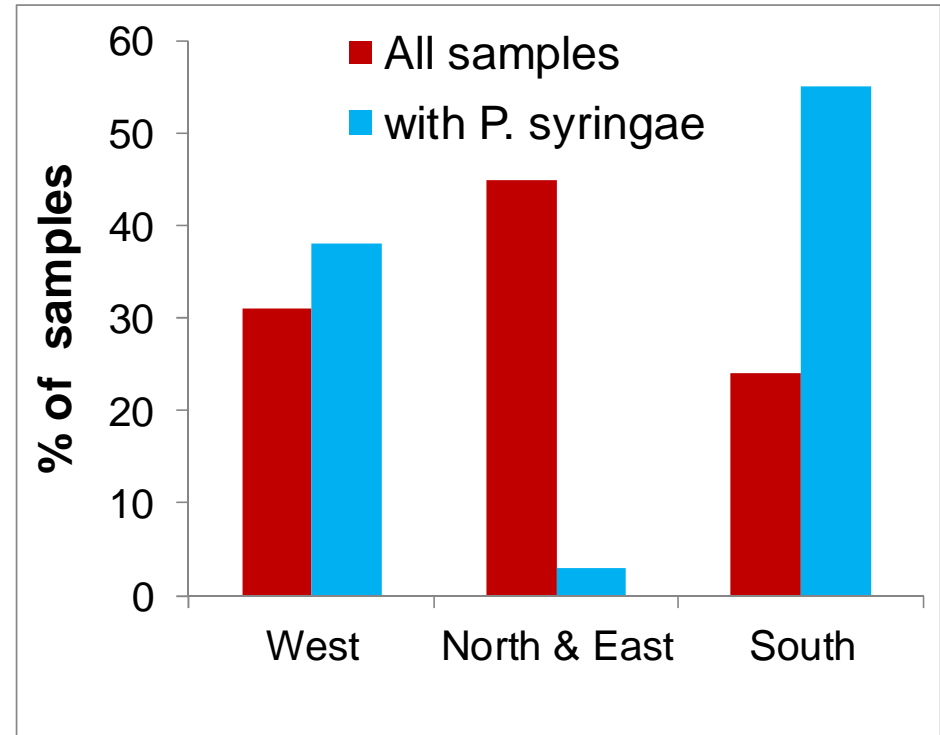
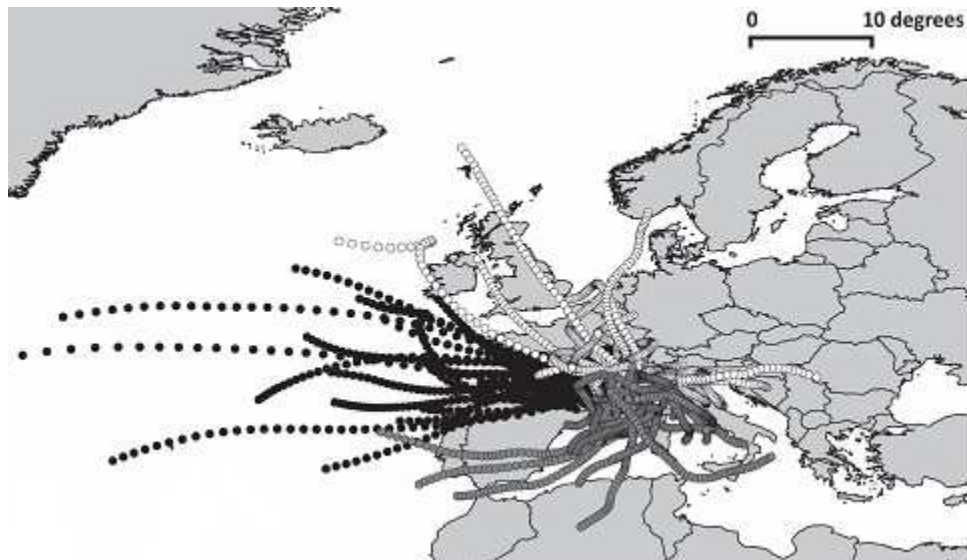


Falling rain and snow carry *P. syringae*, depending on the origin of air masses and precipitation chemistry.

In southern France:

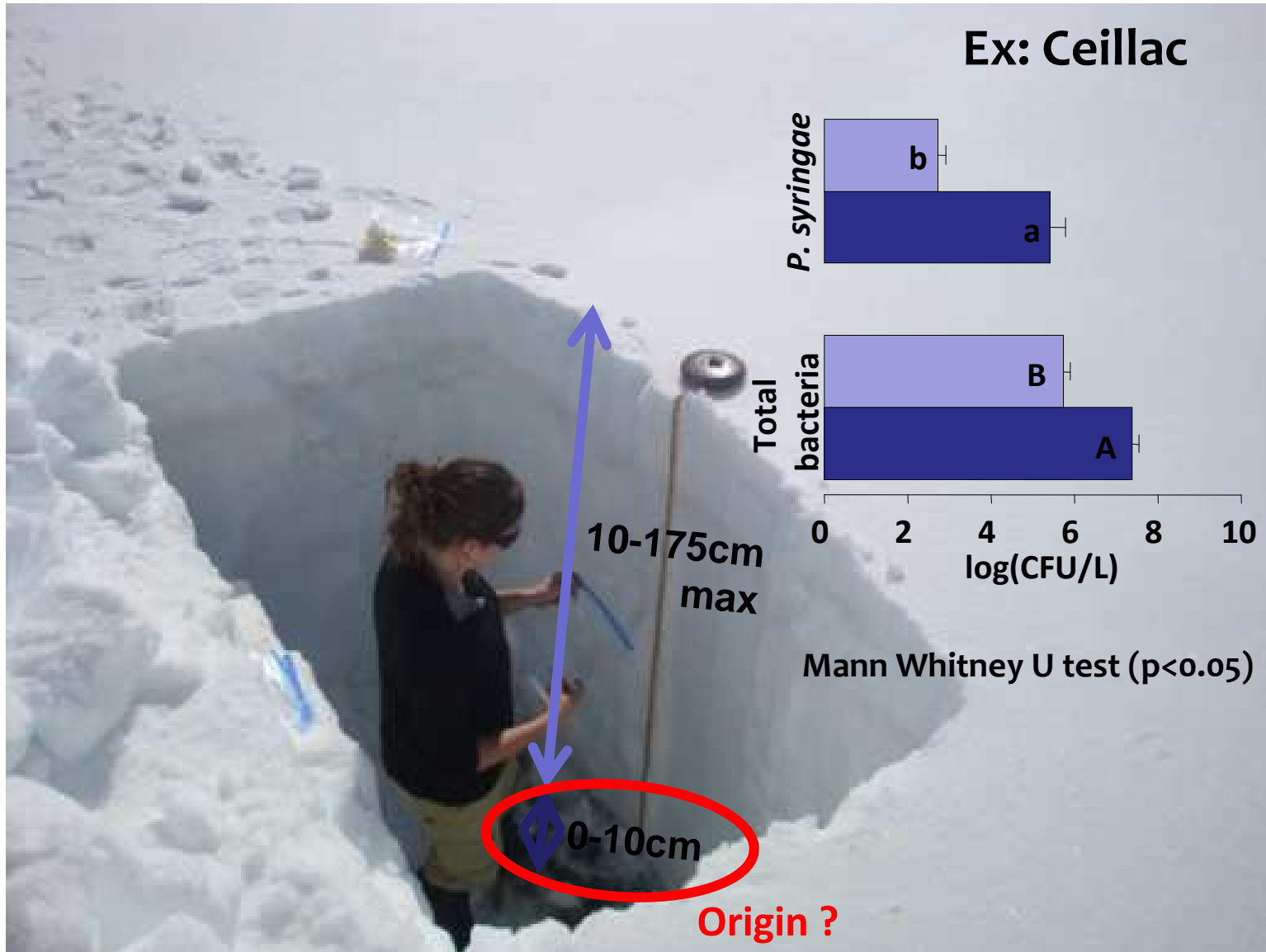
	% w/Ps	<i>P. syringae</i> / L
Rain	65.2	500
Snow	12.3	5000

88 precipitation events 2006 - 2010

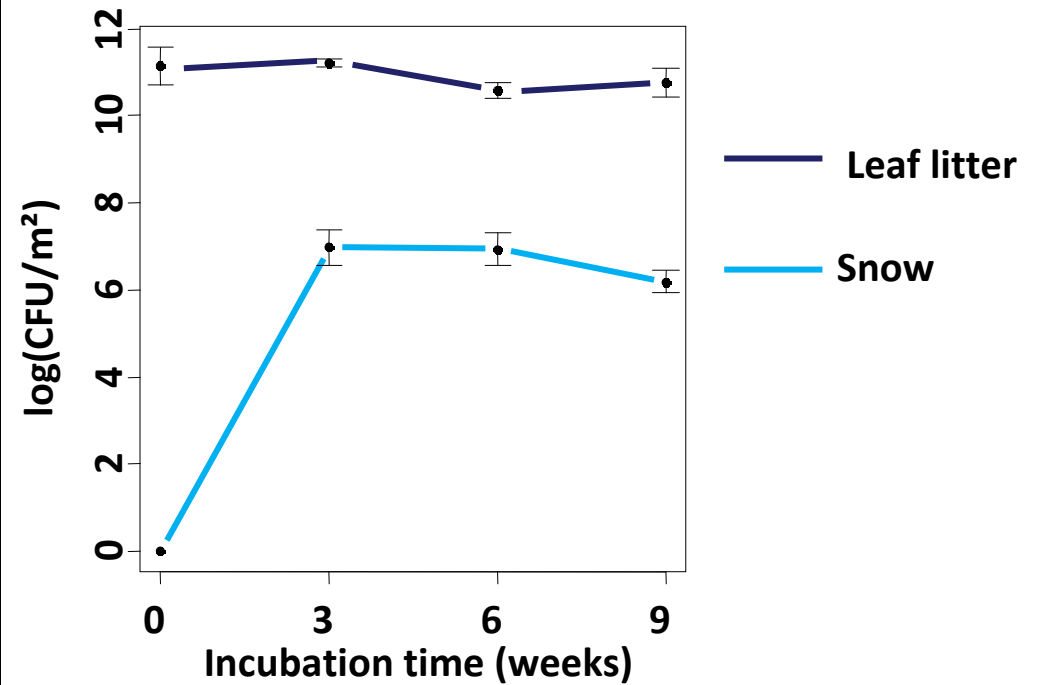
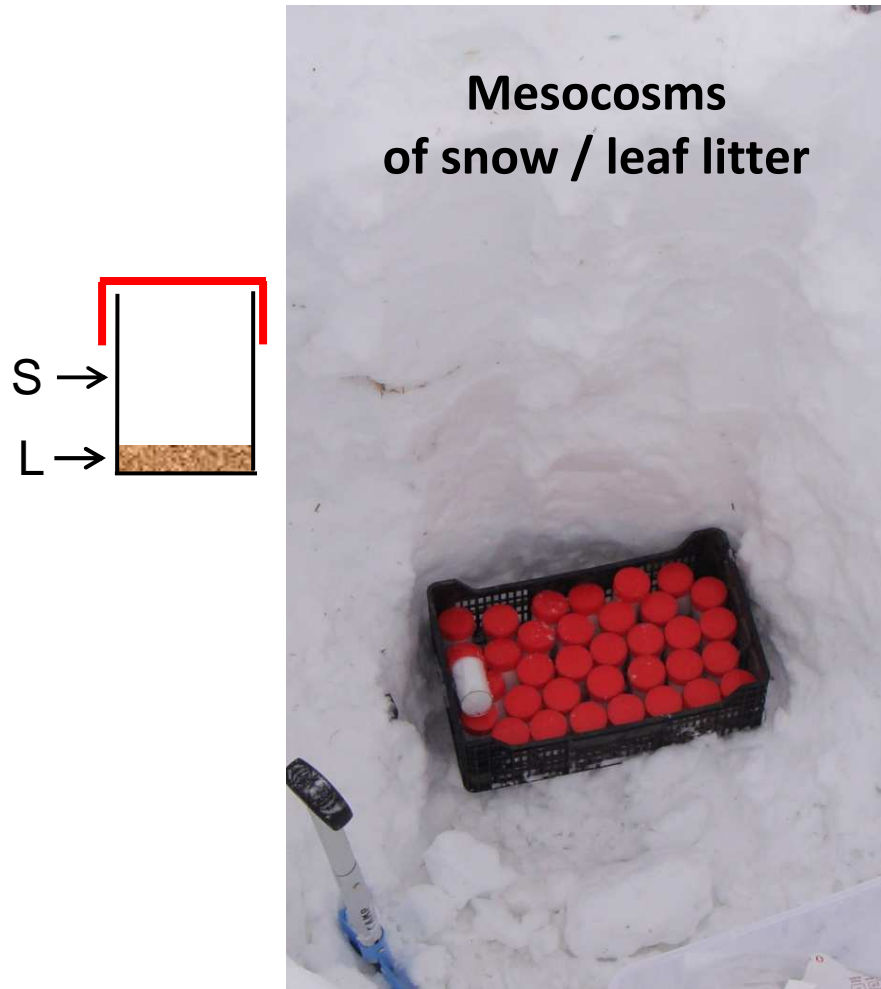


	pH*	Conductivity*
Present	6.21	13 $\mu\text{S}\cdot\text{cm}^{-1}$
Absent	5.76	8 $\mu\text{S}\cdot\text{cm}^{-1}$

P. syringae survives in snowpack during the winter
The population density in snowpack is stratified



Emigration of *P. syringae* from litter to snowpack contributes to a stratified population density



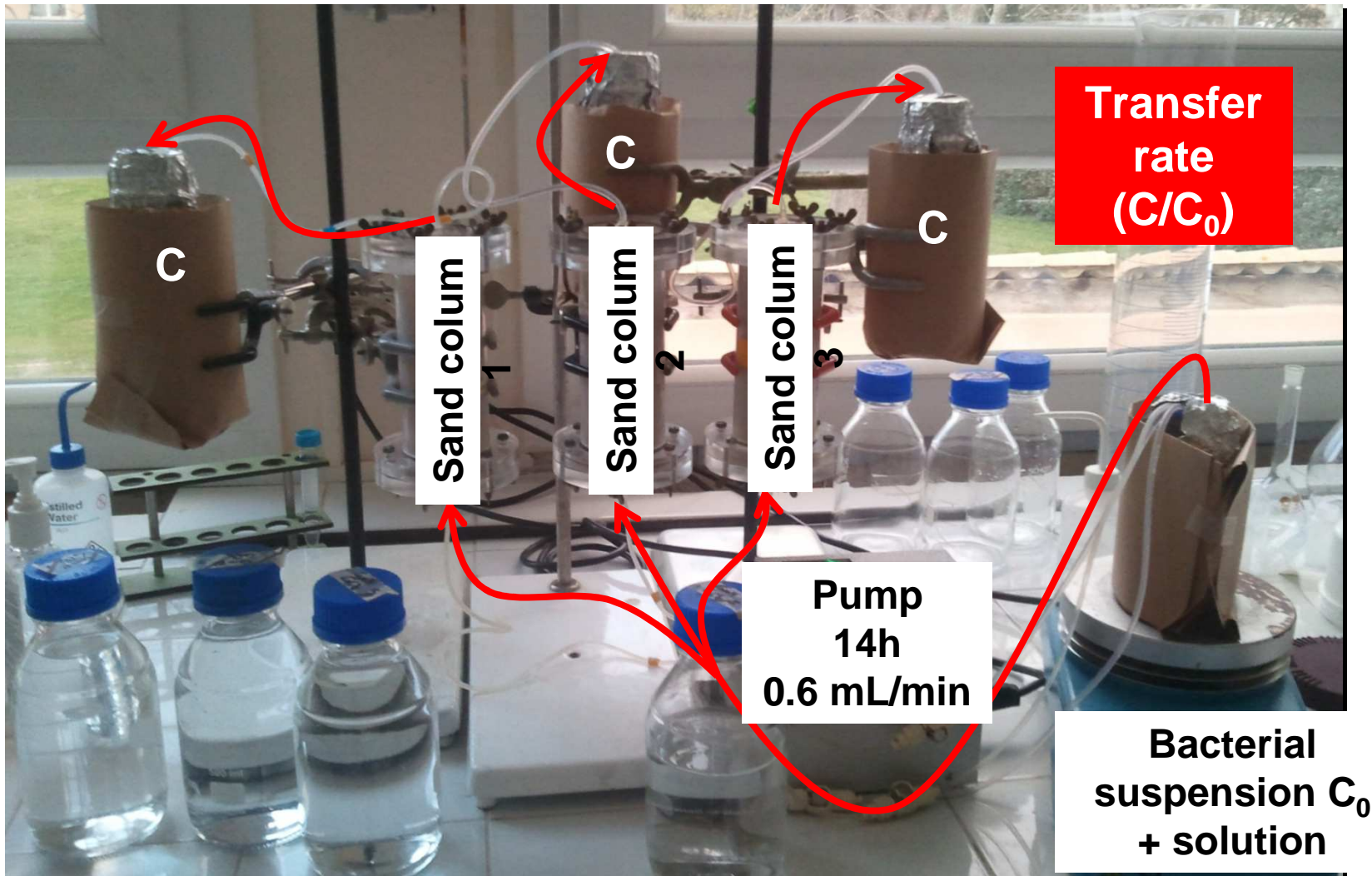
Kruskall Wallis ($p < 0.05$)

P. syringae passes through the soil into surface and ground waters.

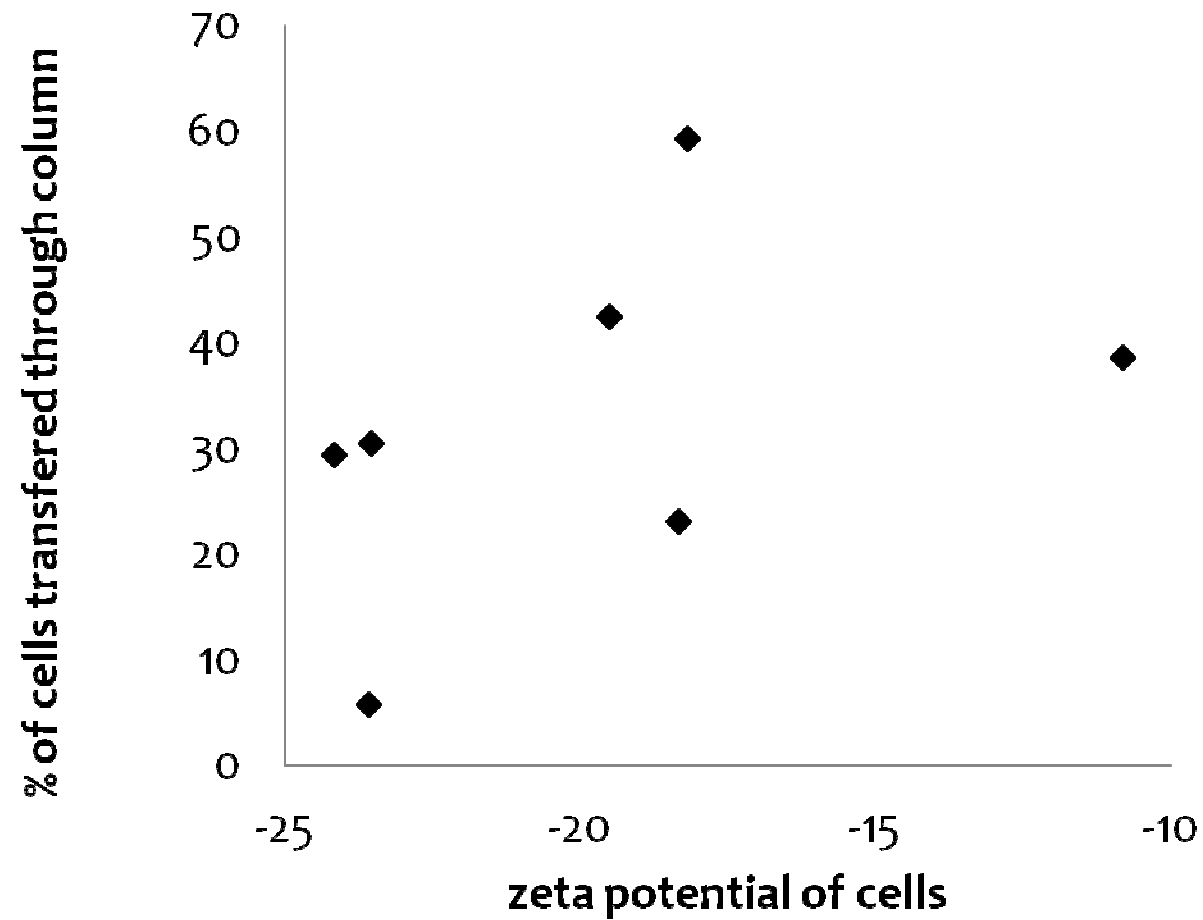


Monteil et al. 2013
Environ. Microbiol. **16**:2038–2052..

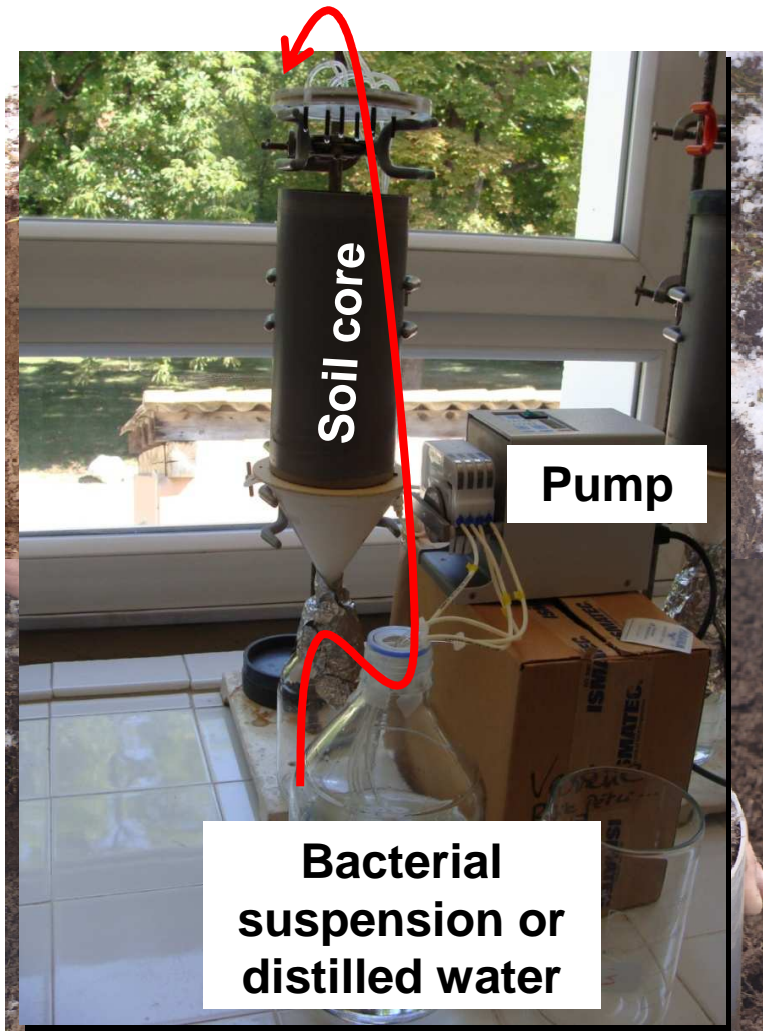
P. syringae transport rate through sand columns



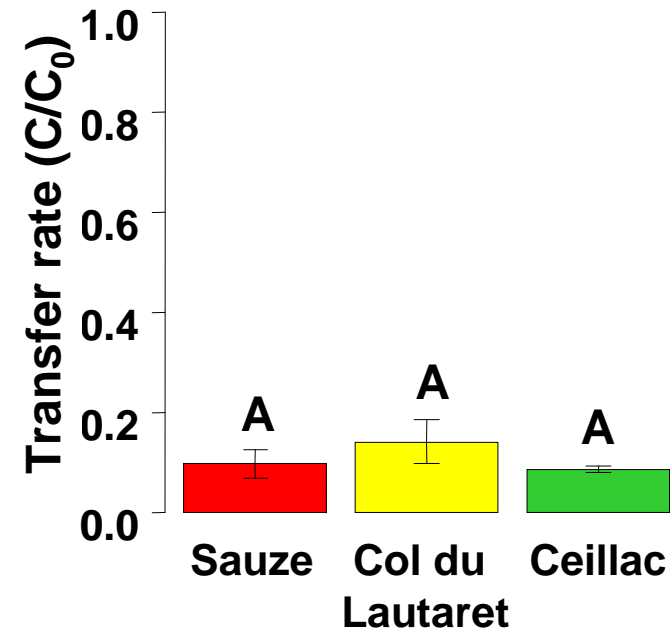
P. syringae transport rate through sand columns:
7 strains representing the range of zeta potential



The rate of transfer of *P. syringae* through soil columns is comparable to that in sand columns



Strain TA022
resistant to rifamycin



Kruskall Wallis test ($p < 0.05$) 20

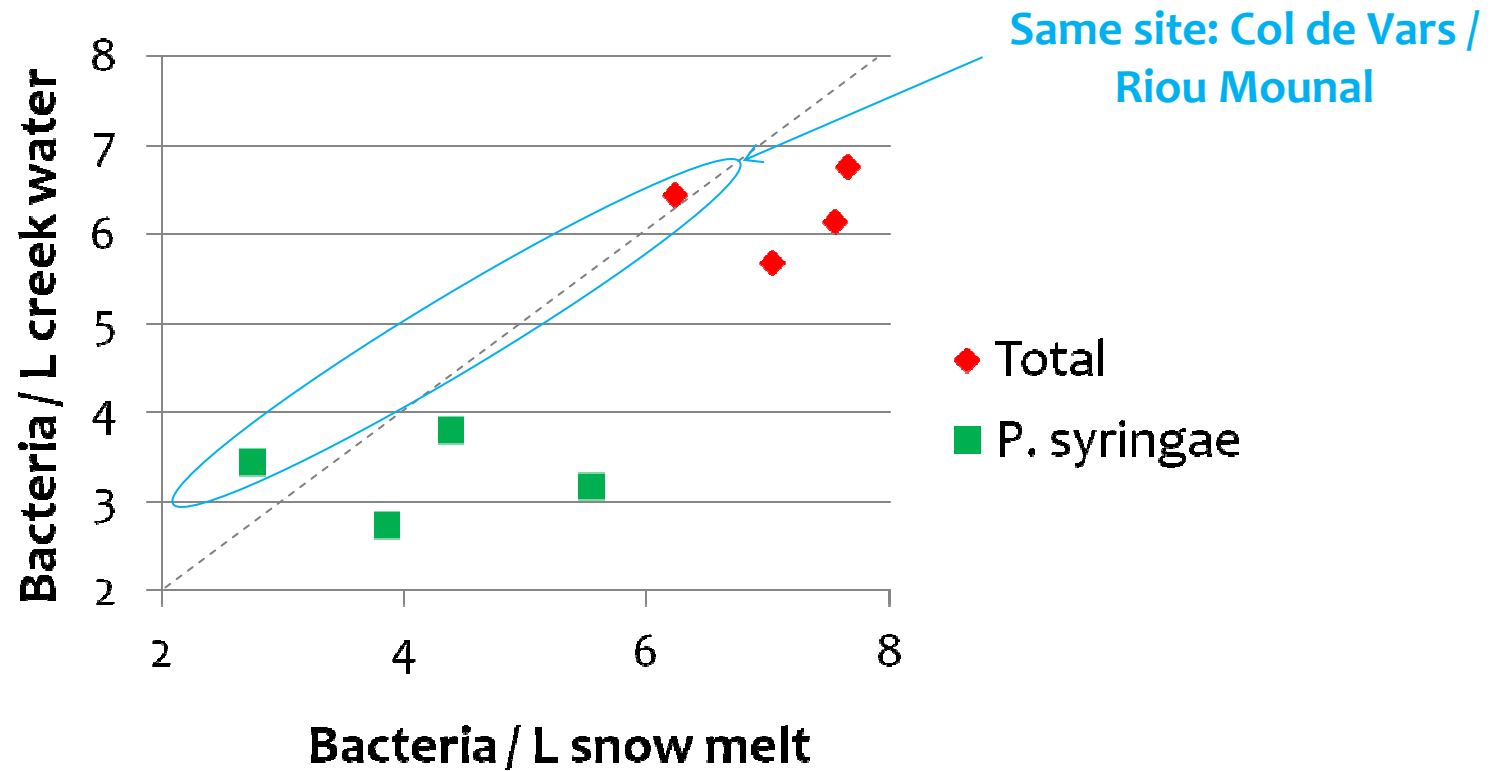
Soil core also leached indigenous
P. syringae

Concentrations of bacteria in creeks are usually less than those in snowmelt

Catchment basin, creek	Substrate	pH	EC ($\mu\text{s}/\text{cm}$)	Population densities (CFU/L)	
				Total bacteria	<i>P. syringae</i>
Ceillac, Pisse					
	Snow	5.62 ^c	3 ^e	7.03 ^a	3.85 ^{bc}
	Water	7.90 ^a	609 ^a	5.67 ^{dvv}	2.73 ^e
Col de Vars, Riou Mounal					
	Snow	5.40 ^{bcd}	4 ^{de}	6.23 ^{abcd}	2.74 ^{bc}
	Water	7.89 ^a	216 ^c	6.46 ^{cvv}	3.43 ^{ce}
Col du Lautaret, Roche Noire					
	Snow	6.30	6	7.55 ^a	5.54 ^a
	Water	7.78 ^{ab}	143 ^d	6.13 ^{bcd}	3.16 ^{cde}
Super Sauze, Soudane					
	Snow	5.30 ^d	4 ^e	7.65 ^a	4.38 ^b
	Water	7.88 ^a	324 ^b	6.75 ^{bvv}	3.79 ^c

Mann Whitney ($p < 0.05$)

Concentrations of bacteria in creeks are usually less than those in snowmelt – but not always



Association of *P. syringae* with the water cycle contributes to pandemic and endemic biogeography

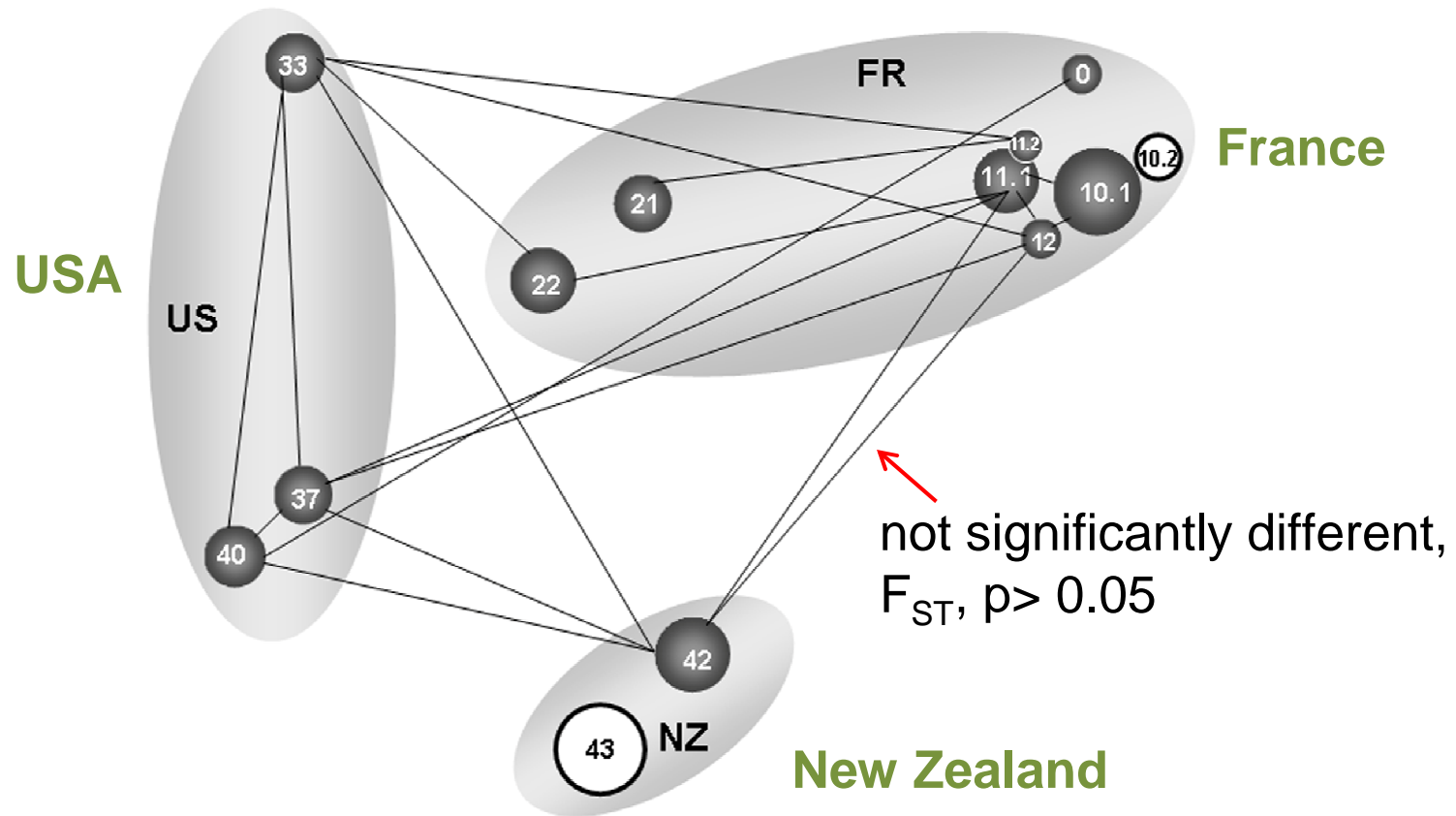
Population structure of *P. syringae* populations in water:

236 strains, 11 sites, 3 continents

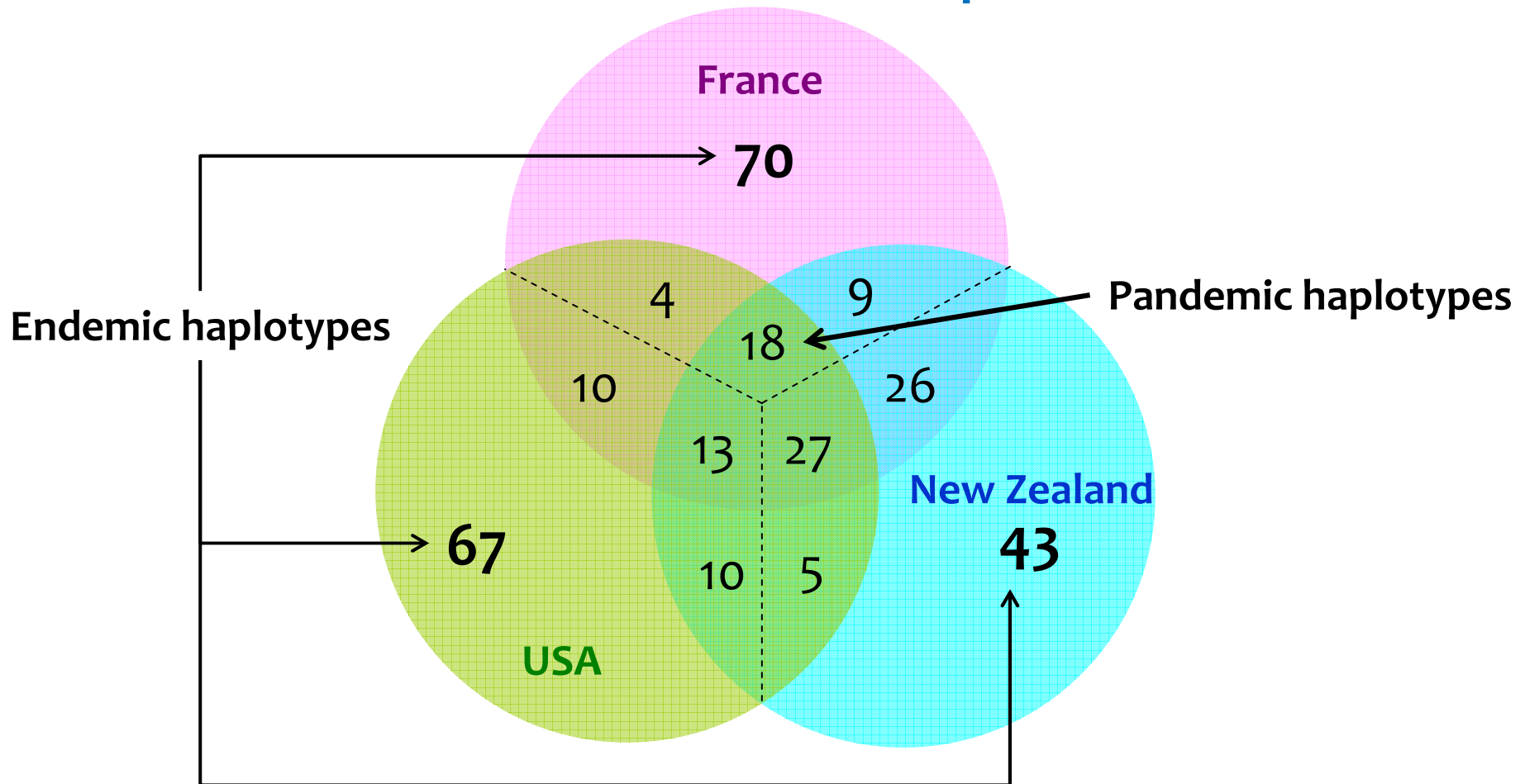
population genetics based on conserved housekeeping genes



Pandemic genetic structure



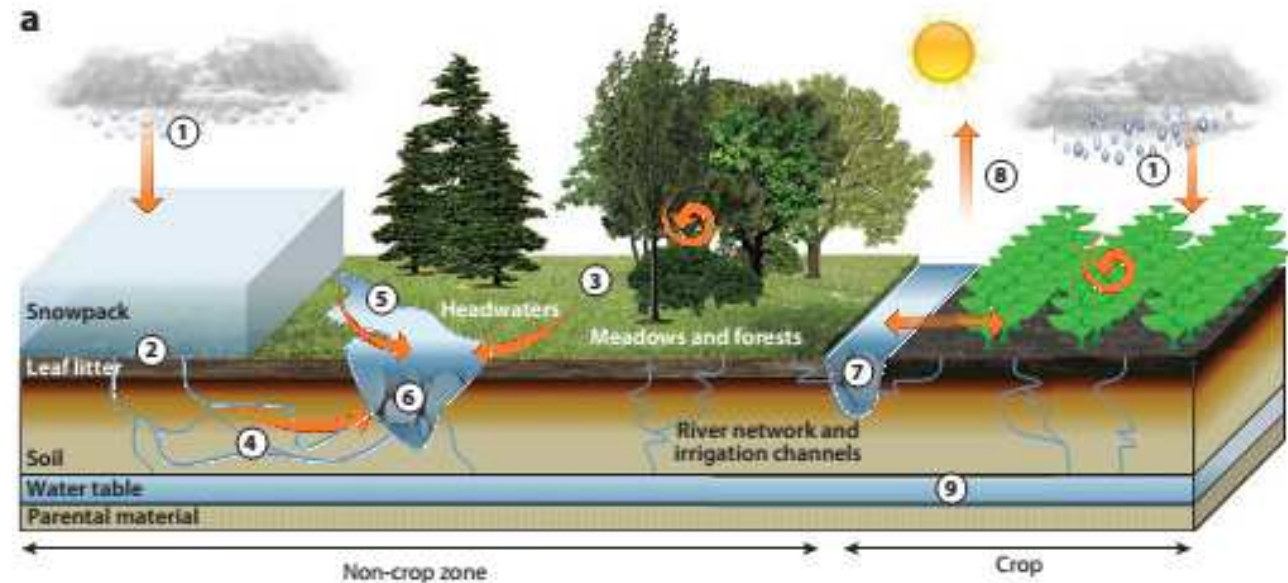
... with an endemic component



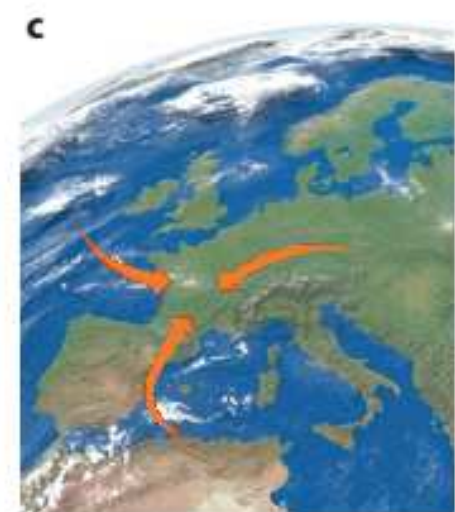
Percent of haplotypes that are unique to each region and in common between regions.

The water cycle links agricultural and natural habitats

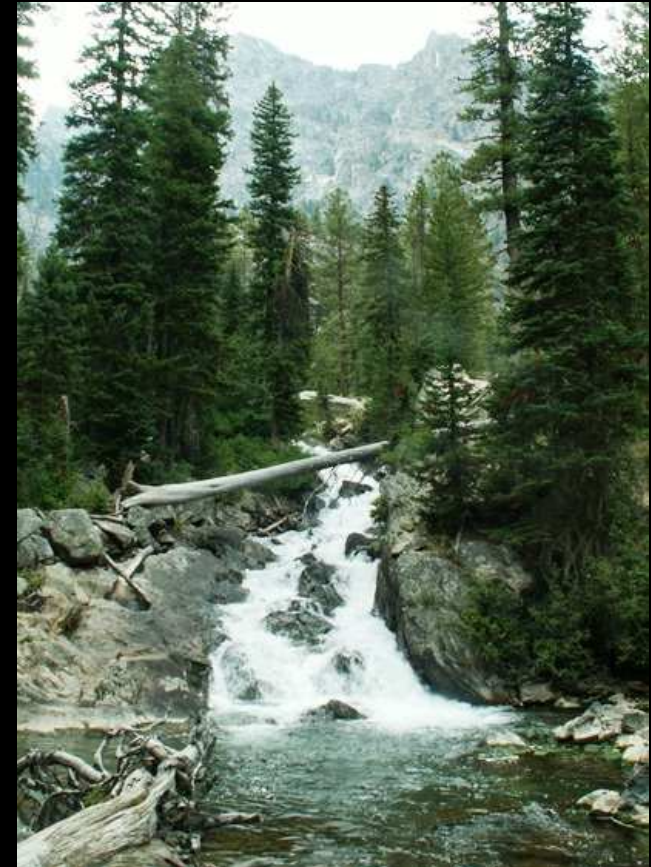
at a range of scales



We have detected *Ps* in the water table at 7m depth



Fresh water: a major reservoir of *P. syringae*



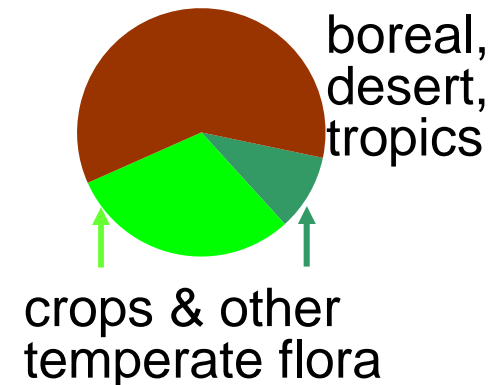
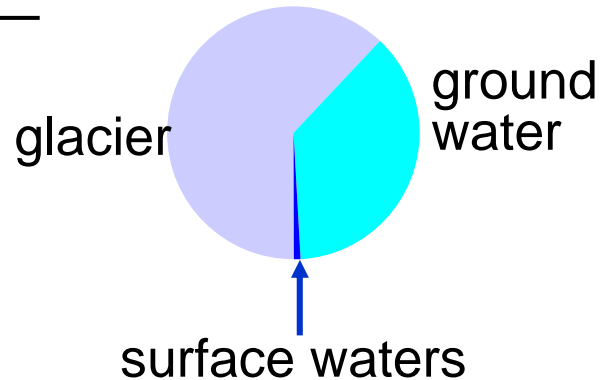
Comparison of metapopulation sizes of *P. syringae*: fresh-water vs. plant-associated



Total potential habitat:

$5 \times 10^7 \text{ km}^3$

10^9 km^2



Known to be colonized:

$2 \times 10^5 \text{ km}^3$

$4 \times 10^8 \text{ km}^2$

Density:

1000 cfu / L

100 – 10⁴ cfu / cm²

10^{15} cfu / L

$10^{12} - 10^{14} \text{ cfu / km}^2$

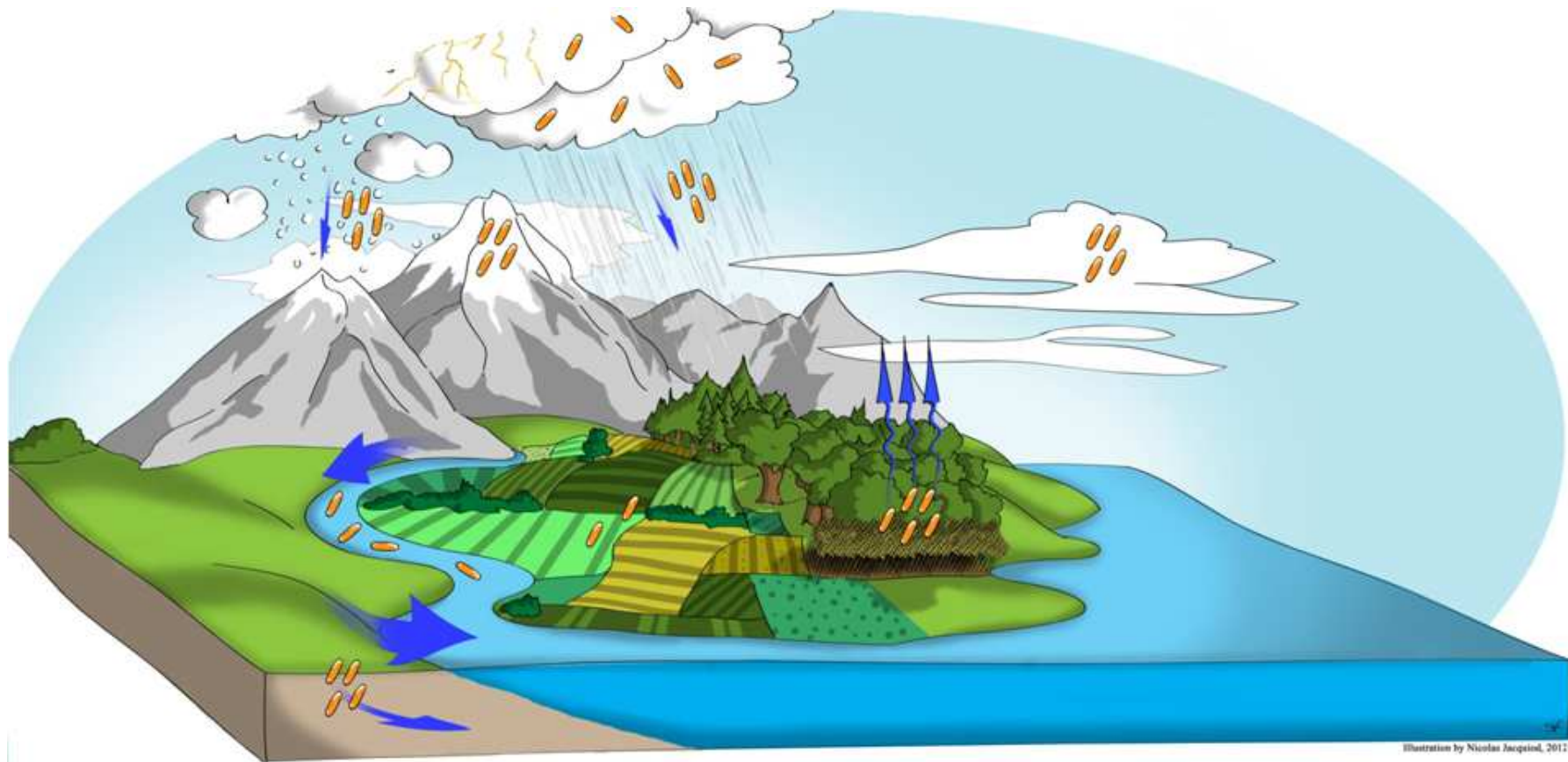
Population size:

$2 \times 10^{20} \text{ cfu}$

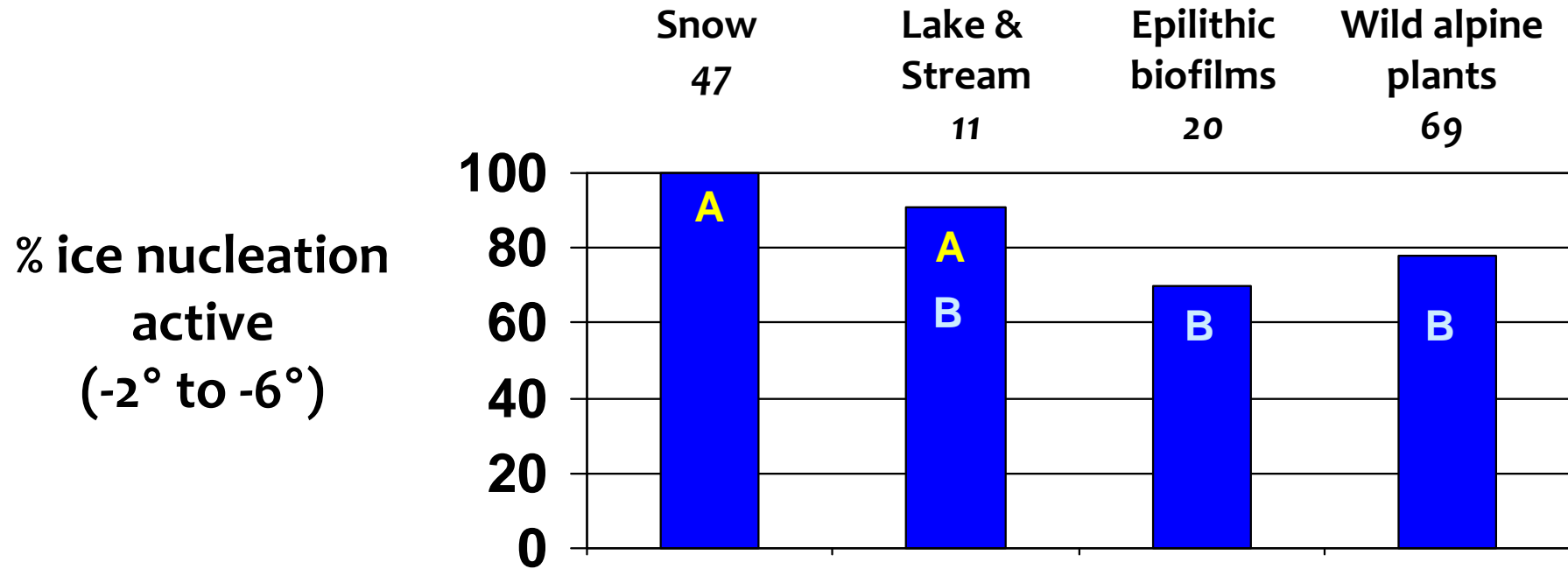
$4 \times 10^{20} - 10^{22}$

cfu

5. Ice nucleation activity links pathogen dissemination and evolution – with potential impact on the water cycle



Precipitation positively selects for INA *P. syringae*



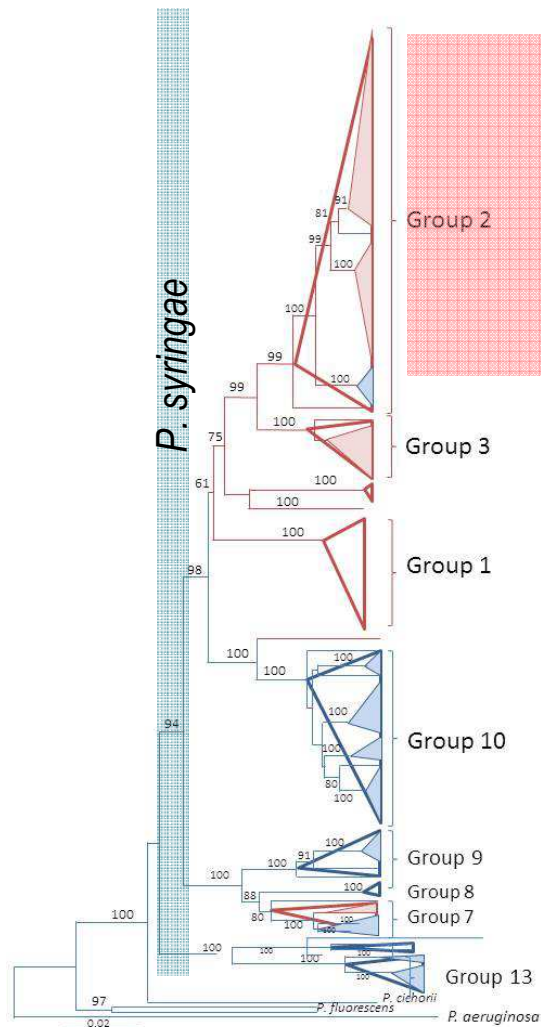
pair-wise χ^2 , $p \leq 0.05$

Morris et al. 2008. ISME Journal 2:321-334

In clouds, only ca. 10% of *P. syringae* strains are INA

Joly et al. 2013. Atmospheric Environment 70:392-400

Precipitation positively selects for INA and highly aggressive *P. syringae* with broad host ranges





Phylogroup 2 contains

- the most ubiquitous strains
- with the broadest host range and
- the most frequently INA

Broad host range is correlated with ice nucleation activity

Number of strains

		Ice nucleation activity		
		0	1	2
Host range (proxy)				
	0	24	22	37
	1	3	10	35
	2	3	12	54



$p < 0.0001$, Fisher's exact test

Broad host range is correlated with ice nucleation activity

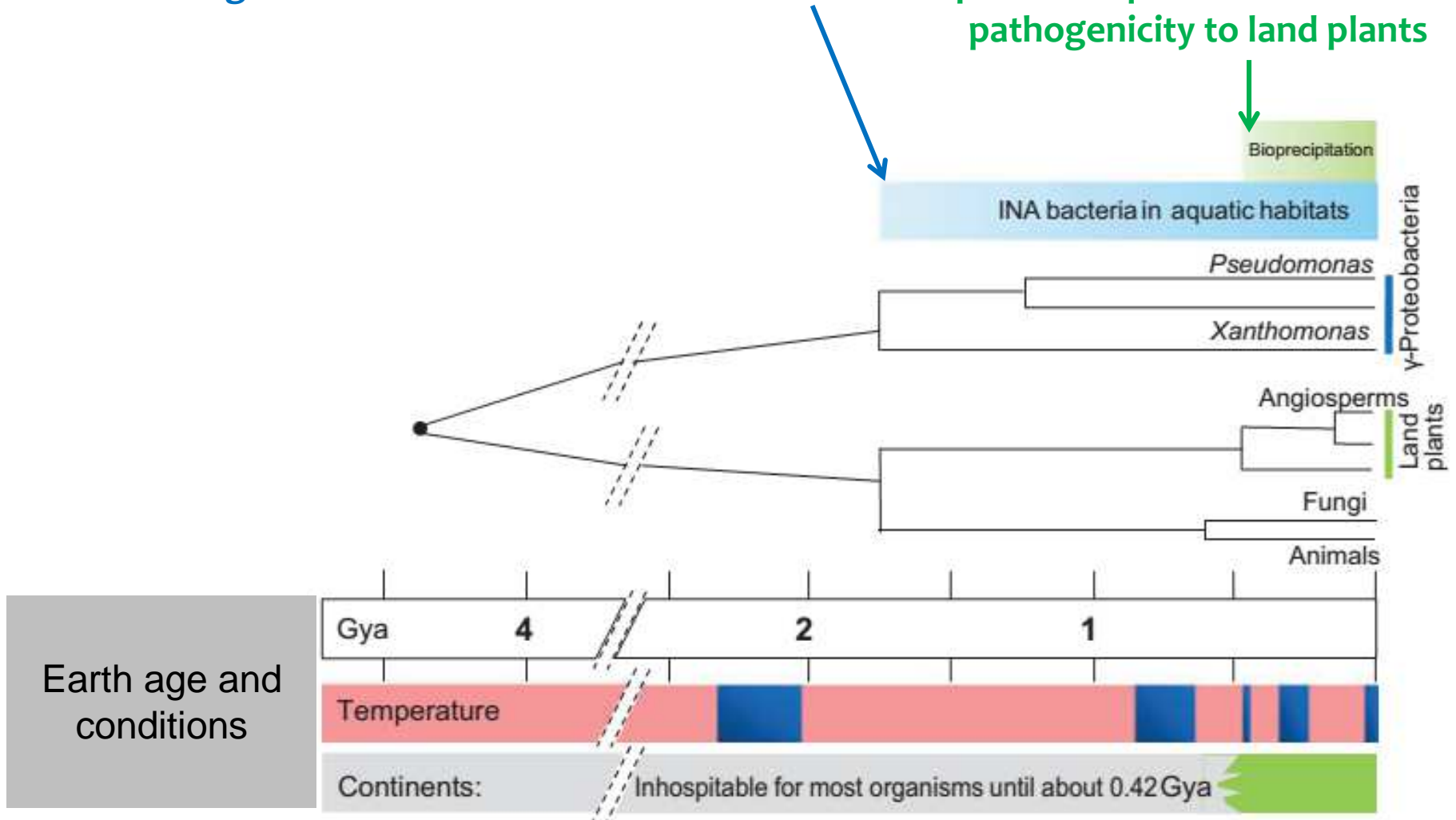
Pathogenicity and INA are NOT related mechanistically.

How have they co-evolved?

Evolution of ice nucleation activity and pathogenicity

gene for INA in common ancestor

Repeated acquisition of factors for pathogenicity to land plants



Bioprecipitation: a feedback cycle linking Earth history, ecosystem dynamics and land use through biological ice nucleators in the atmosphere

Global Change Biology

Morris et al. 2014. Global Change Biology 20:341-351



