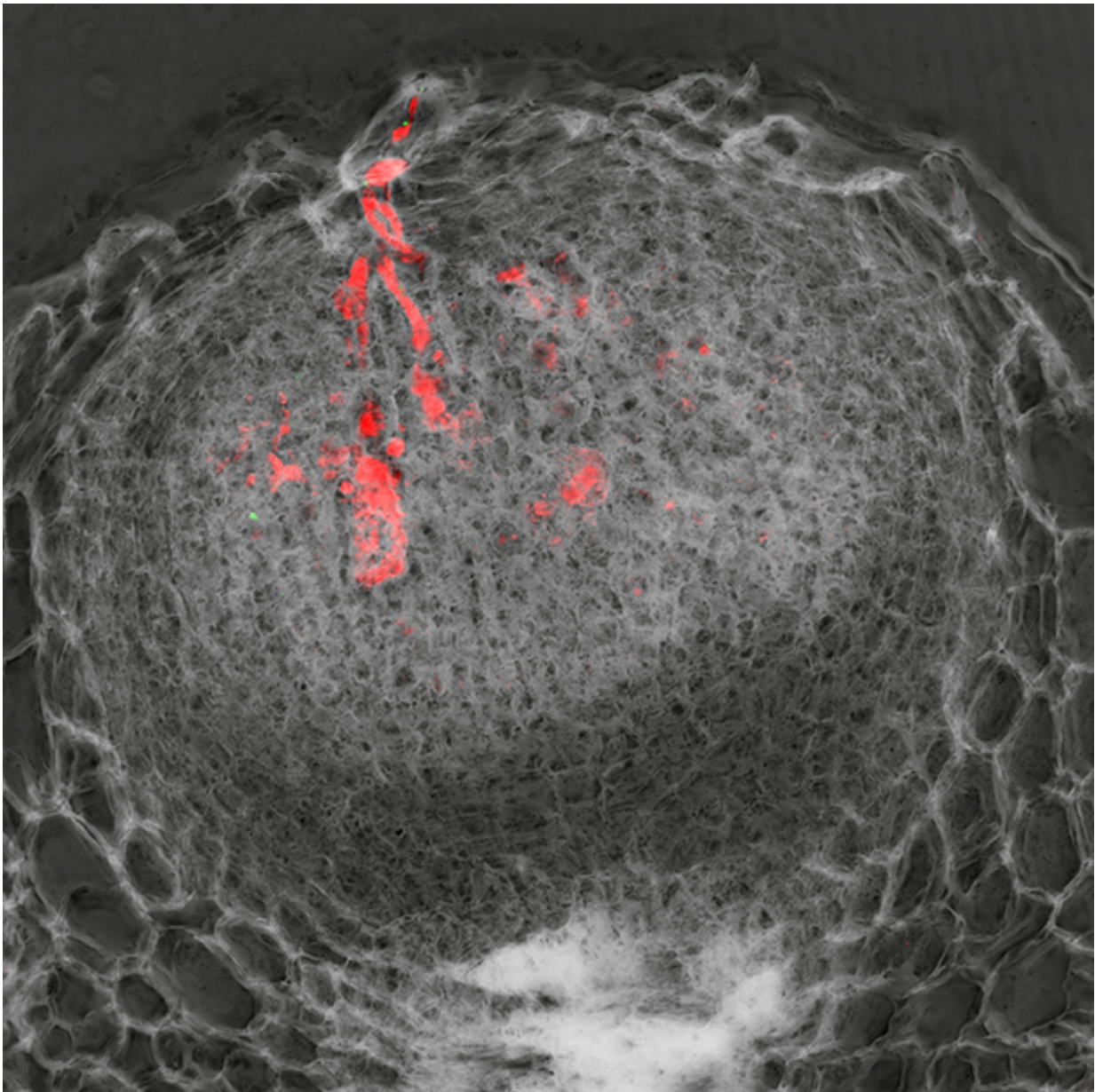


Understanding images: A genetic framework in legumes controls infection of nodules

July 31 2015, by Simon Kelly And Simona Radutoiu



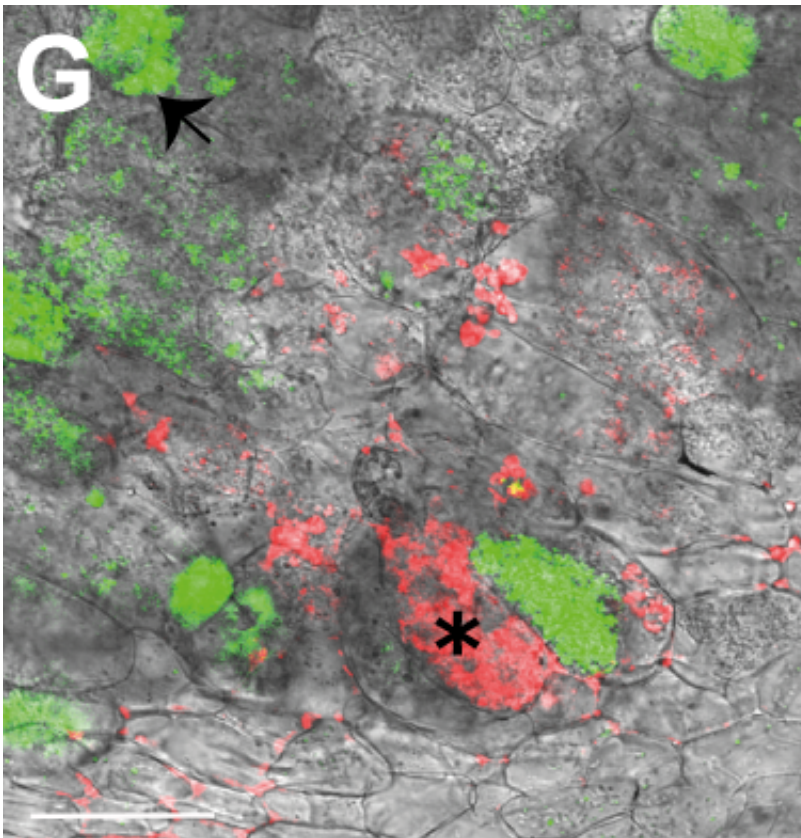
Confocal laser scanning microscopy image of a nodule section illustrating the internal infection pattern of the endophyte *Rhizobium mesosinicum*, strain KAW12 (red) and of the incompatible symbiont *Mesorhizobium loti*, strain exoU (green). Image Credit: Rafal Zgadza

The soil environment harbors a diverse range of bacteria, many of which could potentially be detrimental if they are able to gain entry to plant tissues. We are interested in determining how the host plant selects which bacteria are able to colonize its tissues and to identify important endophyte factors that allow them to be accommodated by the host plant. In this issue of *PLOS Genetics* we investigate the genetic components and molecular signals that allow the endophyte *Rhizobium mesosinicum* strain KAW12 (KAW12) to colonize symbiotically induced nodules on the model legume *Lotus japonicus*. We have used different symbiotic and endophytic strains and performed mixed inoculations of wild-type or symbiotic *L. japonicus* mutants in order to identify the respective contributions of the different interacting partners – legume host, symbiont and endophyte.

Colonisation of *Lotus japonicus* Nodules by Endophytic Bacteria

The legume root nodule is a unique environmental niche induced by [symbiotic bacteria](#), where multiple symbiotic and endophytic bacterial species can co-exist. Several endophytes were tested for their ability to colonize *L. japonicus* nodules that were induced by its usual symbiotic rhizobia (*Mesorhizobium loti*) in co-inoculation experiments. Our study identified KAW12 as an [endophyte](#) that uses the symbiotically induced infection threads to co-colonize *L. japonicus* nodules without inducing nodule necrosis, providing us with a system to study host and endophyte genetic features that are important for such interactions.

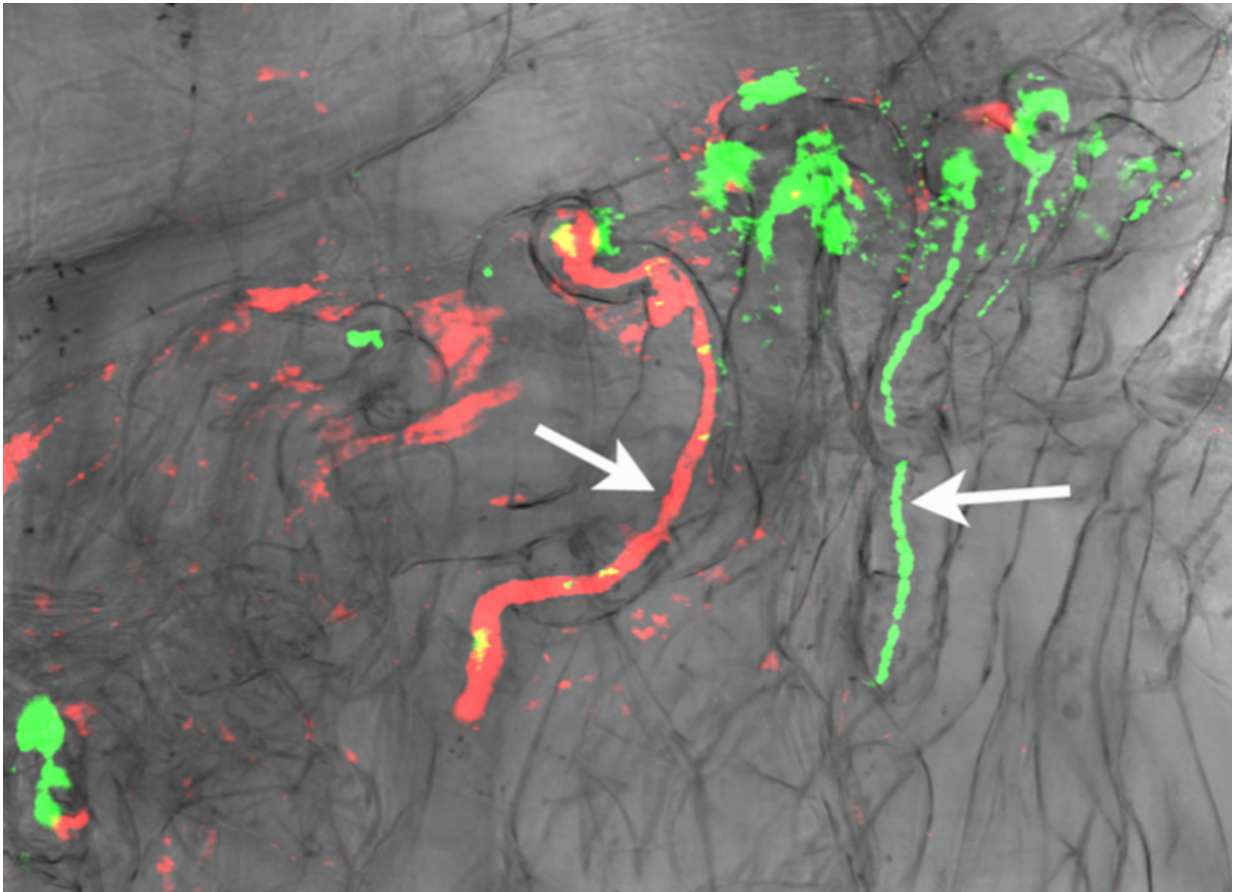
Initiation of the symbiotic process requires Nod-factor signaling. Nod factors are signal molecules produced by symbiotic rhizobia and recognized by the plant through Nod-factor receptors. This perception triggers the initiation of the infection pathway, through which rhizobia enter the plant, as well as initiation of nodule organogenesis. Using symbiotic strains that produce different types of Nod factors as co-inoculating partners, we determined that intact Nod-factor signaling provided by the symbiont is required for nodule colonization by the endophytic KAW12 bacteria.



Section of an *M. loti* nodZ-induced nodule presenting KAW12 (*) and *M. loti* nodZ (arrow) infection. Credit: Zgadzaj et al.

Exopolysaccharides are Key for Chronic Infection

Exopolysaccharide (EPS) production is important during symbiosis⁵. The *M. loti* *exoU* strain utilized in this study is affected in EPS biosynthesis and is thus impaired in symbiosis due to an inability to form infection threads. To further investigate this we isolated an EPS-deficient strain of KAW12 and performed co-inoculation experiments with *M. loti* *exoU*. Strikingly, nodule colonization by the KAW12 EPS mutant was absent, indicating that EPS is a key molecule required by the endophyte to allow for nodule colonization.



Root hair infection threads (arrows) colonised by *M. loti* *exoU* (green) and KAW12 (red). Credit: Zgadzaj et al.

Endophyte Nodule Occupancy is Host-Controlled

To study the role of the legume host in mixed inoculations we took advantage of the large collection of *L. japonicus* mutants available to identify host genetic components required for nodule colonization by the endophyte. Our results revealed that the mutation of genes required for infection thread formation prevented nodule colonization by the KAW12 endophyte. In contrast, KAW12 colonization of nodules formed on plant mutants in genes required for supporting nitrogen-fixation within nodules was not impaired.

The presence of endophytes within legume nodules may restrict the occupancy of effective nitrogen-fixing symbionts and therefore represents a major challenge that contributes to limiting legume cultivation. Our study shows that the well-established genetic resources available for the model legume *L. japonicus* can be utilised in co-inoculation studies to identify genetic and molecular factors important for determining compatibility with soil bacteria, providing further avenues to address this issue.

More information: ¹ Broghammer, A. et al. Legume receptors perceive the rhizobial lipochitin oligosaccharide signal molecules by direct binding. *Proc Natl Acad Sci U S A* 109, 13859-13864, [DOI: 10.1073/pnas.1205171109](https://doi.org/10.1073/pnas.1205171109) (2012).

² Radutoiu, S. et al. Plant recognition of symbiotic bacteria requires two LysM receptor-like kinases. *Nature* 425, 585-592 (2003).

³ Oldroyd, G. E. D., Murray, J. D., Poole, P. S. & Downie, J. A. The rules of engagement in the legume-rhizobial symbiosis. *Annu. Rev.*

Genet. 45, 119-144, [DOI: 10.1146/annurev-genet-110410-132549](https://doi.org/10.1146/annurev-genet-110410-132549) (2011).

⁴ Kawaharada, Y. et al. Receptor-mediated exopolysaccharide perception controls bacterial infection. *Nature* 523, 308-12 (2015).

⁵ Kelly, S. J. et al. Conditional requirement for exopolysaccharide in the Mesorhizobium-Lotus symbiosis. *Mol Plant Microbe Interact* 26, 319-329, [DOI: 10.1094/mpmi-09-12-0227-r](https://doi.org/10.1094/mpmi-09-12-0227-r) (2013).

This story is republished courtesy of PLOS Blogs: blogs.plos.org.

Provided by PLOS Blogs

Citation: Understanding images: A genetic framework in legumes controls infection of nodules (2015, July 31) retrieved 23 April 2024 from <https://phys.org/news/2015-07-images-genetic-framework-legumes-infection.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.