

Community ecology of foliar fungi and oomycetes of *Pseudotsuga menziesii*

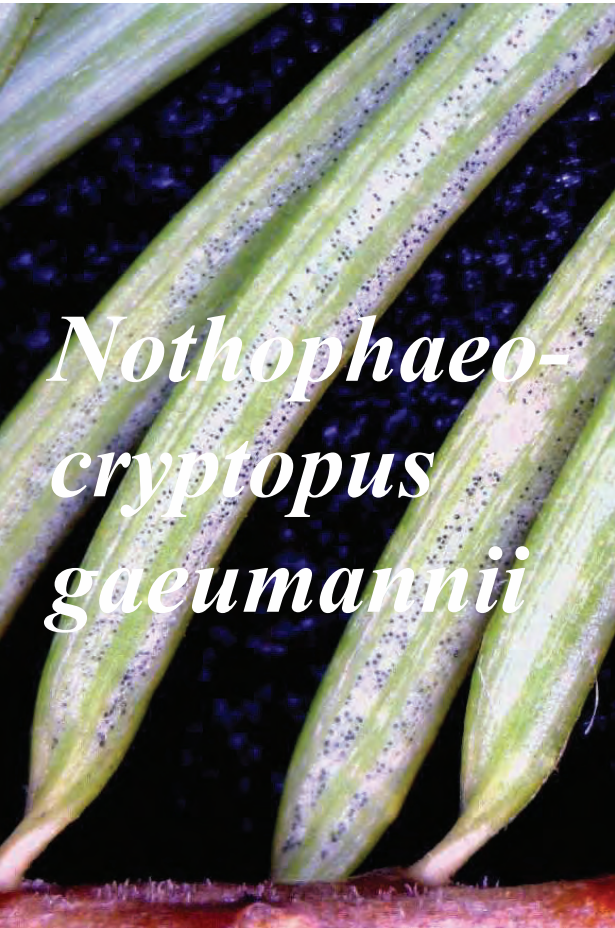
Lilah Gonen

MS Defense

Dec 9, 2020

Advised by Jared
LeBoldus and Andy Jones

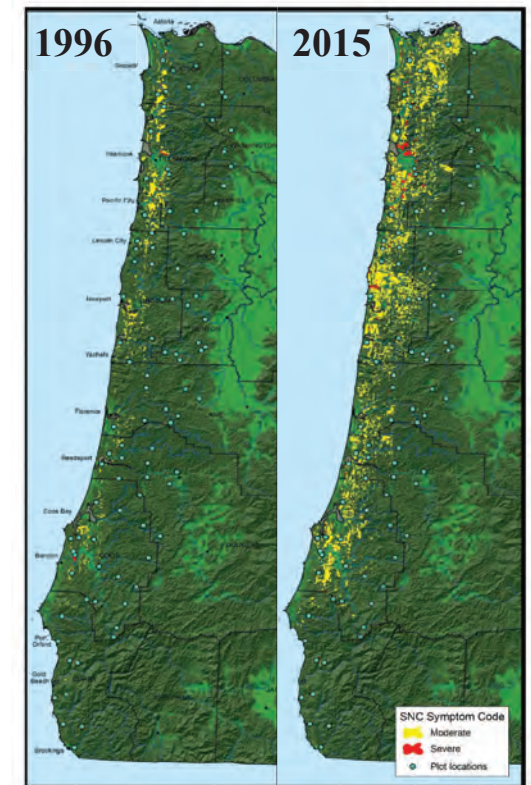




*Nothophaeo-
cryptopus
gaeumannii*

Fungal endophyte/latent pathogen

- Ubiquitous, obligate, specific to *Pseudotsuga*
- Causal agent of Swiss needle cast (SNC)
 - Climate change (Manter et al. 2005)
 - Management (Stone et al. 2008)
 - Climate transfer distance (offsite planting) (Wilhelmi et al. 2017)



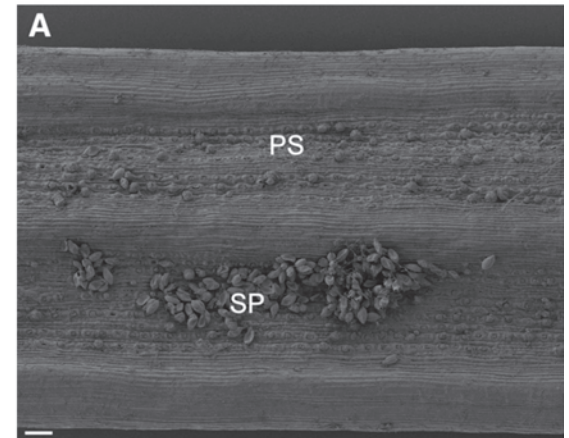
Adapted from Ritóková et al. 2016



Reeser et al.
2013

Only foliar oomycete described in *P. menziesii*

- Cryptic symptoms
- Episodic, geographically isolated needle casting events
- Inversely abundant to *N. gaeumannii* (Gómez-Gallego et al. 2019)



Gómez-Gallego et al. 2019

Objectives

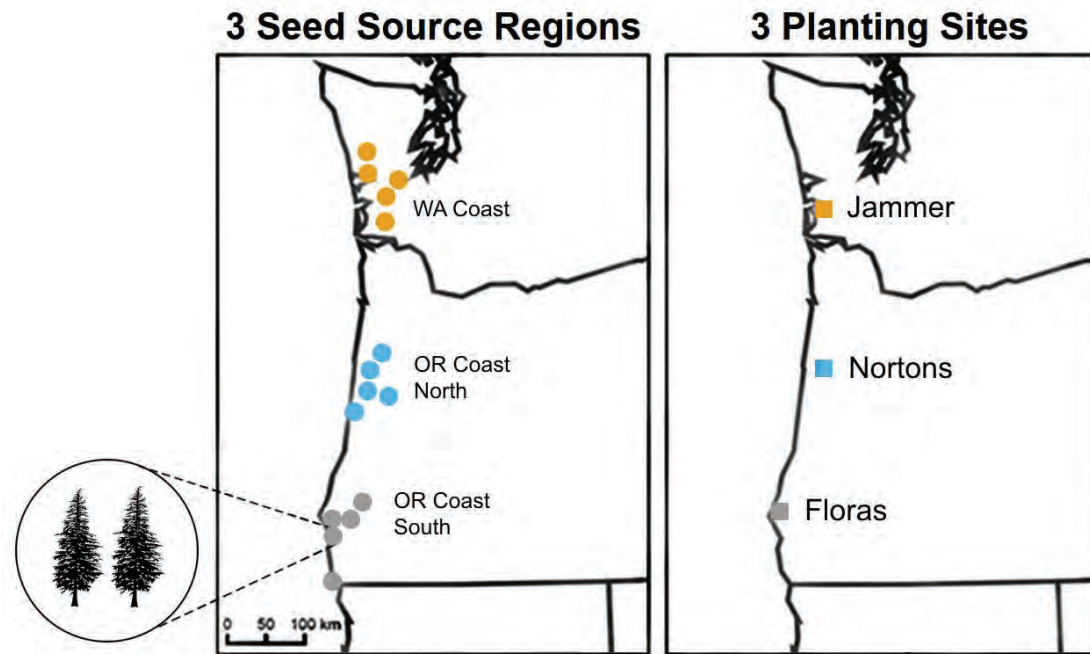
- 1) Describe community dynamics of foliar fungi and oomycetes of *P. menziesii* on the PNW coast
- 2) Characterize microbial communities in response to effects of host location and genetic lineage
- 3) Model fungal diversity in response to climate transfer distance and oomycete diversity
- 4) Investigate associations between *N. gaeumannii* and oomycetes

Methods

- **Sample needles from trees in reciprocal transplant provenance study (SSMT)**
- **Culture**
 - Isolate and sequence oomycetes from needles
- **Next-generation sequencing (NGS)**
 - Sequence all fungal and oomycete DNA extracted from needles
- **Statistical analyses**
 - Analyses of community dissimilarity
 - Generalized linear mixed modeling (GLMM)

Study Sites

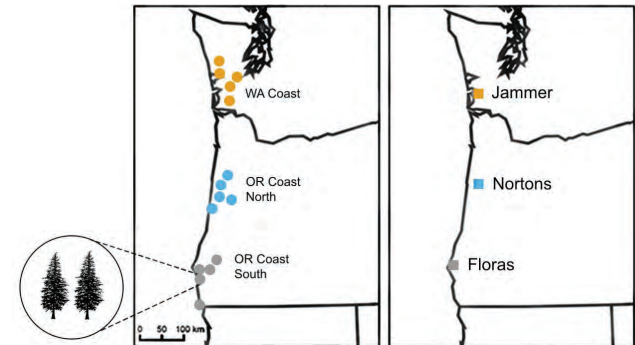
USFS Douglas-Fir Seed Source Movement Trial (SSMT)



*Adapted from Wilhelmi et al.
2017*

Study Sites

Block 4		Block 3	
43	42	31	30
44	41	32	29
45 WACST	40 ORCSTS	33 WACST	28 ORCSTS
46	39 ORCSTN	34 ORCSTN	27
47	38	35	26
48	37	36	25
19	18	7	6
20 WACST	17	8	5
21	16	9 ORCSTS	4 WACST
22 ORCSTN	15	10	3 ORCSTN
23 ORCSTS	14	11	2
24	13	12	1
Block 2		Block 1	



7201	8027	7202	8028
8027	8026	7205	7206
7201	8542	8028	8541
8026	7206	7205	8025

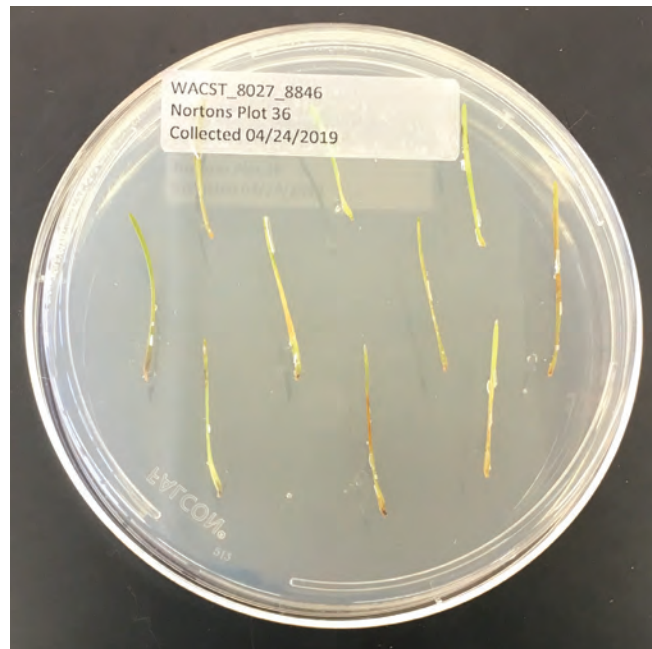
n =
338

Sample Collection



Sample Collection

Plate 10 needles



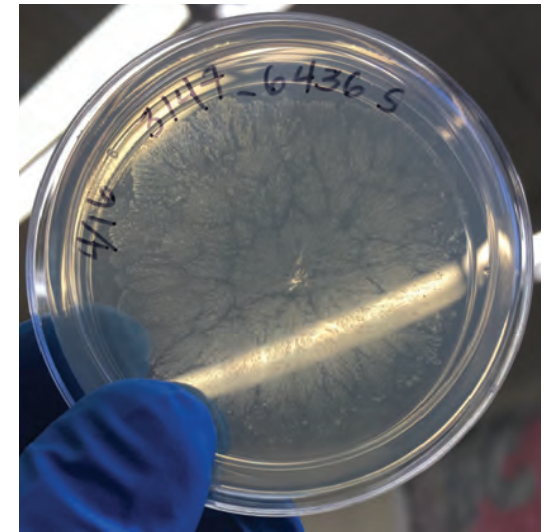
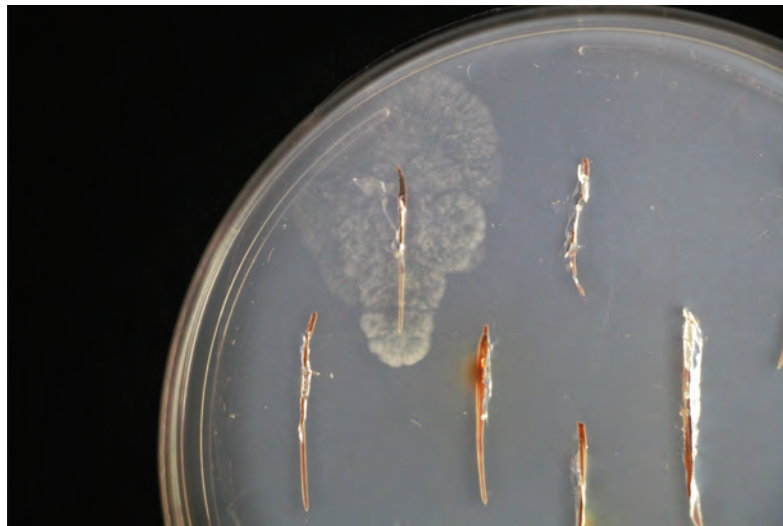
Sequence 10 needles



Culture

18 isolates

Nortons: 16 Jammer: 2 Floras:
0



Culture

12 isolates

Nortons: 11 Jammer: 1 Floras:
0



ITS

- ITS5 & ITS4 (White et al. 1990)

COI

- OomCoxI-Levup & OomCoxI-Levlo (Robideau et al. 2011)

Rps10

- Prv9r-M & Prv9f-M (Foster 2020)

NGS

Library prep and sequencing

- **2-stage PCR**
 - Fungi: *ITS2*
 - Oomycetes: *Rps10*
- **Illumina MiSeq 3000**
 - 300-bp paired end reads
 - Fungi and oomycetes sequenced on same lane



Illumina
a

NGS

Fungi

- n = 246
- OTUs = 105

Oomycetes

- n = 181
- OTUs = 55

All microbes

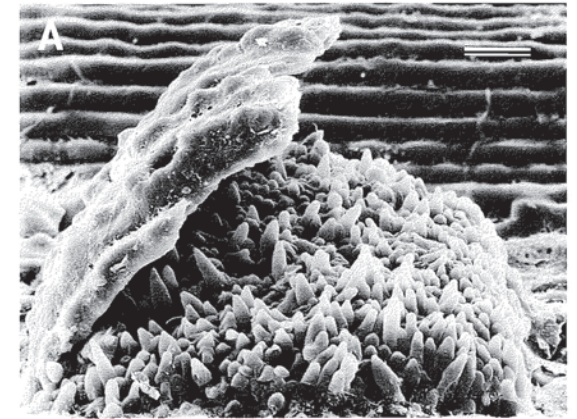
- n = 282
- OTUs = 157
- Rarefied to 1000 min reads

	OTU.1	OTU.2	OTU.3	OTU.5
F12-3180-1757.f	4.009315	2.432123	3.163402	2.50086
F12-6059-2776.f	3.902164	3.292552	3.396323	0.00000
F12-6059-7502.f	3.965152	3.228200	2.447291	0.00000
F12-8552-4829.f	4.004358	0.000000	3.008734	0.00000
F12-8552-9359.f	3.780196	3.568014	2.988478	0.00000
F14-3147-1661.f	3.889971	0.000000	2.458972	0.00000

1) Describe community dynamics of foliar fungi and oomycetes of *P. menziesii* on the PNW coast

Fungi (105

OTU(s) OTU	Taxon
OTU 1	<i>Rhabdocline parkeri</i>
OTU 2	<i>Rhizosphaera</i> sp.
OTU 3	<i>Nothophaeocryptopus gaeumannii</i>
OTU 4	<i>Rhizosphaera</i> sp.
OTU 5	<i>Curvibasidium cygneicollum</i>
OTU 7	<i>Cladosporium</i> sp.
OTU 8	<i>Penicillium penicilliodes</i>
OTU 9	<i>Hormonema macrosporum</i>
OTU 10	<i>Curvibasidium cygneicollum</i>
OTU 12	Didymellaceae



Sherwood-Pike et al. 1986



*OSU SNC
Cooperative*

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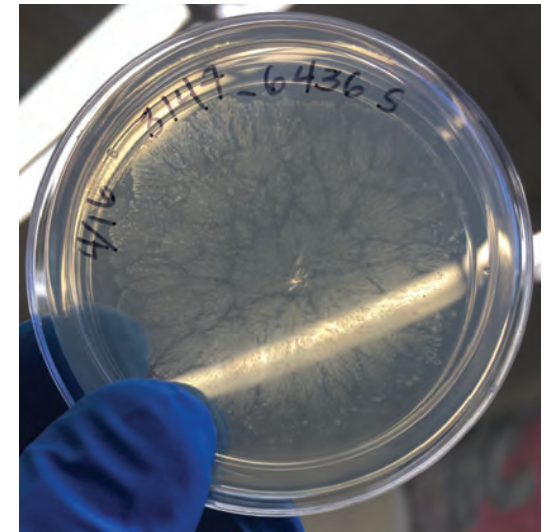
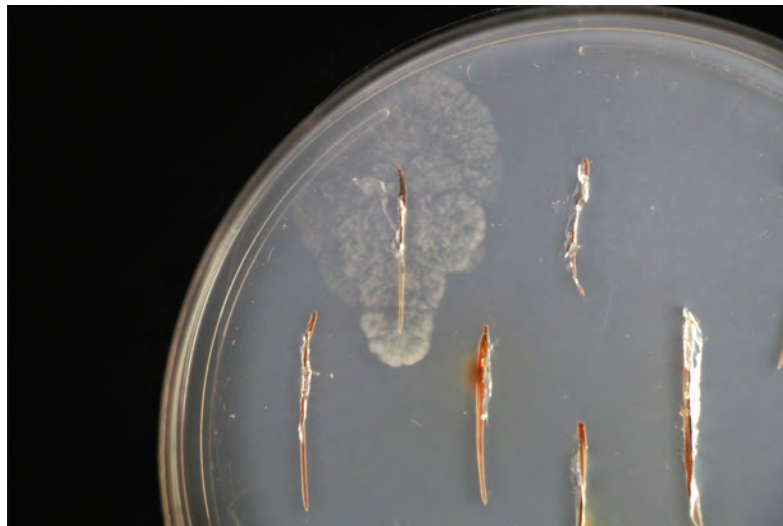
Oomycetes (55 OTUs)

OTU	Taxon
OTU 1	<i>Phytophthora cacuminis</i>
OTU 4	Pythiaceae
OTU 7	Oomycete
OTU 8	Oomycete
OTU 11	Oomycete
OTU 13	Oomycete
OTU 16	Oomycete
OTU 18	Oomycete
OTU 22	Pythiaceae
OTU 26	Oomycete

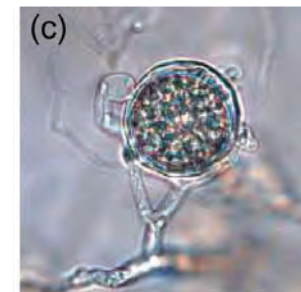
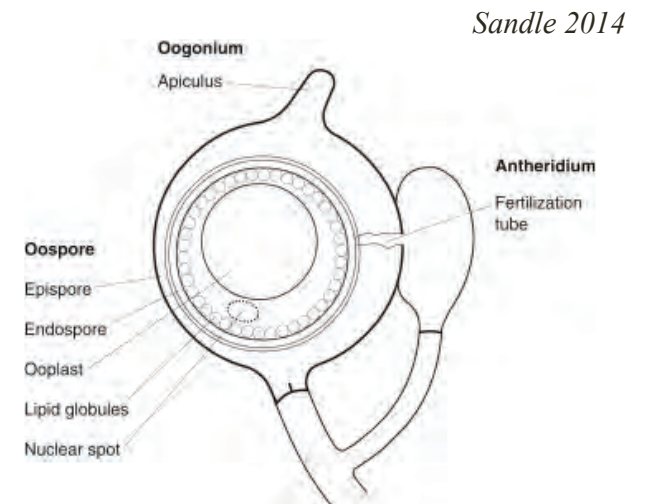
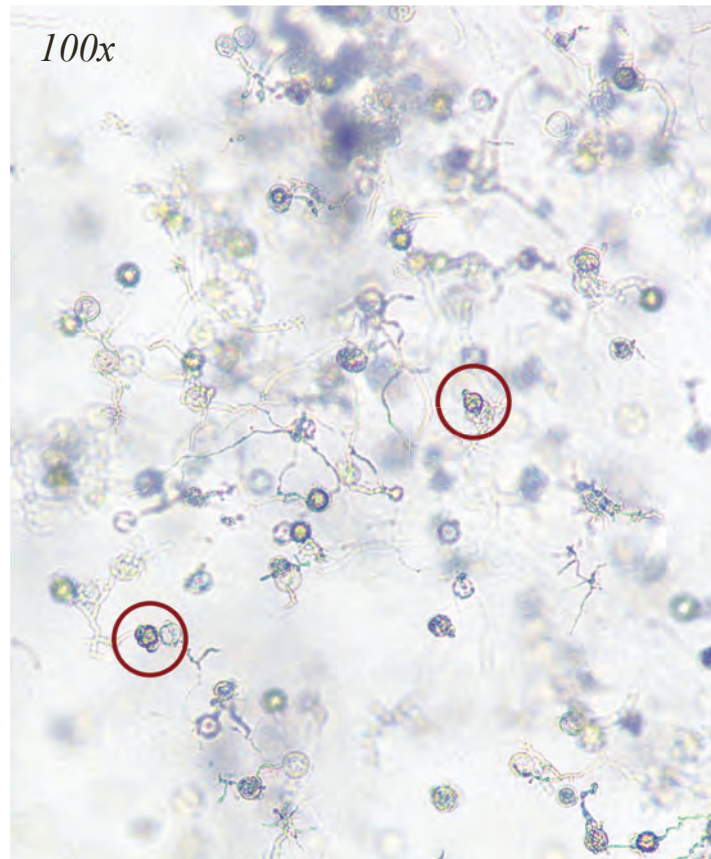
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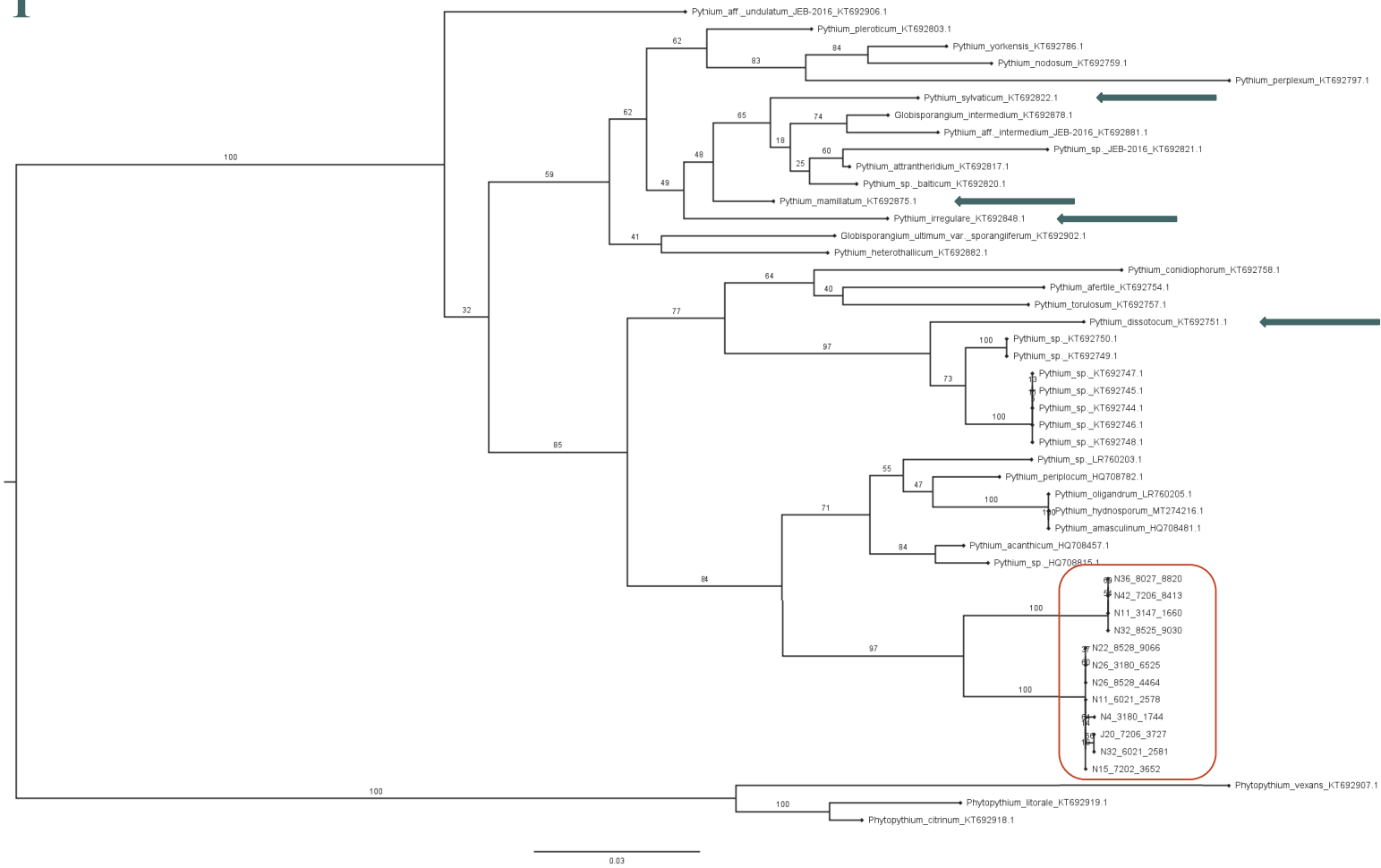


Rahman et al. 2015



de Cock et al. 2015

CO I



IT
S



Rps10



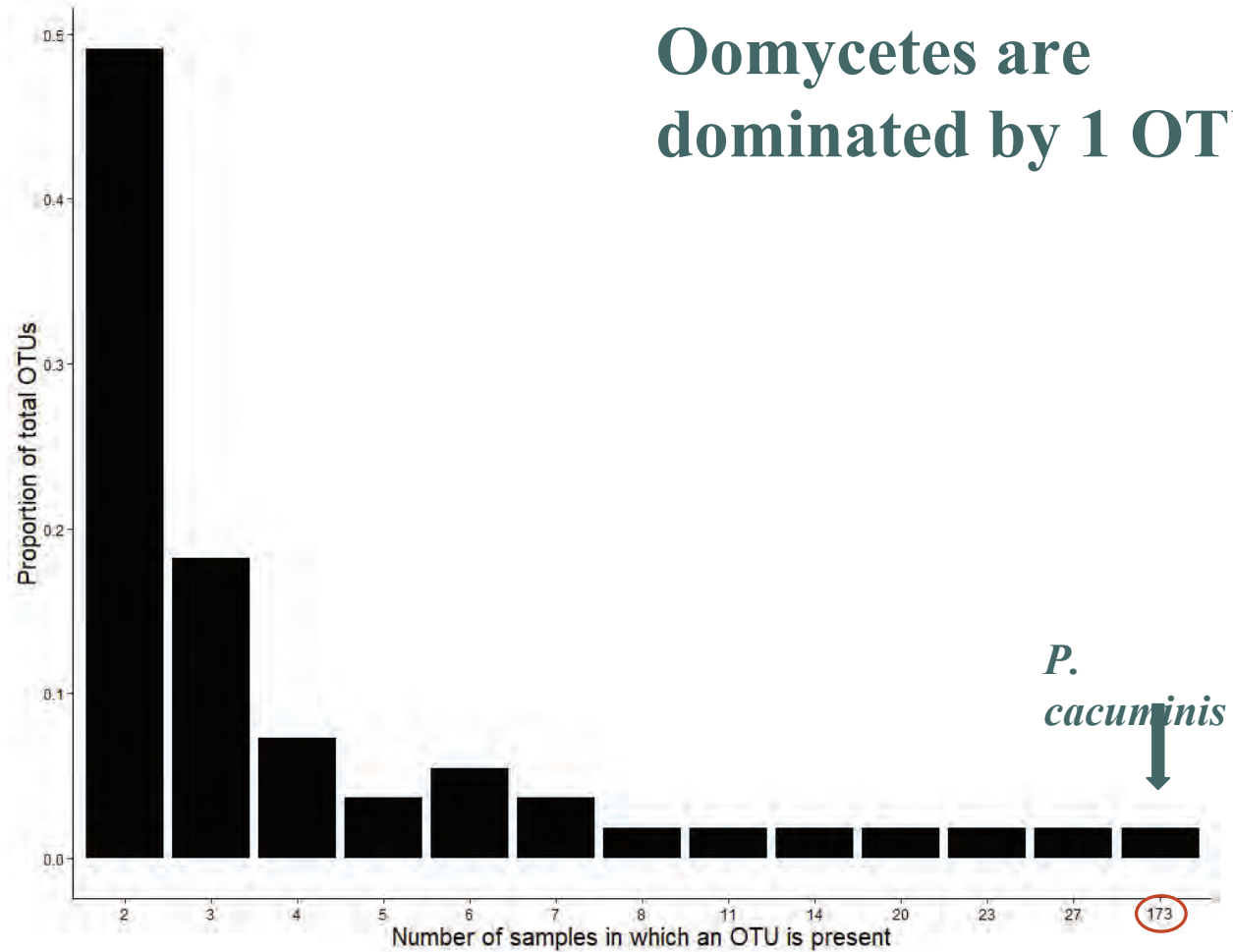
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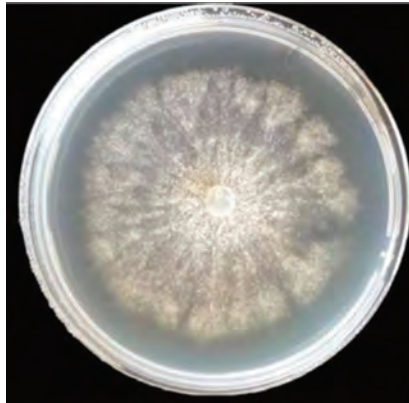
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Oomycetes are dominated by 1 OTU



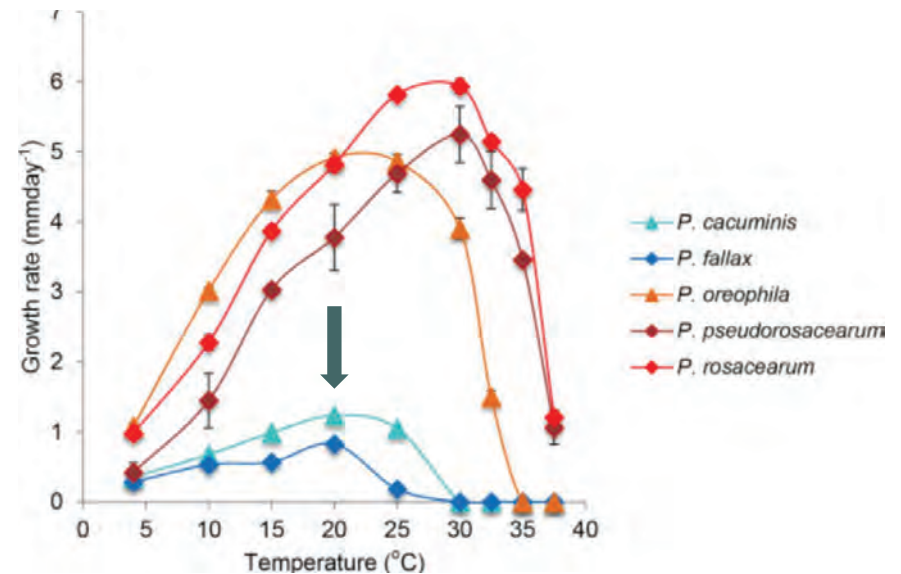
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Phytophthora cacuminis



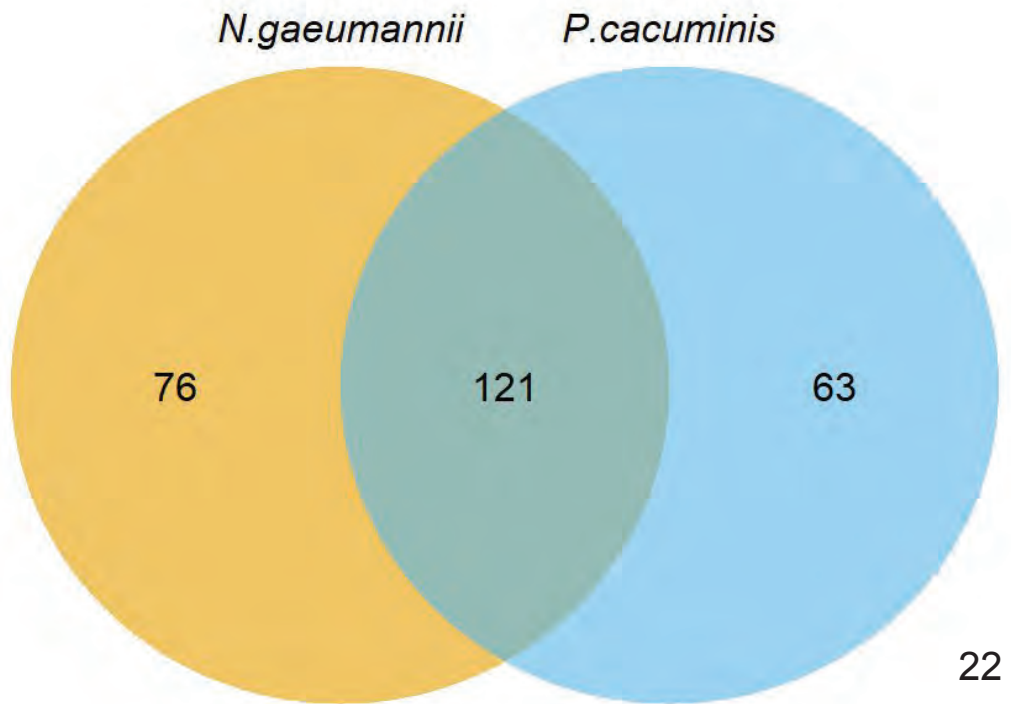
Functional traits suggest adaptation to cold environments (Redondo et al. 2018)

Isolated from asymptomatic *Eucalyptus* root tissue in alpine environments in **Australia** (Khaliq et al. 2019)



Figures from Khaliq et al. 2019

4) Investigate any associations between *N. gaeumannii* and oomycetes



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Logistic regression (log link)

Response: *N. gaeumannii* presence/absence

Fixed effect: *P. cacuminis* presence/absence

Random effects: Site, Block(Site)

Term	Estimate	Std Error	z value	<i>p</i>
Intercept	1.393	0.458	3.047	0.002
<i>P. cacuminis</i>	-0.654	0.309	-2.117	0.03

P. cacuminis present: $\pi_1 = \mathbf{0.677}$

P. cacuminis absent: $\pi_0 = \mathbf{0.801}$

Thank you

Committee

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