



Jutta Ludwig-Müller and Susann Auer Clubroot Steering Committee Meeting, 30 April 2020

Biocontrol potential for clubroot by Acremonium *alternatum* – chances and challenges



The defense compound salicylic acid can be derivatized by an effector of *P. brassicae*





Ludwig-Müller et al., 2015, Mol. Plant Pathol.

PbBSMT is among the highly expressed putative secreted effectors

Schwelm et al., 2015, Scientific Report

Annotation of the most abundant *P. brassicae* proteins of a proteome analysis based on their orthologs

Veronika Malych, Miroslav Berka (Mendel University, Brno)







PbBSMT can methylate salicylic acid, benzoic acid and anthranilic acid



Ludwig-Müller et al., 2015, Mol. Plant Pathol.



Slide 3



Can we use compounds known to induce defense but not methylated by PbBSMT?



Disease Index	Infection rate (%)
83	98
84	100
79	100
79	100
72	93
	Disease Disease Index 83 83 84 79 79 79 79 72 72

no treatment







INA





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Slide 4



Can we use plants with constitutively activated SA-mediated defense?



Lovelock et al, 2016, Mol. Plant Pathol.





Defense can be induced in Arabidopsis by *Acremonium alternatum* via the salicylic acid pathway





Aa = Acremonium alternatum

Pb = Plasmodiophora brassicae



Effect of Acremonium on canola cultivars

Co-inoculation with *Aa* results in a positive effect on Disease Index (DI) and biomass of plants

(medium effect; Cohen´s d_{AaPb vs Pb}= 0.62)

Results for tested *Brassica napus* cultivars



Auer and Ludwig-Müller (2014)







Clubroot suppression in Brassica napus cv. Ability by Acremonium





uninfected small galls large galls





27% reduction in DI after treatment with *Aa* cell wall extract (CWE)

 \Box uninfected \Box small galls \blacksquare large galls





Acremonium increases stem length and yield of "Ability"



Brassica napus cv. Ability, 5 months old







Clubroot resistance is cultivar-dependent!

active genes in canola vary and are largely dependent on parent plants → implications for resistance response!

Cultivars have their own microbiomes each.

high microbial diversity suppresses pathogens → plants can exploit microbial consortia from soil for protection

breeding can hamper possibilities for positive BCA interaction

Assessment of a panel of B. napus varieties by mRNAseq





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What do we think is important for clubroot research in the future

Conditions more close to field tests Field soil testing if only greenhouse trials are possible

Testing of more canola cultivars with more *P. brassicae* isolates should be made

Identification of current clubroot differential sets that are active in soils right now \rightarrow timely testing of isolates

Easier (faster) access to genome/transcriptome data

More collaboration possibilities between remote labs





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BIOCONTROI POTENTIAI FOR CIUDROOT DY ACREMONIUM AITERNATUM – CHANCES AND CHAIIENGES Clubroot Steering Committee meeting, 30 April 2020



