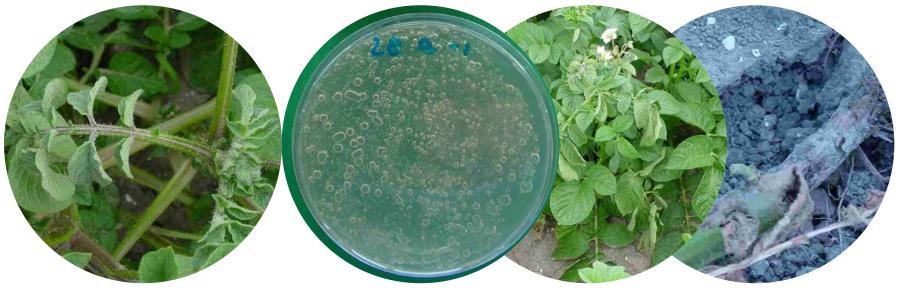
Use of disease suppressiveness in the battle against potato blackleg

EAPR Pathology & Pest symposium, Neuchatel, Switzerland, 1–5 September 2019

Jan van der Wolf







# Background study

- Bacterial diseases caused by Dickeya and Pectobacterium species are still the most important problem in seed potato cultivation
- There are no pesticides available to control the disease
- There is no (full) resistance present in commercial potato varieties
- Infections are hard to prevent
- Disease expression is difficult to predict

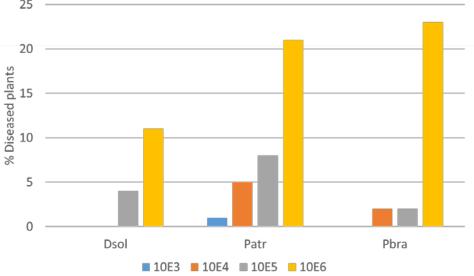








# Factors determining expression of potato blackleg



- Potato cultivar
- Pathogen density
- Pathogen variant
- Environmental conditions (temperature, humidity)
- Suppressiveness seed tubers??

(Ability to prevent disease expression)







## Suppressiveness in seed lots

#### Research questions

- Does suppressiveness against soft rot *Pectobacteriaceae* (SRP) exist in seed lots?
- If yes, what is the basis?
  - Physical-chemical properties?
    - Dry matter, moisture content?
    - (micro)nutrients?
  - Metabolites?
  - Microbiome?









#### Set up experiment

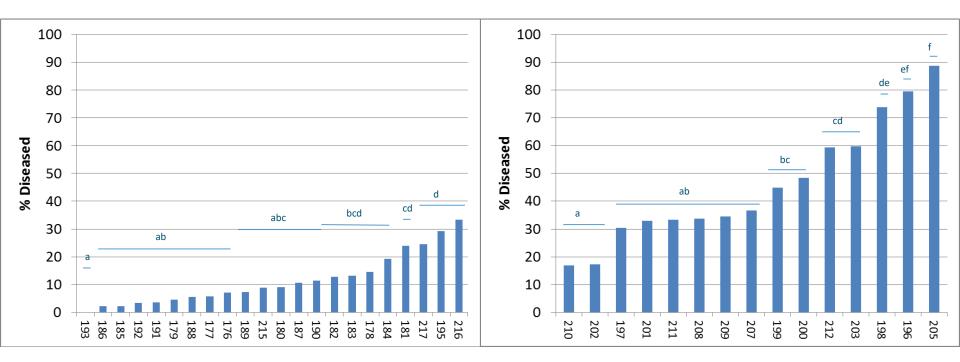
- Seed lots of two cultivars (ca. 20 per cultivar per year) collected from different locations and growers in the NL
  - Tested for a low level of SRP's
  - Analysed for dry weight content and minerals
  - Inoculated with 10<sup>6</sup> cfu/ml of *D. solani* or not inoculated
  - 100 tuber per treatment were planted in a (randomized block design) in a sandy soil in the North of the NL







#### % diseased plants



Cultivar Spunta

Cultivar Kondor







#### Does suppressiveness against SRP exists?

#### Seed lots

- of same cultivar (same genetic background)
- with the same infection level (*Dickeya solani*, 10E6 cells/ml)
- planted in the same field (same environmental conditions)

can remarkably differ in disease incidence!

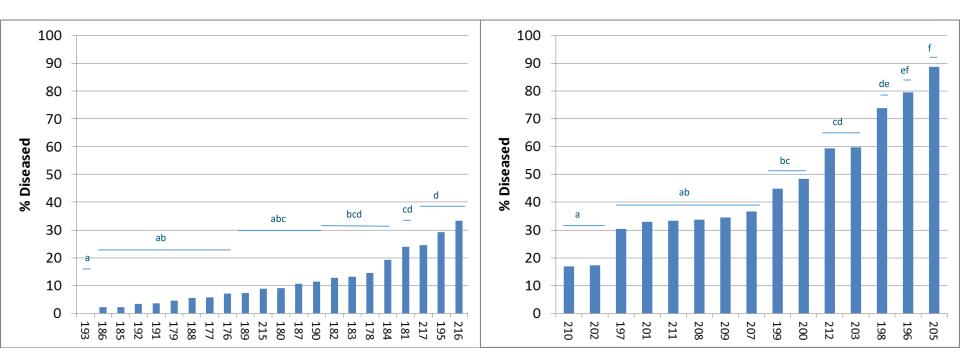
#### Yes, suppressiveness against SRP exists







# Consequences for management and research (including breeding)



**Cultivar Spunta** 

Cultivar Kondor

#### Use a minimum number of seed lots in your studies

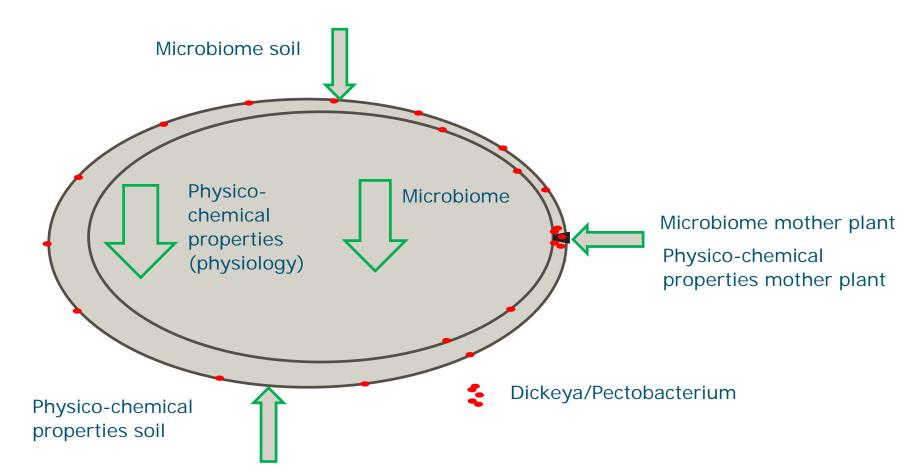






## Background suppressiveness





Microbiome = the micro-organisms living in a particular environment WAGENINGEN



#### Role of minerals/dry weight

Infection prevalence ~ Year + Cultivar + DS + Na + Ca + P + Mn + Cu + B + N Intercept = KONDOR in 2015, when all minerals and dry weight have average values

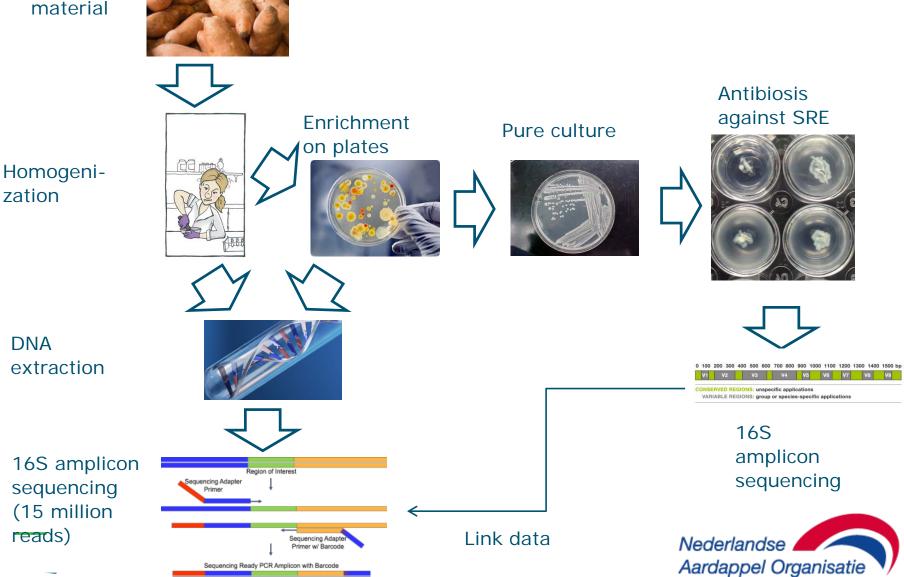
term	p.value		prevalence	net_effects
(Intercept)	0.000196	***	39	0
Year2016	0.000119	***	28	-11
CultivarSPUNTA	5.80E-58	***	12	-27
DS	7.76E-07	***	36	-3
Na	3.20E-07	***	40	1
Ca	2.78E-06	***	42	3
Ρ	0.0063	**	38	-1
Mn	0.074895	-	40	1
Cu	0.063117	-	38	-1
В	0.003335	**	37	-2
Ν	9.33E-13	***	43	4



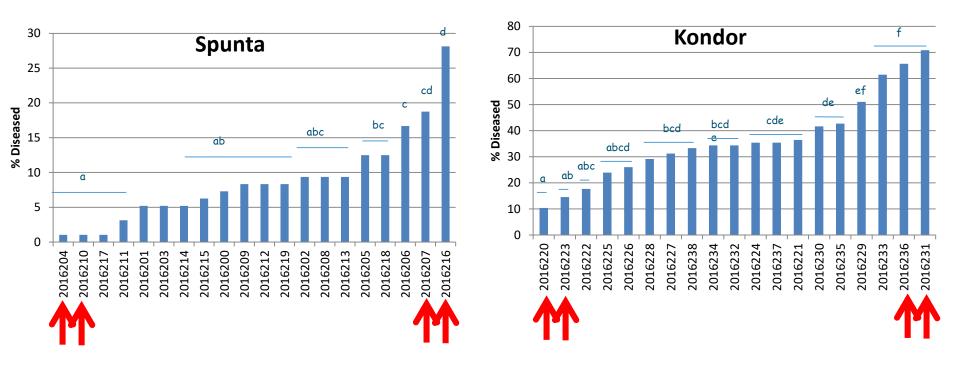
Nederlandse Aardappel Organisatie

#### Microbiome analysis

Sampling plant material



# Selection of seed lots for microbiome analysis

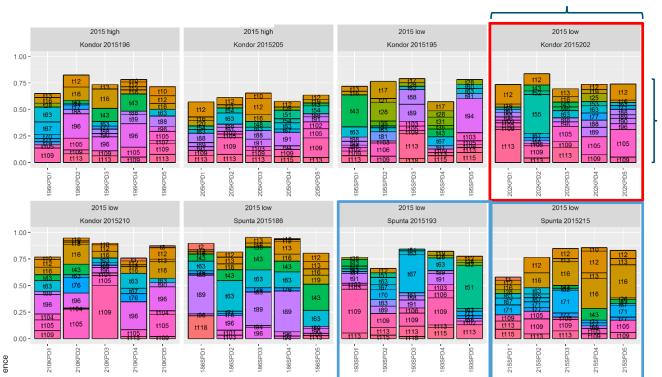








## Results microbiome analysis



#### 5 tubers/lot

10 dominant taxonomical groups (OTU's)

Large variation in microbiome between tubers within a lot Significant differences in microbiome between lots

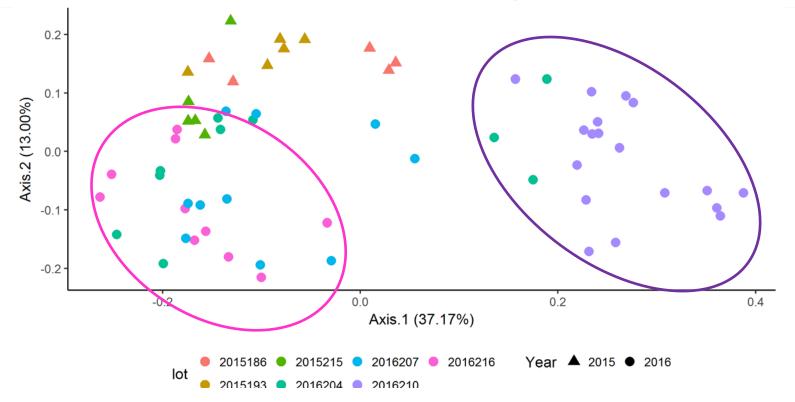




#### PCoA for microbiome periderm cv. Kondor



PCoA wUniFrac otu Hellinger

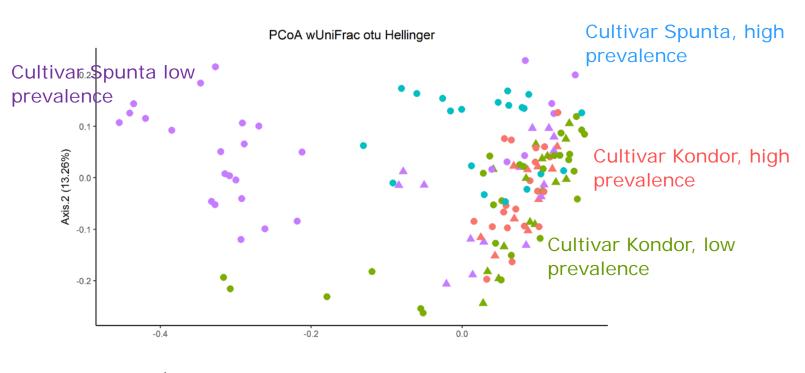


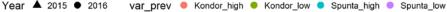
#### Significant differences in microbiome between lots





#### PCoA for periderm of all seed lots (preference groups)

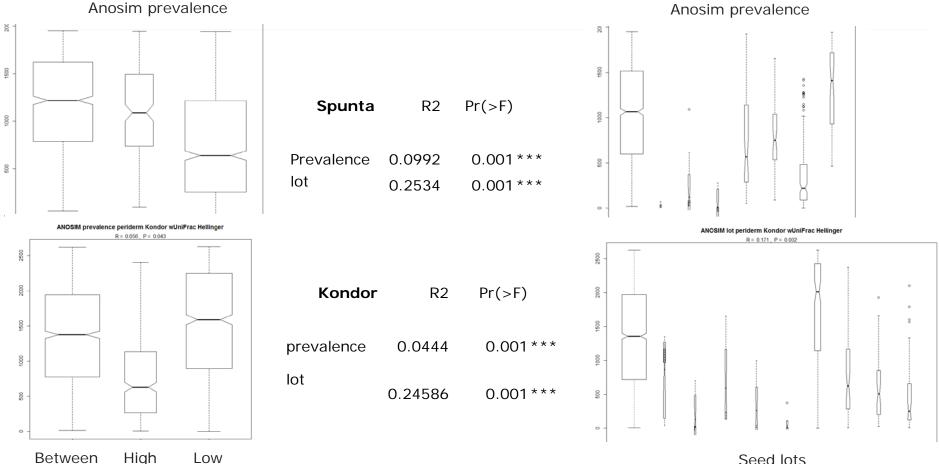








## Proportion of variance explained by the prevalence and lot – per variety



Seed lots

Nederlandse

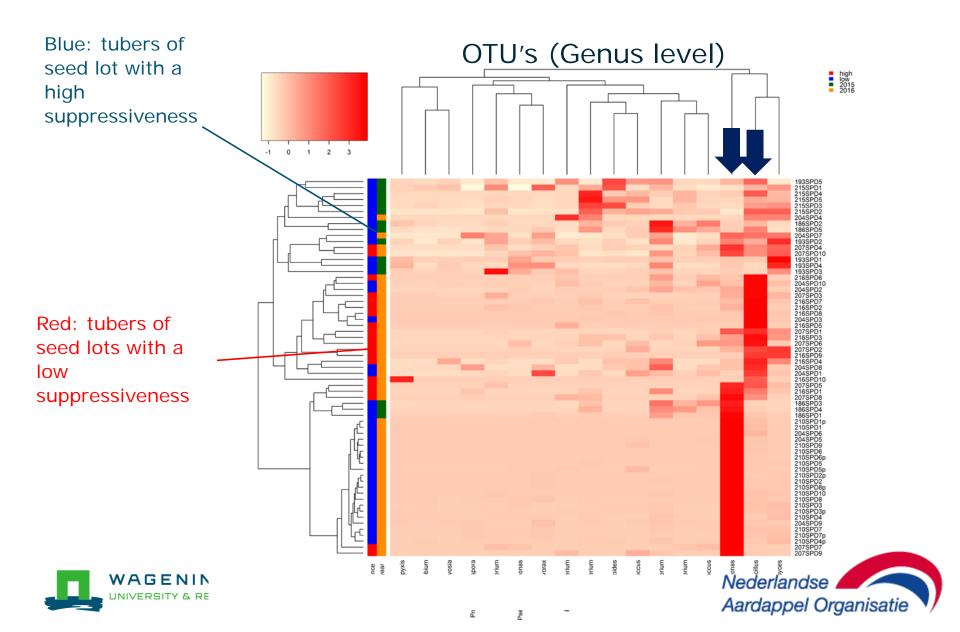
Aardappel Organisatie

The microbiome of the tuber tissue is to some extent correlated with the infection prevalence The microbiome differ largely per seed lot





#### Indicators suppressiveness cv. Spunta



#### How to use the information



- Identification of indicators for suppressiveness
- Selection of seed lots with a high suppressiveness
- Improvement of suppressiveness
  - Soil amendments
  - Tuber treatments
    - Biocontrol agents (selection via microbiome analysis)







# Follow up researchGenomeTranscriptome $\downarrow$ <

- Selection of isolated bacteria related to suppressiveness
- Characterization of bacteria
  - To determine risk group (based on 16S rDNA sequences)
  - For their ability to inhibit SRP's in planta
- Evaluation in field experiments
  - Individual strains versus communities
  - Ability to reduce blackleg







Proteome

https://www.researchgate.net/figure/The-metabolome..

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