

The role of plant defense proteins during early symbiotic and pathogenic infection in the model legume Medicago truncatula

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HR 晃

Disease infection

Conclusion and Outlook

the role of Rac1 protein in plant growth and development

molecule involved in host defense against microbes

positively influence plant development

Fig 2: Proposed model of infection network of M. truncatula involving symbiosis and pathogenesis

During the initial infection

receptors perceive extra-

triggering a host cellular

binding of Rac1 leading to

phase, the host membrane

cellular stimuli from both the

response. Rboh is activated by

generation of ROS, which can result to cell death or initiation

of second signalling cascade

symbiosis specific genes

and downstream activation of

symbiont and the pathogen

Retarded growth and low nodulation of MtRac1-deficient lines shows

Activation of Rac1 protein confirmed its role as an early signalling

MtRac1 protein is not only involved in host plant defense but also

For identification, functional analysis and evaluation of protein profiles,

proteomic analysis and mass spectrometry are currently carried out

Background

Legumes are among the most economically important crops worldwide. They establish mutualistic symbioses and pathogenic interactions with microbes. Their infection physiology involves protein-protein interactions with the host plant generating an initial defense response, presumed to be similar at the very early infection phase but later split into symbiotic and pathogenic specific cellular defense responses.

We characterized the two host defense response patterns, using the model legume *Medicago truncatula* as the host for nitrogen-fixing rhizobial bacteria Sinorhizobium meliloti and arbuscular mycorrhizal fungi Glomus intraradices as well as root rot oomycete pathogen Aphanomyces *euteiches.* To functionally characterize the role of defense proteins during the very early phases of infection, we used plants deficient of Rac1 protein. Rac1 is a small monomeric GTP binding protein and a key signalling factor in eukaryotic cells. It regulates the membrane-bound Rboh (respiratory burst oxidase homologue) which catalyses the generation of reactive oxygen species (ROS) in plants in response to microbial infection. microbial infection.

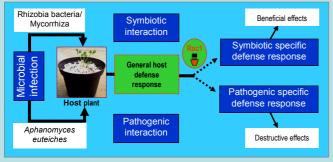


Fig 1: Symbiotic and pathogenic host plant defense response pathways

Phenotyping Medicago truncatula plant lines

Fig 3: (a) Wild type vs. MtRac1- deficient lines and (b) wild type vs. vector control plants



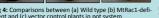


Fig 5: (a) Vector control (b) w type (c) MtRac1-deficient

- Stunted growth in MtRac1-deficient plants compared to wild type plants but the vector control plants developed normally
- More hairy roots/cm root-length in MtRac1-deficient plants compared to vector control
- Tapering and/or deformed leaves in MtRac1-deficient plants but normal development in wild type and vector control lines

Results and Discussion

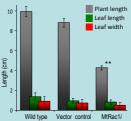
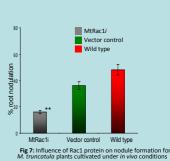


Fig 6: Comparisons of different parameters for plants cultured under in vitro conditions

- Difference in plant length between MtRac1-deficient plants, vector control and wild type plants was significantly high
- MtRac1-deficient lines had smaller leaf size than the wild type and vector control lines
- Significantly higher nodulation in wild type than MtRac1-deficient and vector control lines

us K² and Colditz F^{1*} 2011. Silencing of the Rac1 GTPase MtROP9 in



neficial effects

Gene expression analysis at different time-points intraradices (control) MtRac1

Fig 8: Gene expression analysis at 3 and 24hrs post inoculation via semi -quantitative RT-PCR using Rac1 primer system

Expression level of Rac1 was clearly suppressed in MtRac1 -deficient plants compared to vector control and even after infection with Aphanomyces euteiches

Notes on Materials and Methods

- · Generation of transgenic plants via Agrobacterium rhizogenes-mediated transformation: RNAi - gene silencing approach
- Phenotyping of MtRac1-deficient, vector control and wild type plants under in vitro and in vivo conditions
- Microbial infection and evaluation of infection profiles
- Identification and characterization of protein profiles via 2D-Gel electrophoresis and mass spectrometry



Fig 9: M. truncatula transgenic root cultures and as obs florescence-dsRED to distinguish transgenic hairy roots

ference: Kiirika Leonard M¹, Bergmann H F², Schikowsky C¹, Wimmer D¹, Korte J¹, Udo Schmitz¹, Ni : negatively affects rhizobial infection (*Submitted for publication*) ent: This study was carried out with the support of Deutscher Akademischer Austausch Dienst (DAAD)