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Climate adaptation in the lineages of the Swiss needle cast agent *Nothophaeocryptopus gaeumanii*

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What's up with SNC in British Columbia?

David Rusch will tell you
in the next presentation

Me: I will talk about SNC
lineages & adaptation and
of course some ACGT stuff



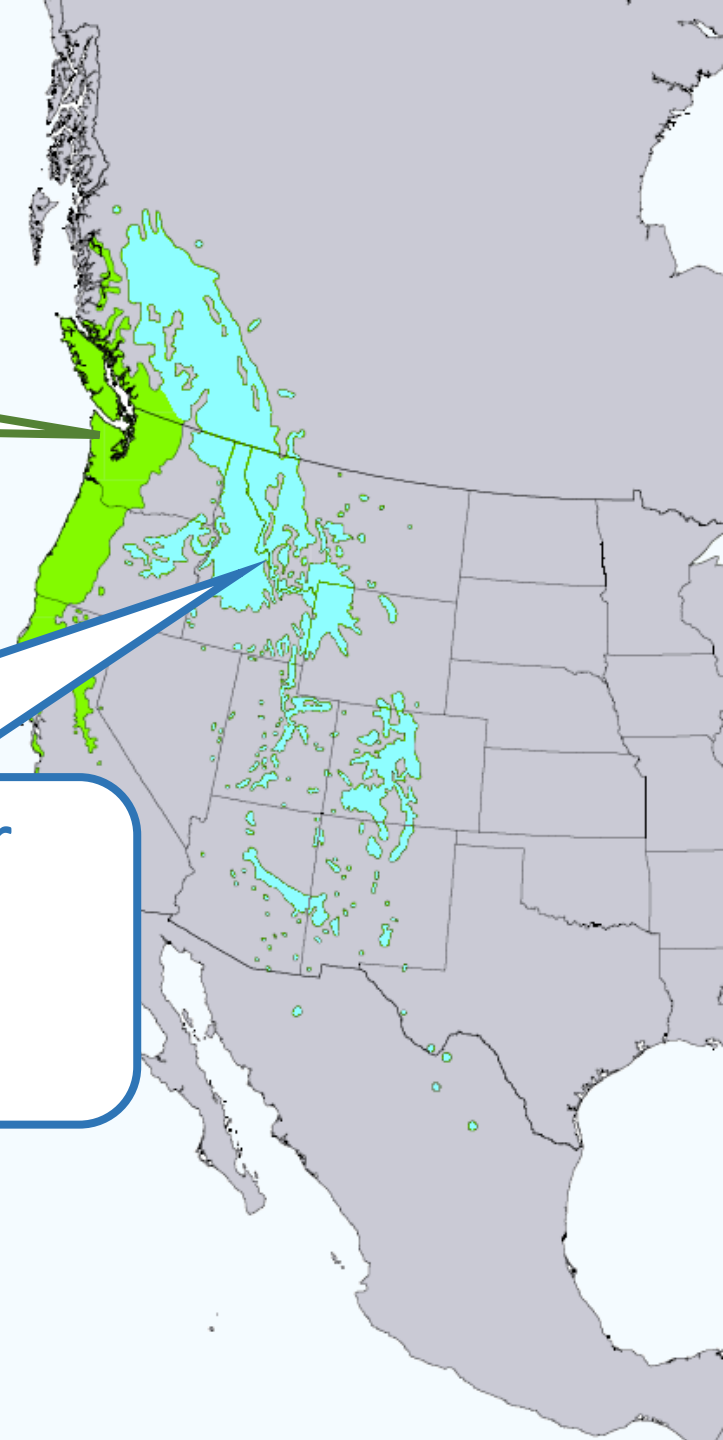
SNC in the Chilliwack river valley in British Columbia

One species, three genetic lineages

Coastal Douglas-fir
(*P. menziesii* var. *menziesii*)
Lineage 1c and Lineage 2

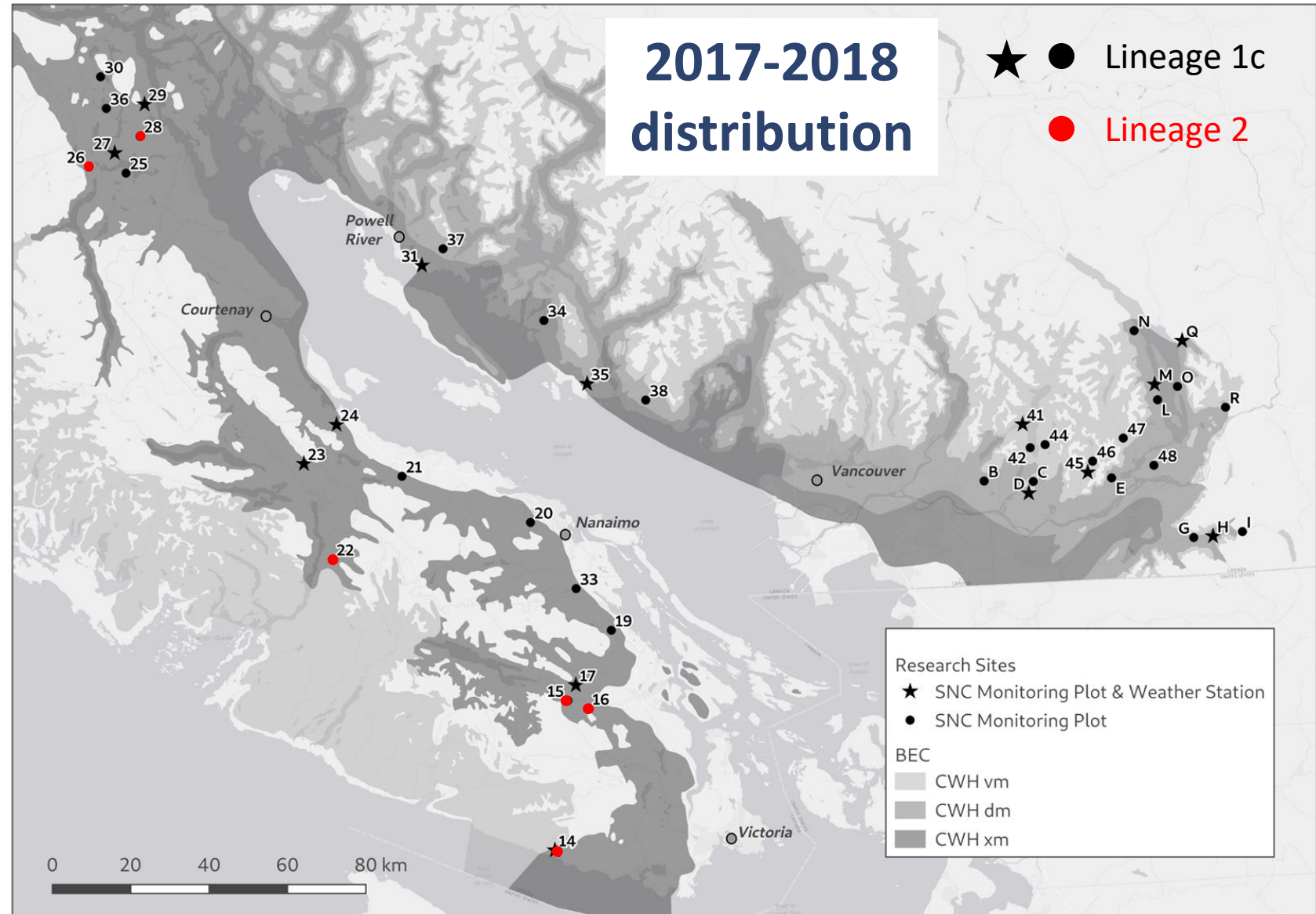
Rocky Mountains Douglas-fir
(*P. menziesii* var. *glauca*)
Lineage 1i

Collection covering OR, WA, ID and BC
200 SNC genomes sequenced
3 lineages morphologically identical

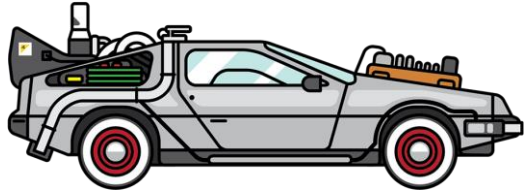


Lineage 2 seems to like the humidity and cooler temperatures of the coast

2022-2023's survey suggests that L2 is becoming difficult to find



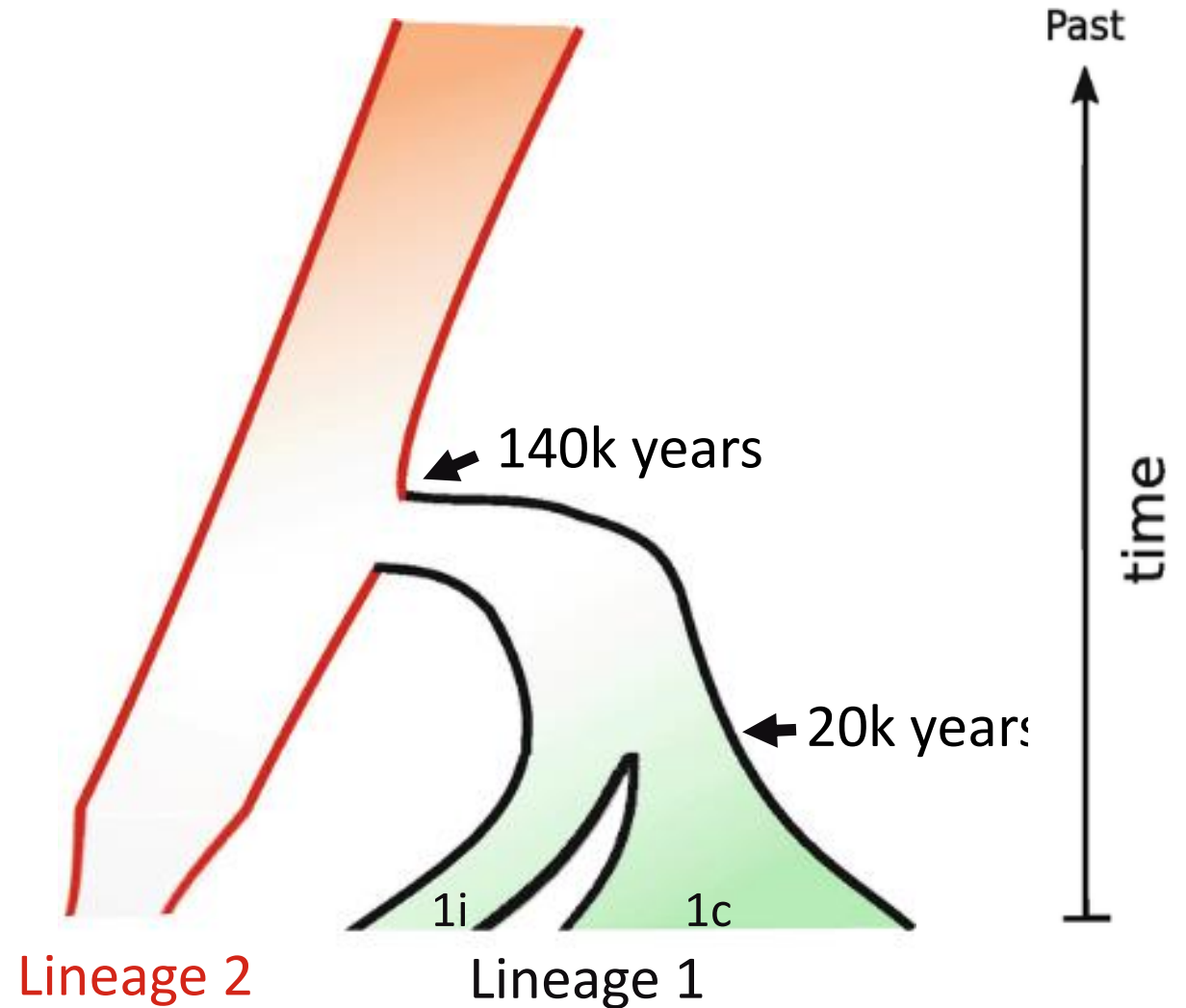
Riding the "genomics time travelling machine"



Genomes were used to infer SNC evolutionary & demographic history

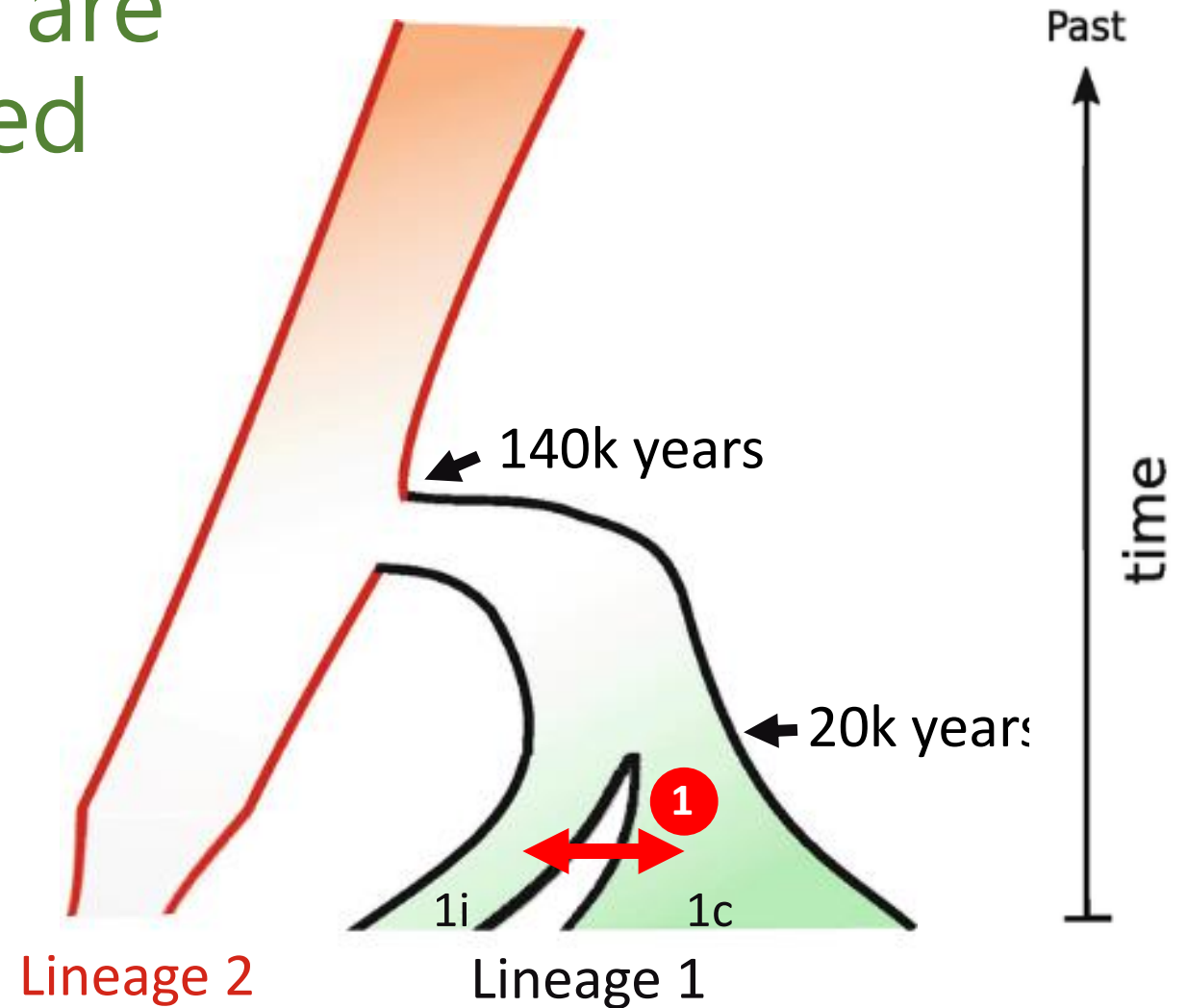
Time estimate of lineages divergence

Demographic expansion of Lineage 1c



The three genetic lineages are not reproductively isolated

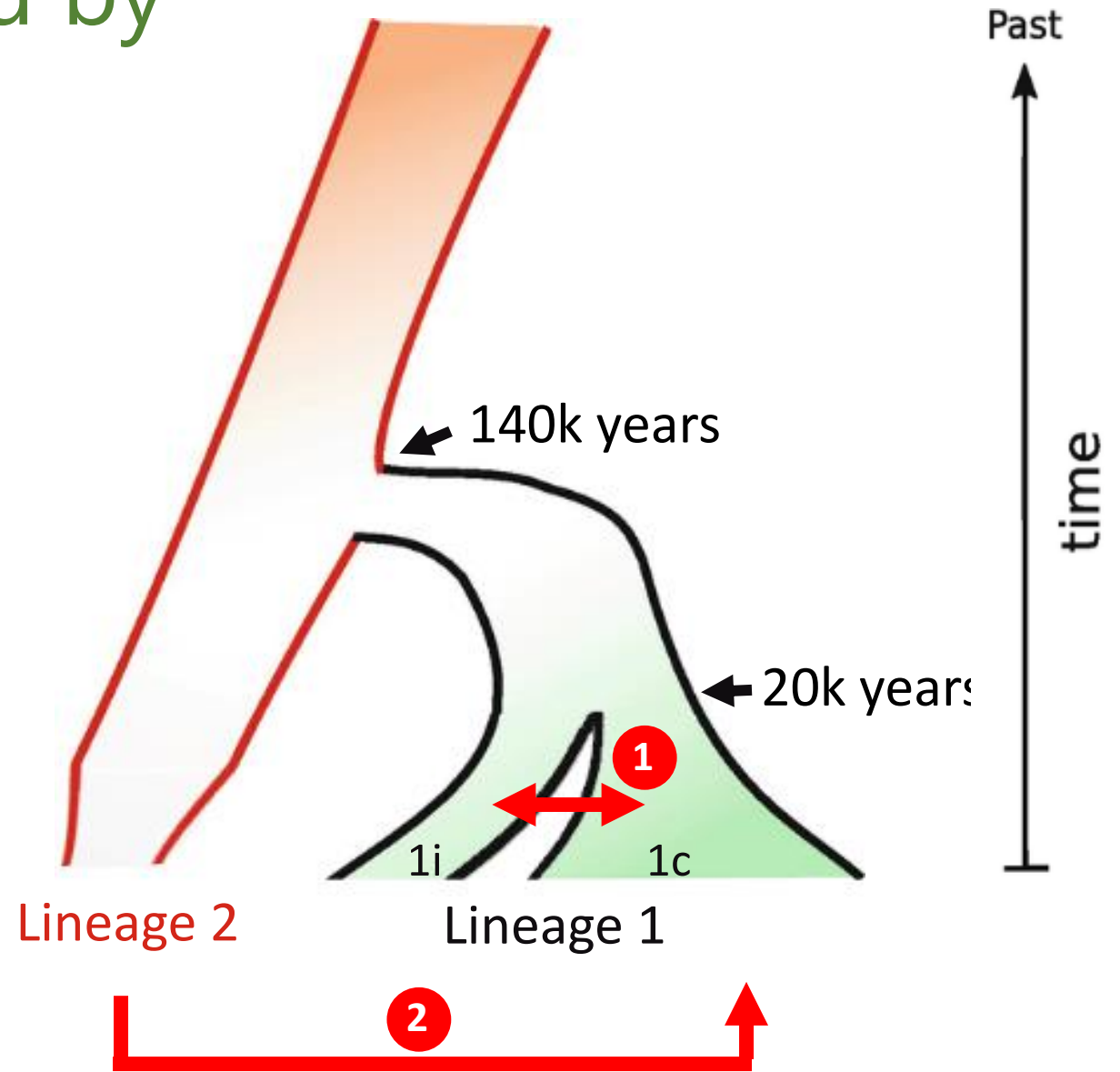
	Model likelihood
No gene flow	-2847.73
1 Lin1c <-> Lin1i	-2559.29



Lineage 1c is introgressed by Lineage 2

	Model likelihood
No gene flow	-2847.73
1 Lin1c <-> Lin1i	-2559.29
2 Lin2 => Lin1c	-2511.67

No barriers to gene flow between the three lineages



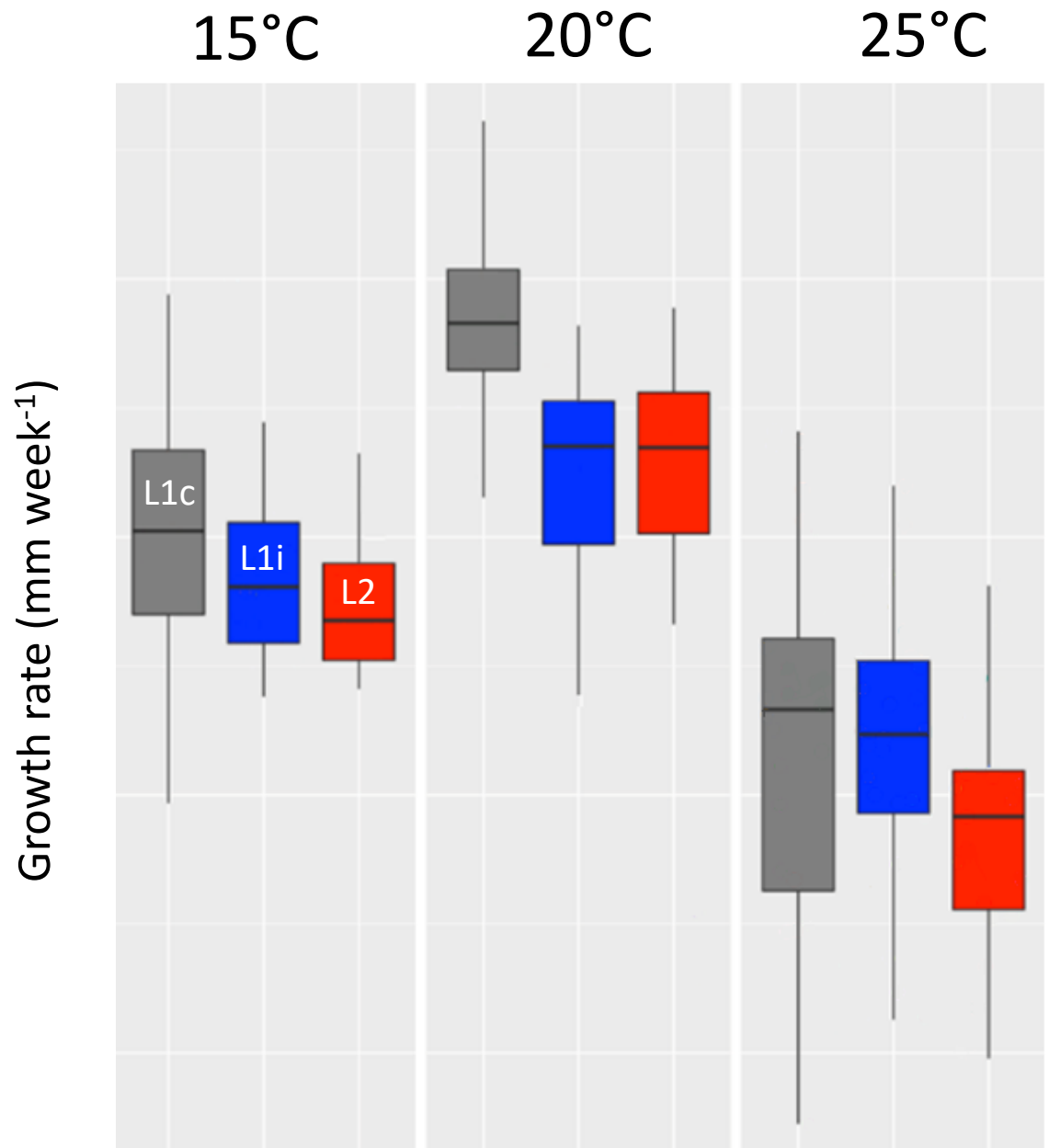
Lineage 1c has a higher adaptive plasticity than Lineage 2

1 - Temperature

Lineage response (30 ind./lineage) to temperature changes differs

Growth rate at different temperatures:

Lin. 1c > Lin. 2 & Lin. 1i



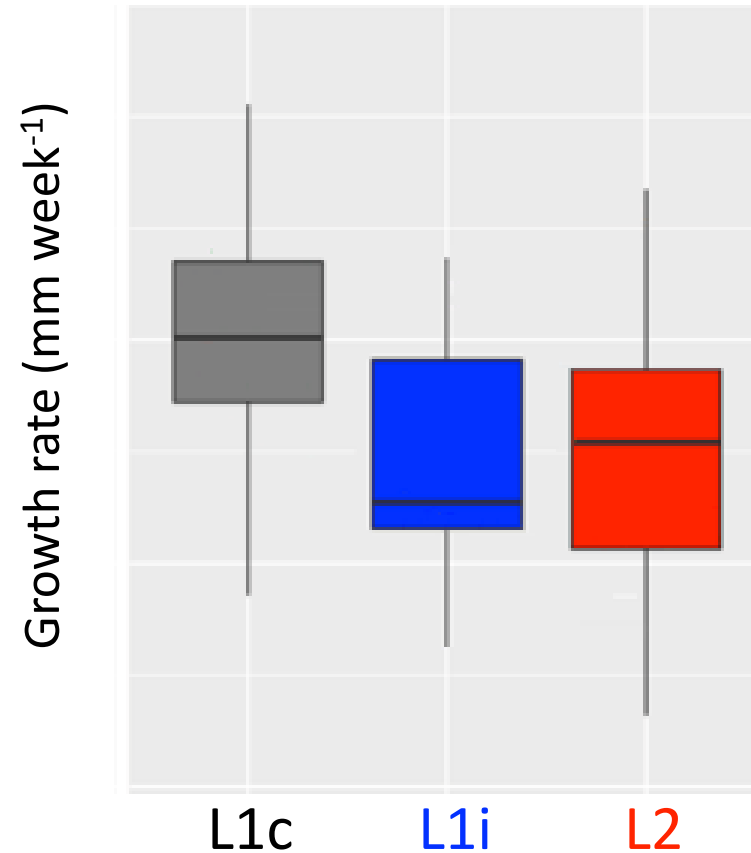
Lineage 1c has a higher adaptive plasticity than Lineage 2

2 - Drought

Growth rate at 20°C under drought stress:

Lin.1c > Lin. 2 & Lin. 1i

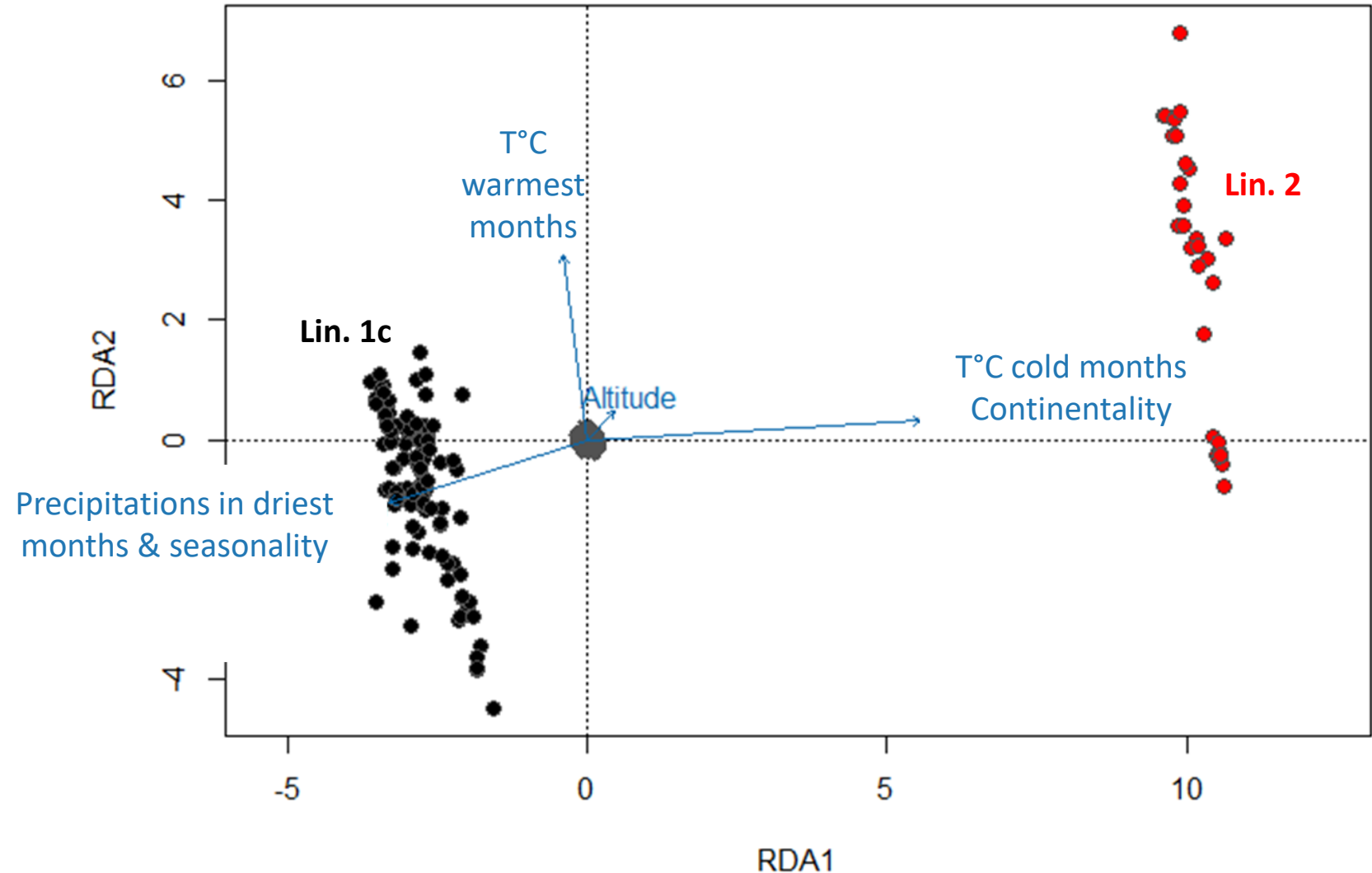
20°C + drought
(3% MEA + 10% glycerol)



The genetic variation between SNC lineages is explained by spatial and environmental variables

Adaptive genetic variation in Lin. 1i explained by host specificity and elevation

Lin. 1c and Lin. 2: can be explained by only few climatic variables ($P < 0.001$)



Climate is triggering some changes in lineages distribution

Modeling of SNC expected distribution (Naomie Herpin-Saunier et al. 2022)

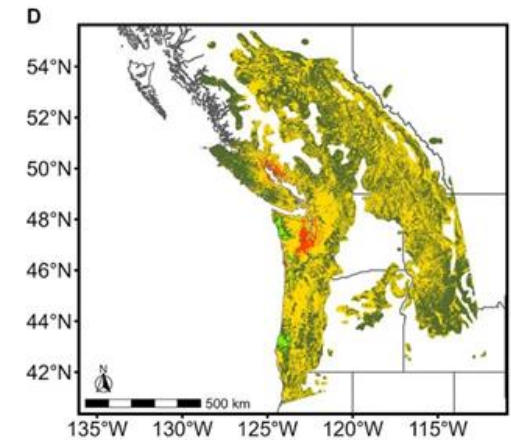
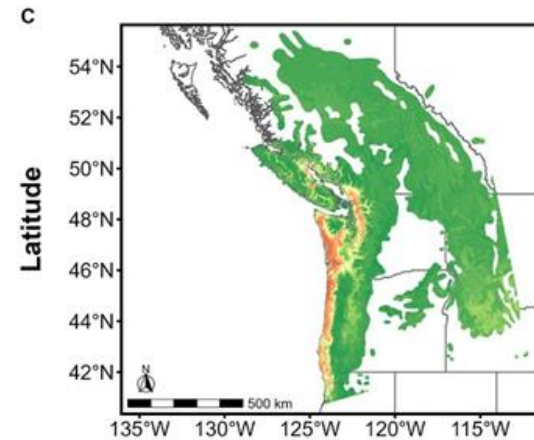
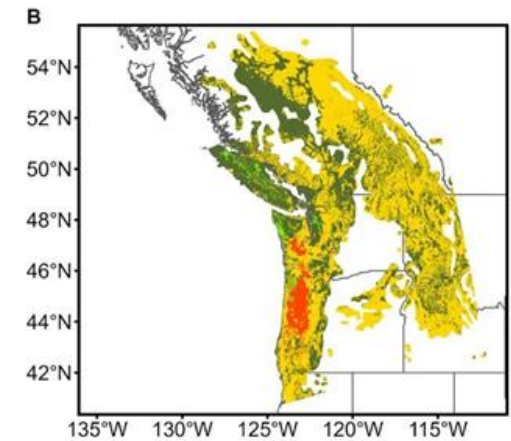
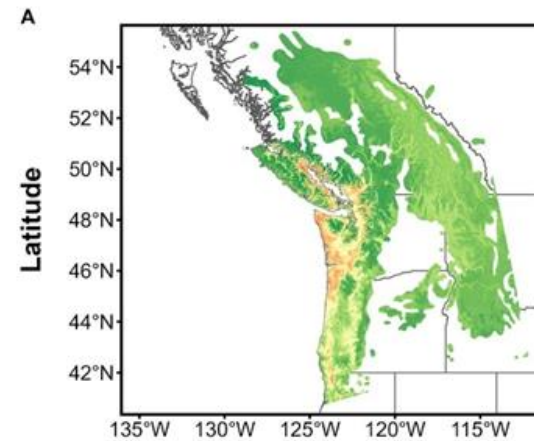
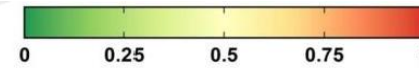
- Environmental tolerance range of Lin. 1c exceeds that of Lin. 2
- Lin. 1c favored by hotter and drier summers
- Lin. 2 has less tolerance to heat

Lineage 1c

Lineage 2

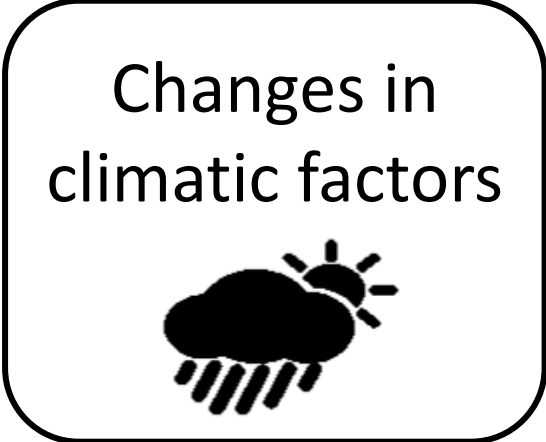
Current probability of occurrence

Probability of change (2050 vs. current)



Longitude

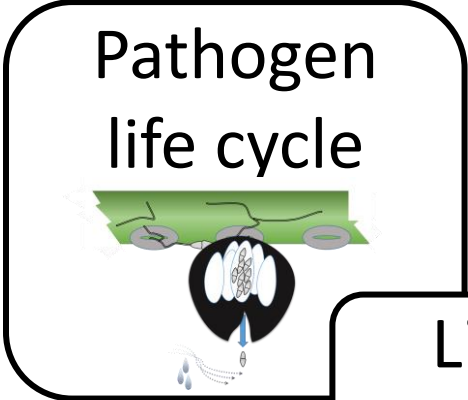
Longitude



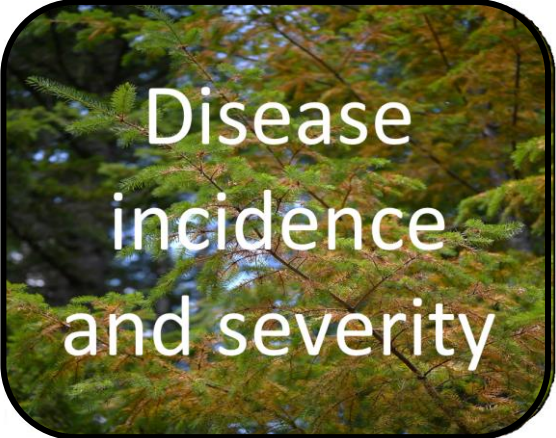
INDUCE

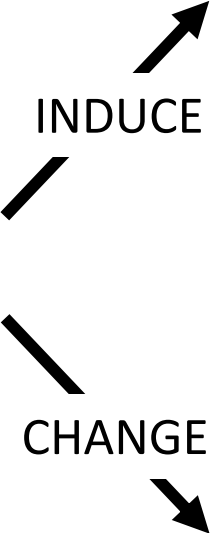
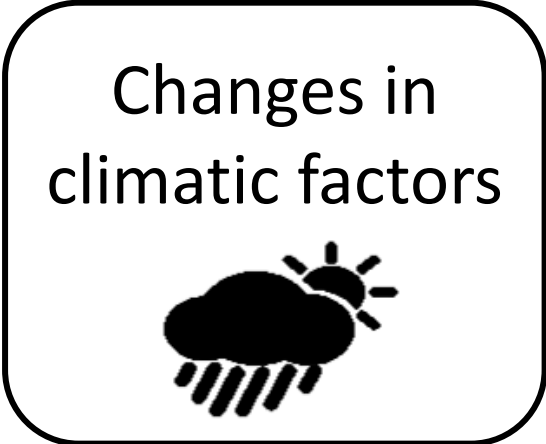


CHANGE



CHANGE



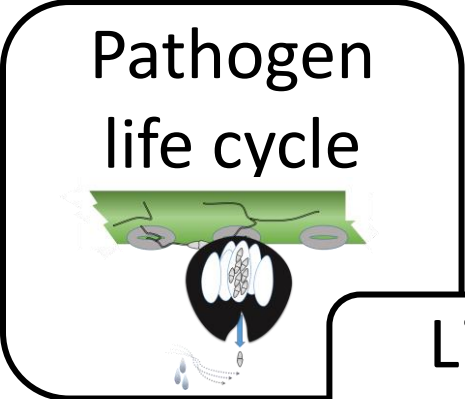
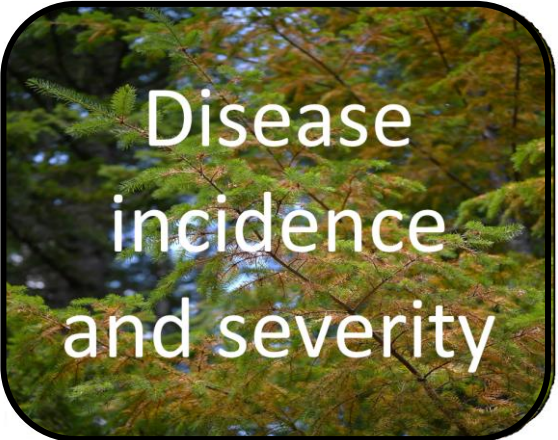
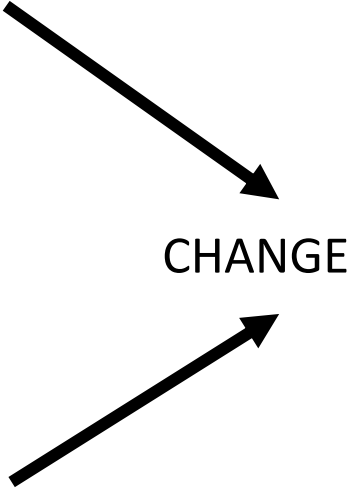


Drought on Df
Drought x SNC ?



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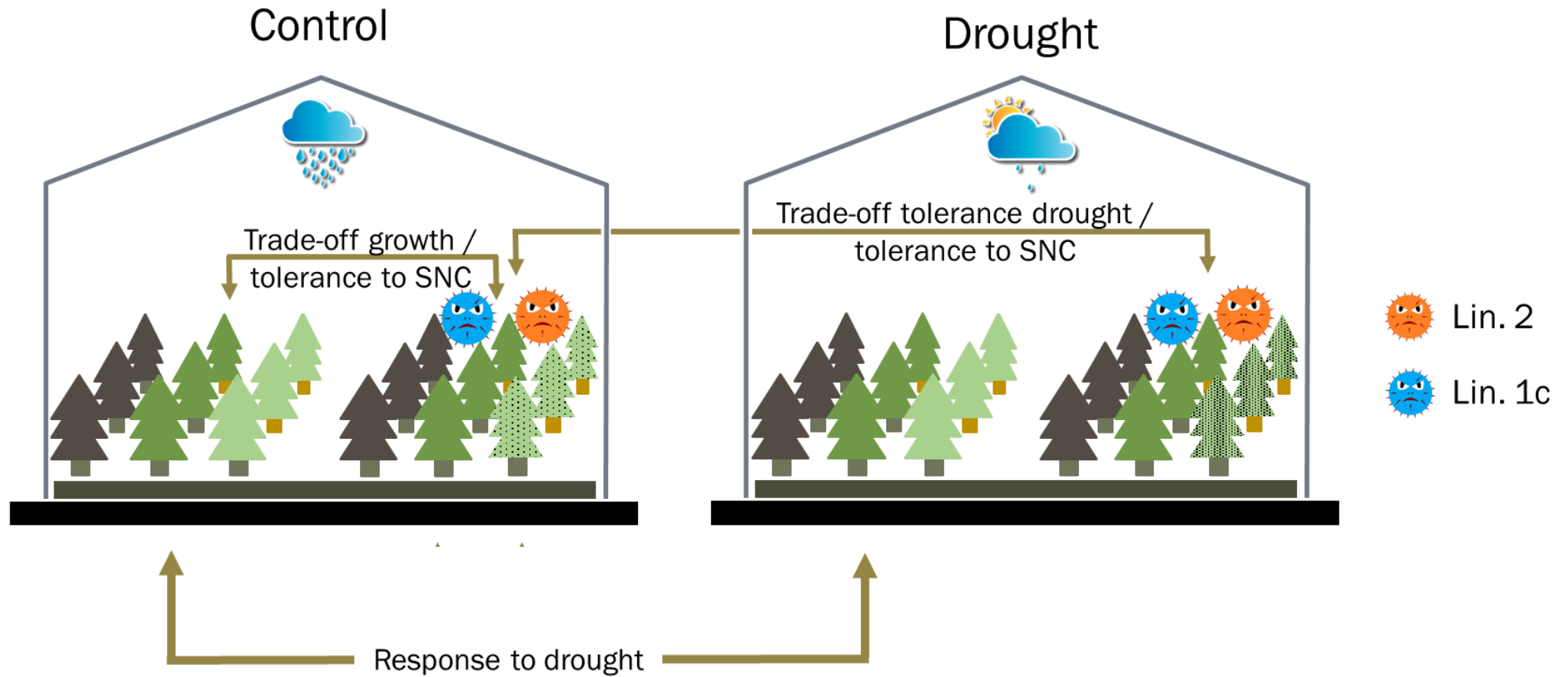
Lineages distribution

CoAdapTree



Healthy Trees for Future Climates

Studying Df x SNC x Drought

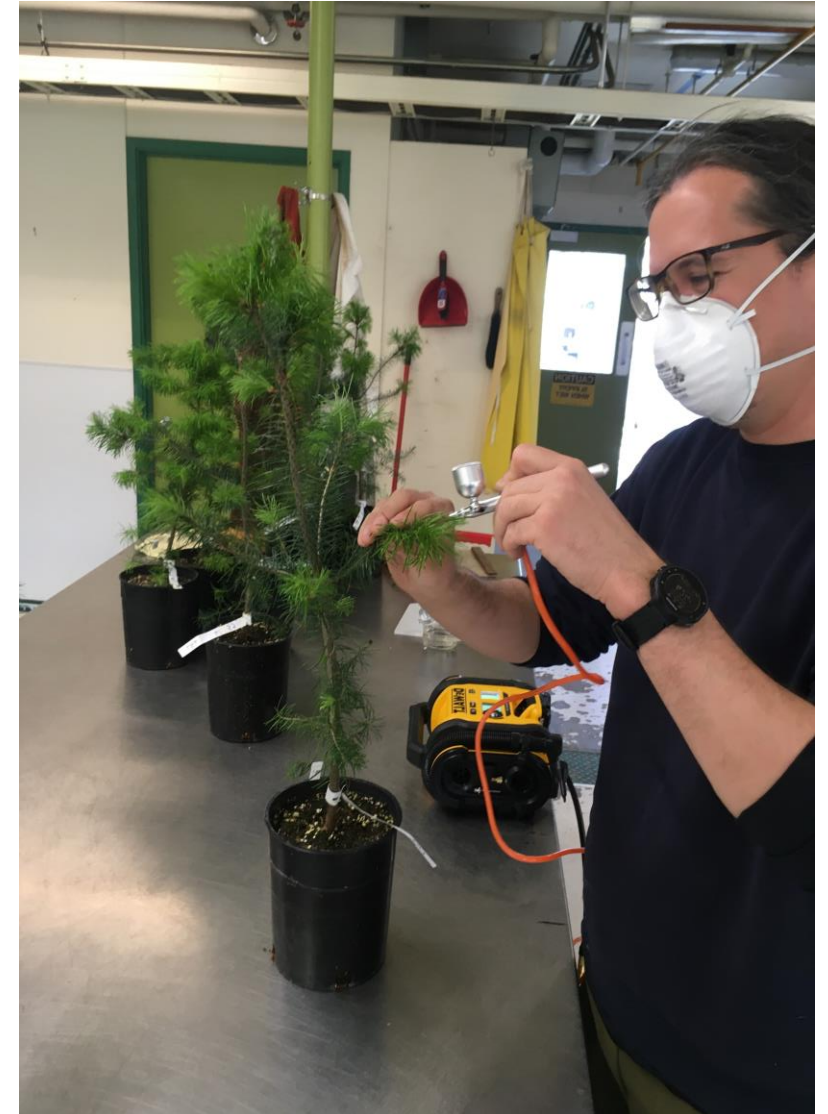
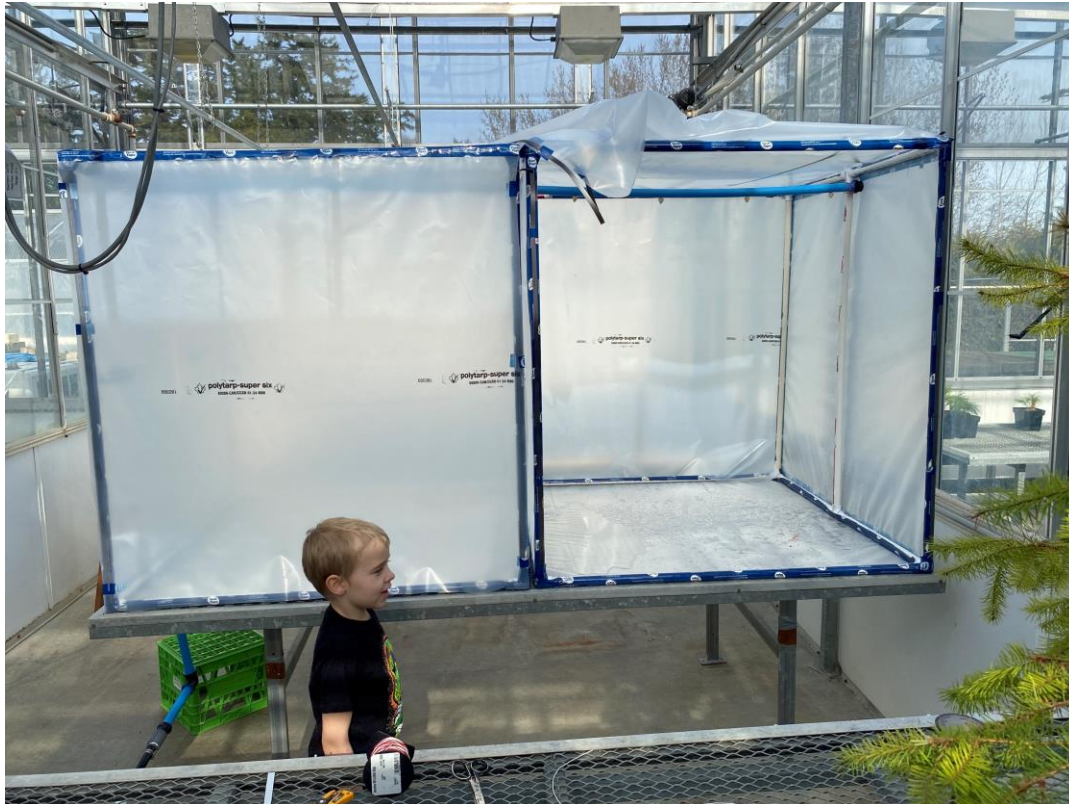


Phenotyping (growth & SNC severity) 4 Douglas-fir families from BC-MoF

Transcriptomics (gene expression)

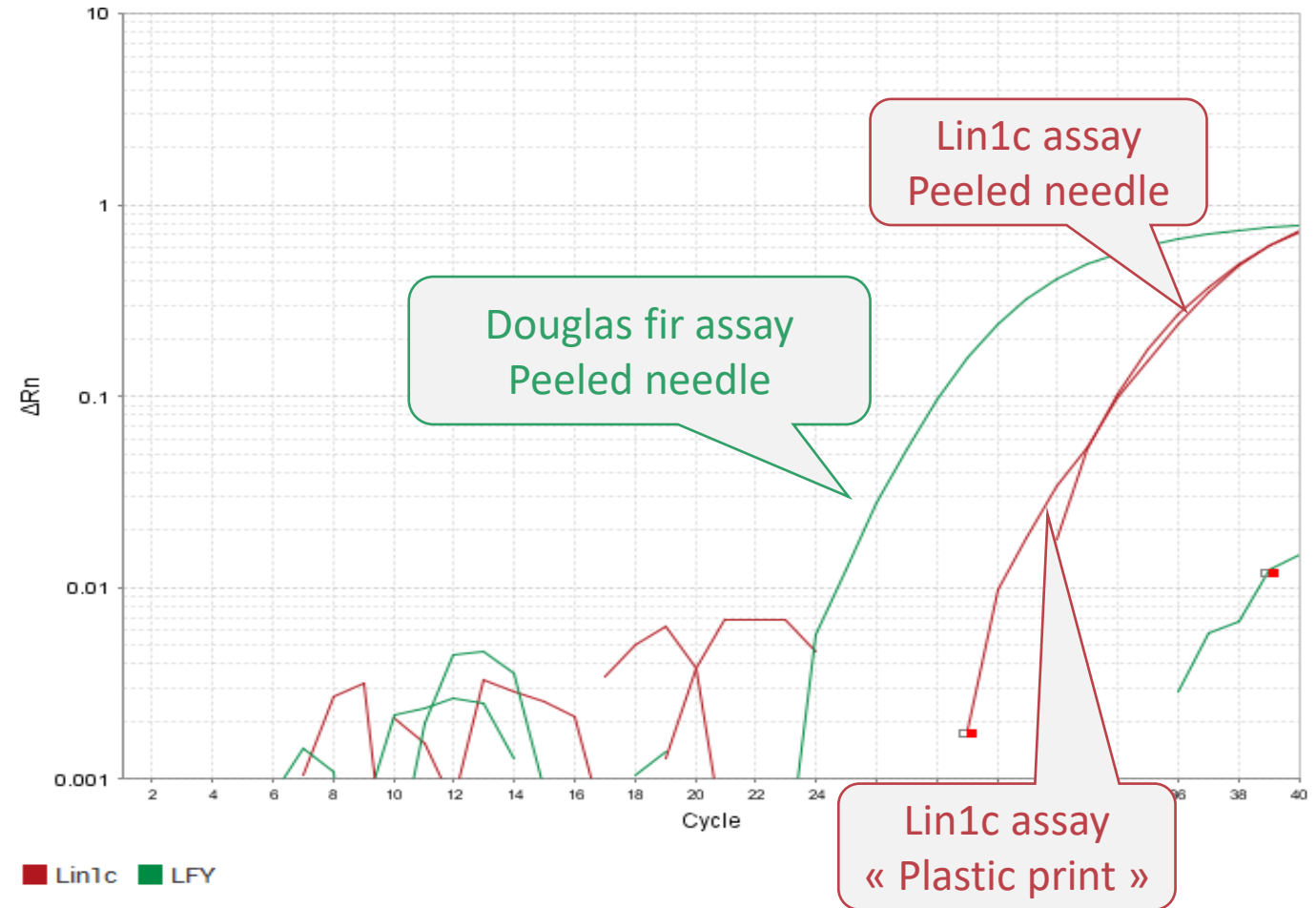
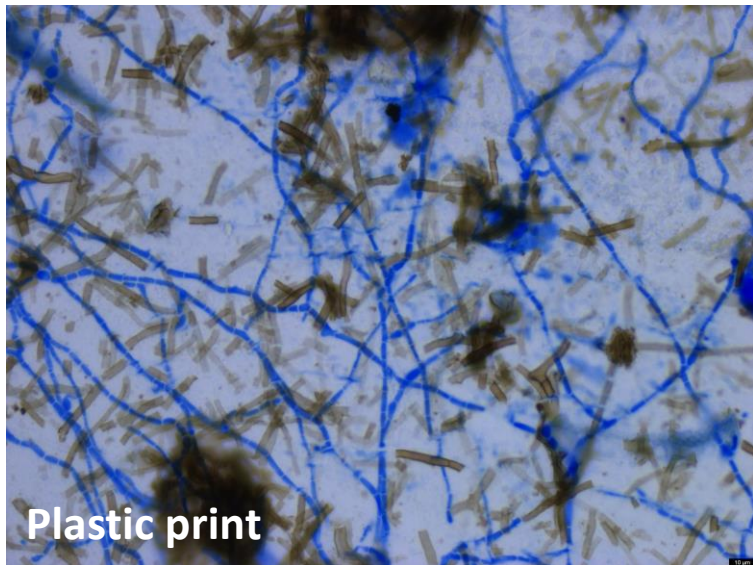
Controlling SNC infections in the lab: Mission impossible?

N. gaeumannii does not sporulate in Petri dish
Sprayed fragmented mycelium (20gr/L);
Mist chambers 17-21°C, >80% humidity



Mission possible!

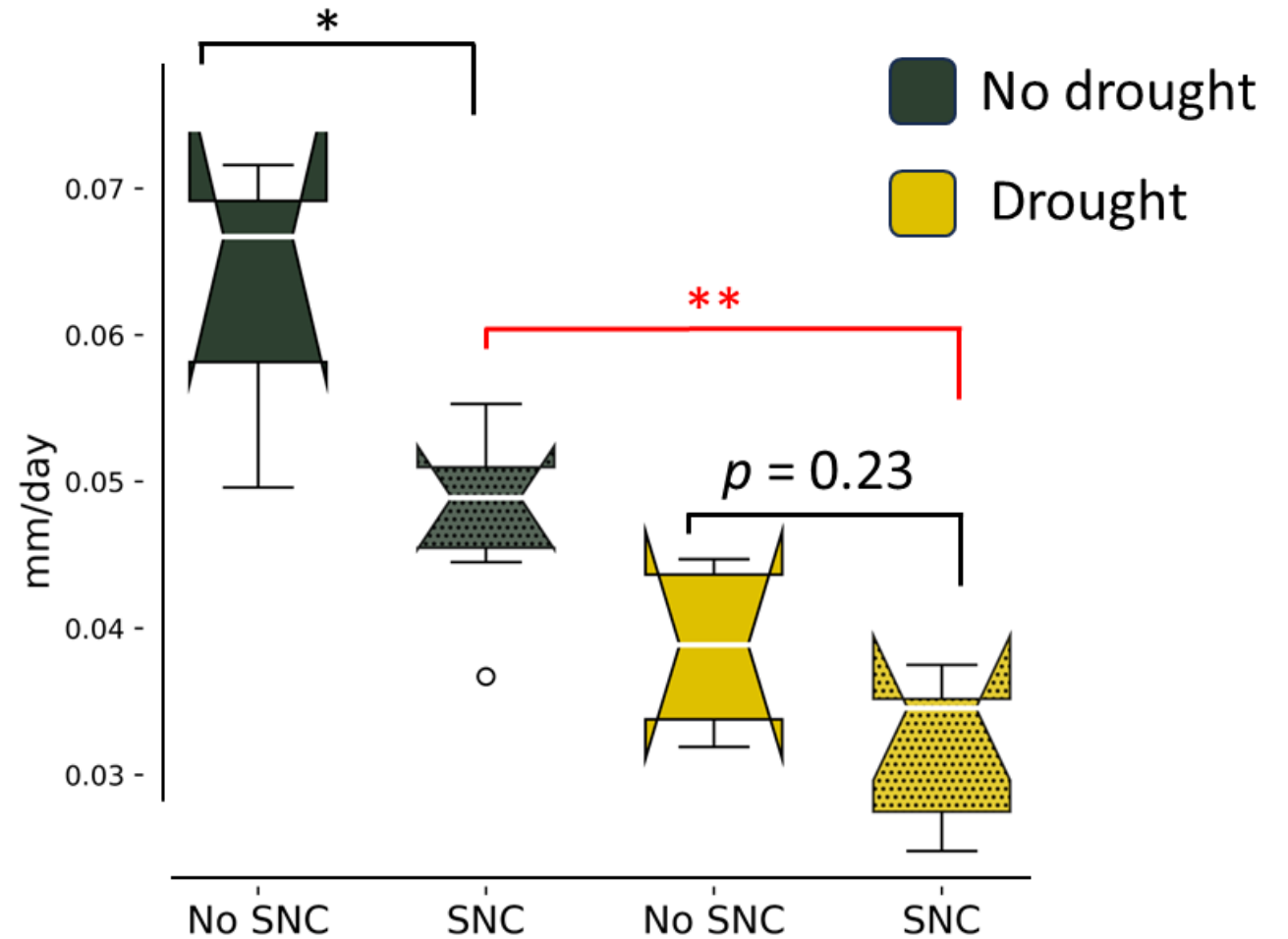
3 weeks post inoculation :



qPCR detection confirmed at 6-months post-inoculation

Six months after the first inoculation :

- Drought impacts tree growth (we knew that)
- **SNC impacts diameter growth**
- **Trend visible (and significant) in 3 families (out of 4 tested)**
- SNC effect not visible on tree length



Diameter growth in Df family 81

Acknowledgments

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CoAdapTree



Sally Aitken
Sam Yeahman
Richard Hamelin



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