

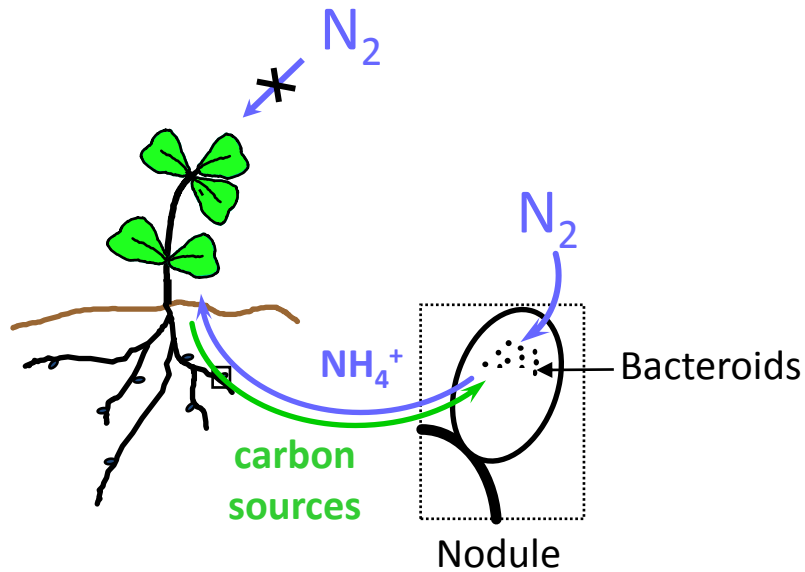
Experimental evolution of rhizobia:

Transient hypermutagenesis accelerates the evolution of legume symbionts following horizontal gene transfer

Delphine Capela

Group of Catherine Masson and Jacques Batut

The rhizobium-legume symbiosis, an agronomical and ecological important association

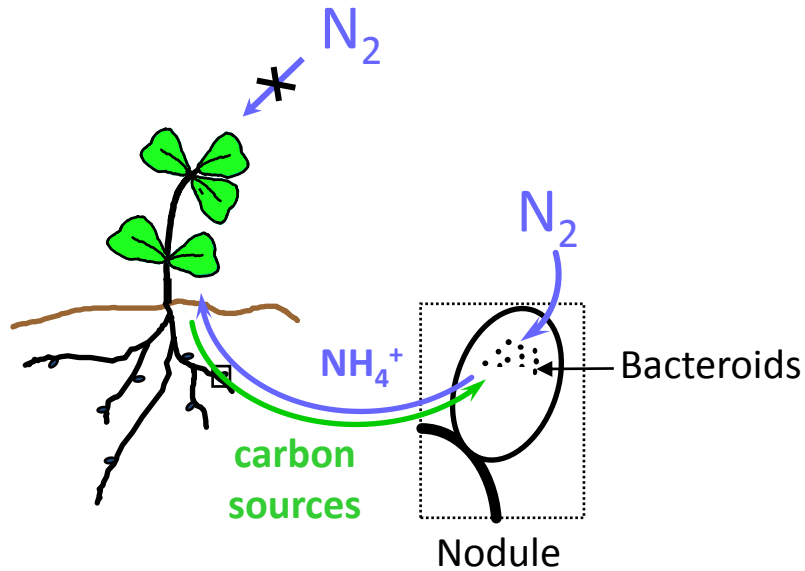


Biological nitrogen fixation by legumes accounts for ~ 100 millions MT of N_2 reduced into NH_4^+

← Not inoculated → ← Inoculated soybean →



The rhizobium-legume symbiosis, an agronomical and ecological important association



Biological nitrogen fixation by legumes accounts for ~ 100 millions MT of N_2 reduced into NH_4^+

Legumes consist in ~17 000 species

← Not inoculated → ← Inoculated soybean →



Mimosa pudica
C. taiwanensis



Sesbania rostrata
A. caulinaudans

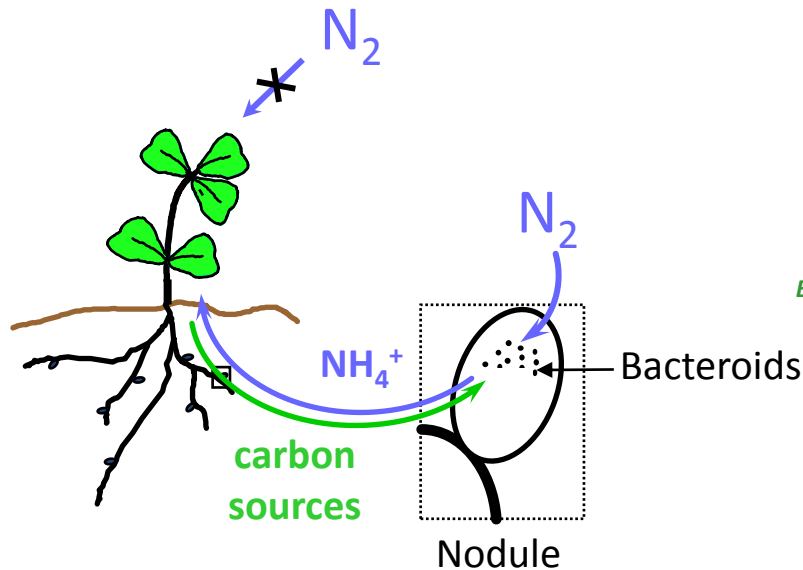


Aeschynomene
afraspera
B. sp. ORS322



Aeschynomene
sensitiva
B. sp. ORS278

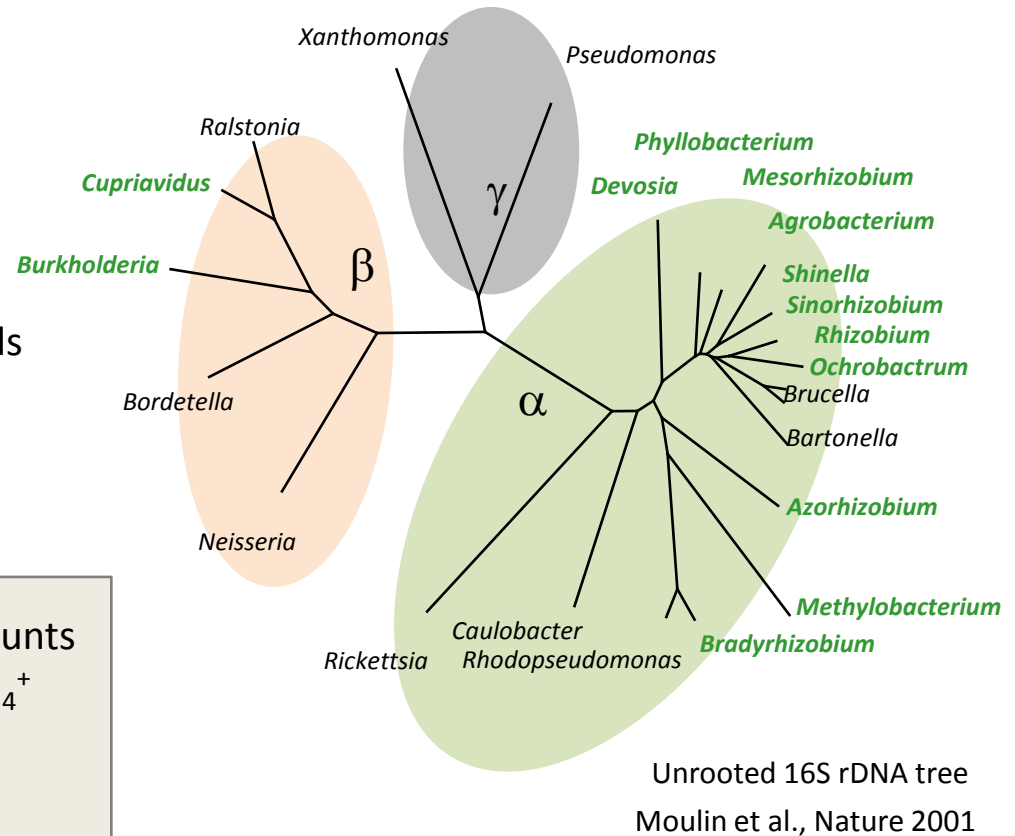
The rhizobium-legume symbiosis, an agronomical and ecological important association



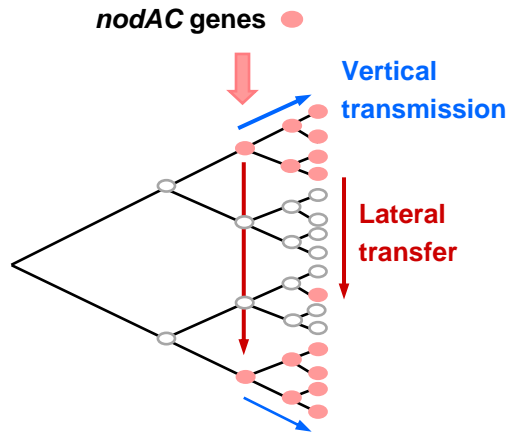
Biological nitrogen fixation by legumes accounts for ~ 100 millions MT of N_2 reduced into NH_4^+

Legumes consist in ~17 000 species

Rhizobia were identified in 13 genera and >100 species



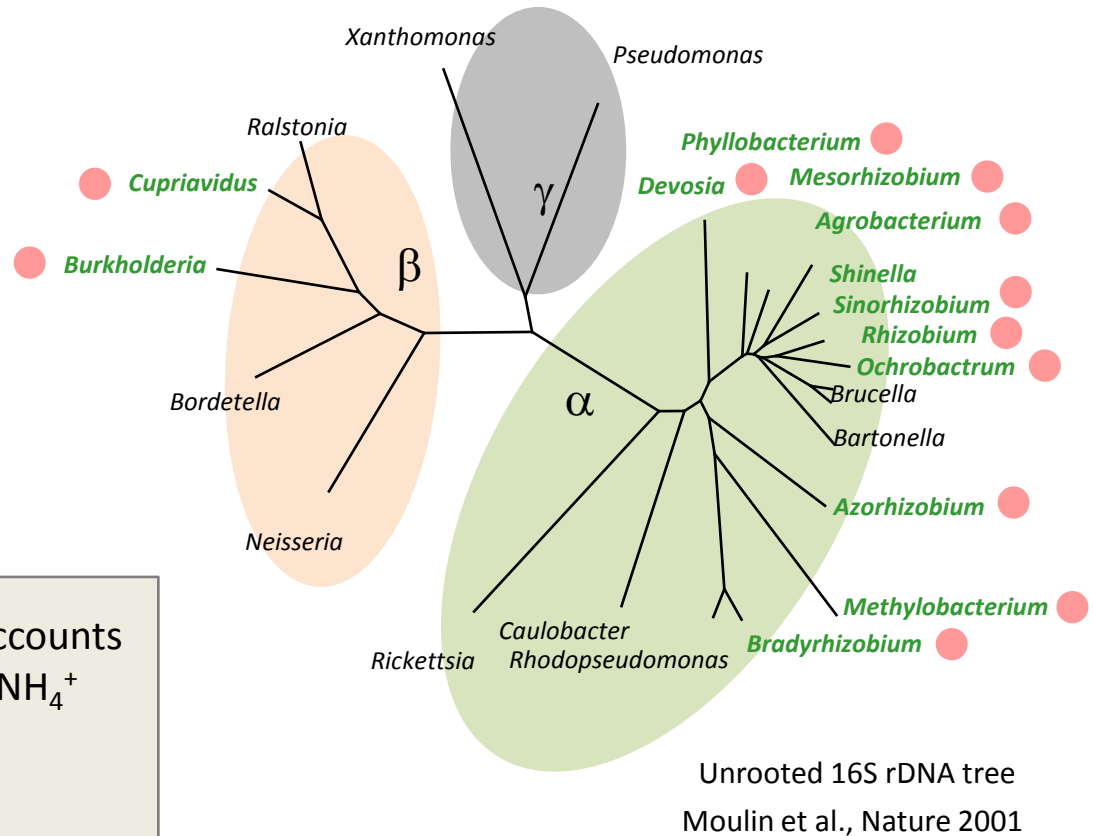
Rhizobia are phylogenetically diverse N₂-fixing legume symbionts



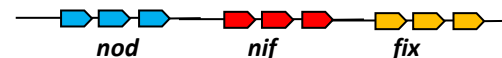
Biological nitrogen fixation by legumes accounts for ~ 100 millions MT of N₂ reduced into NH₄⁺

Legumes consist in ~17 000 species

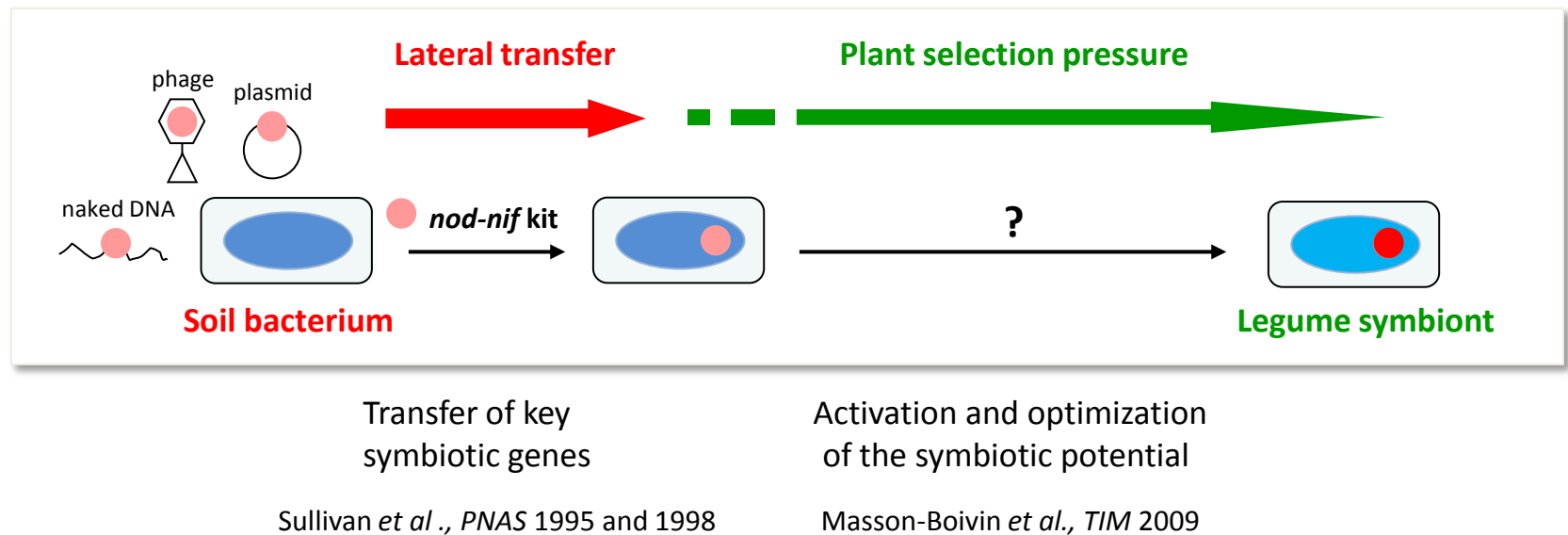
Rhizobia were identified in 13 genera and >100 species



● *nod-nif* kit on MGE

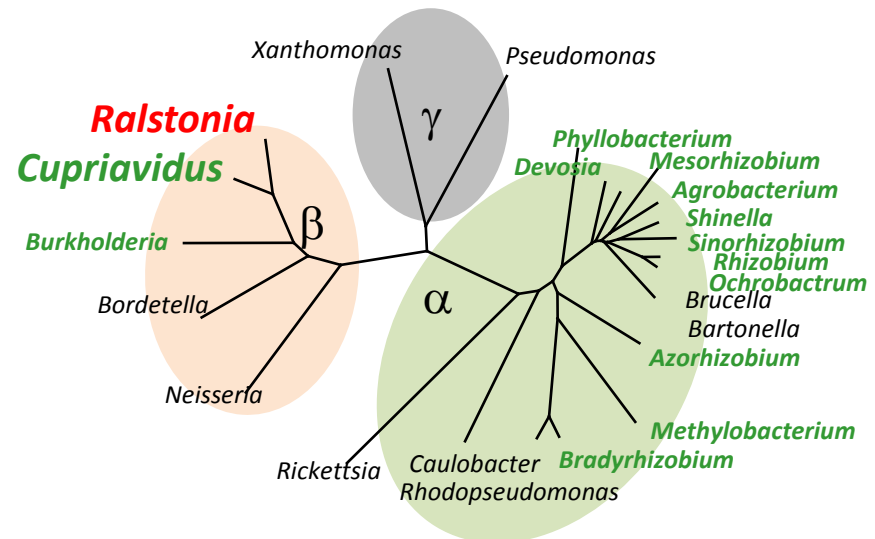
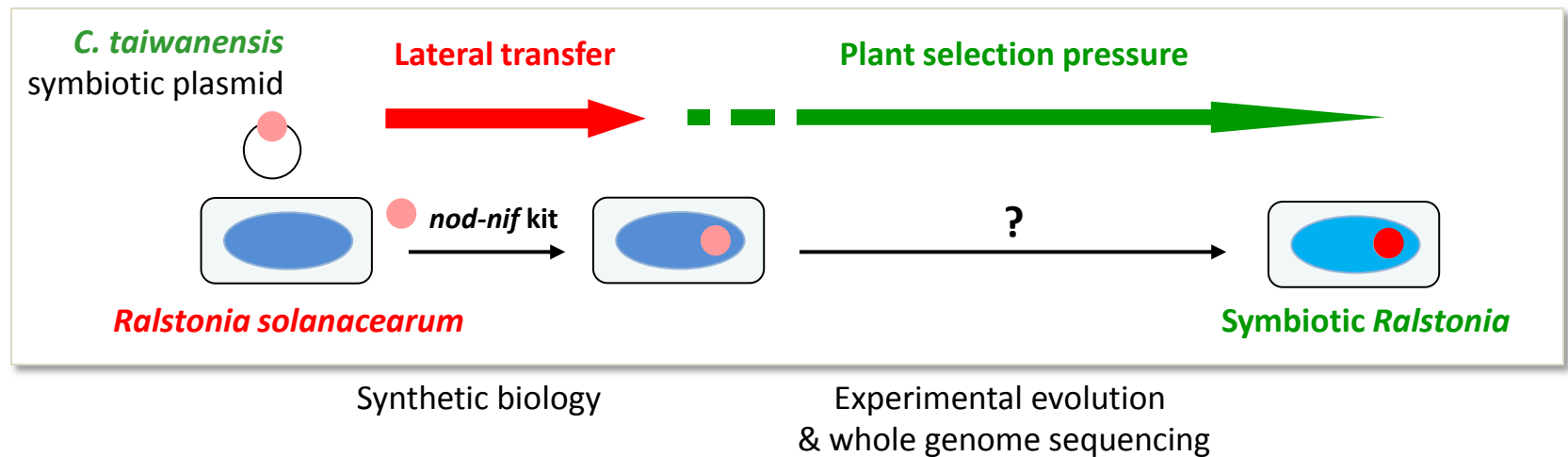


A two step-based scenario for rhizobium evolution



How complex phenotypic traits such as the ability to fix nitrogen with legumes have successfully spread over large phylogenetic distances?

Experimental evolution of a plant pathogen into legume symbionts



Construction of the chimeric strain

Design

C. taiwanensis



pSym



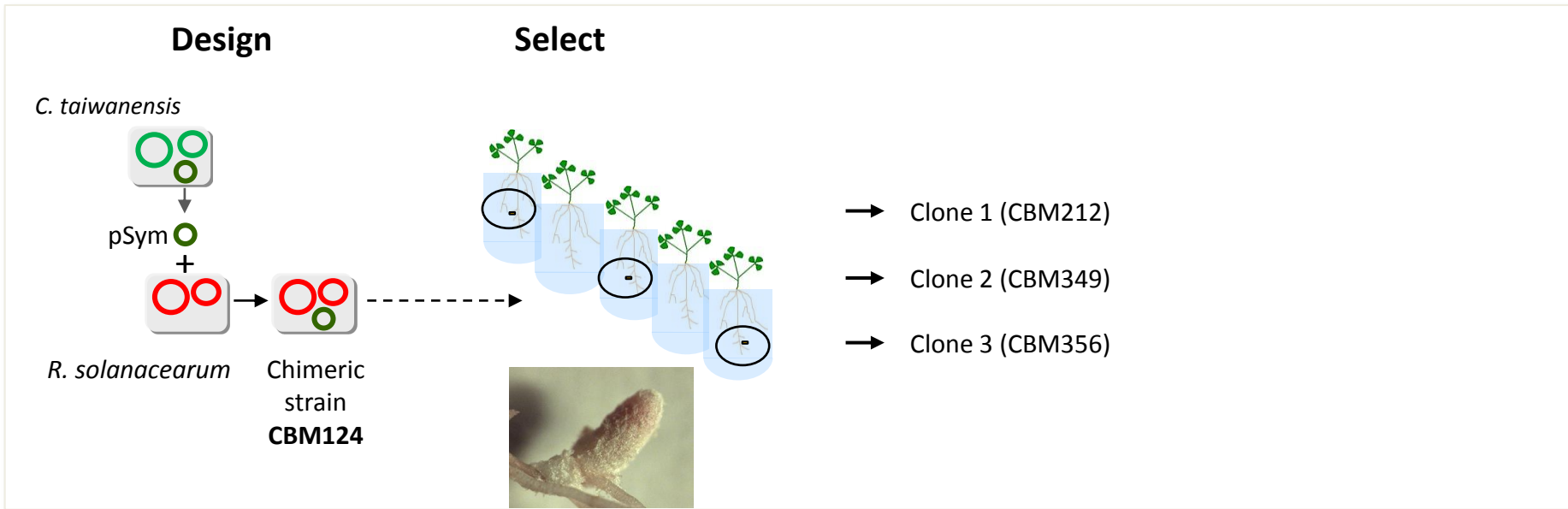
+



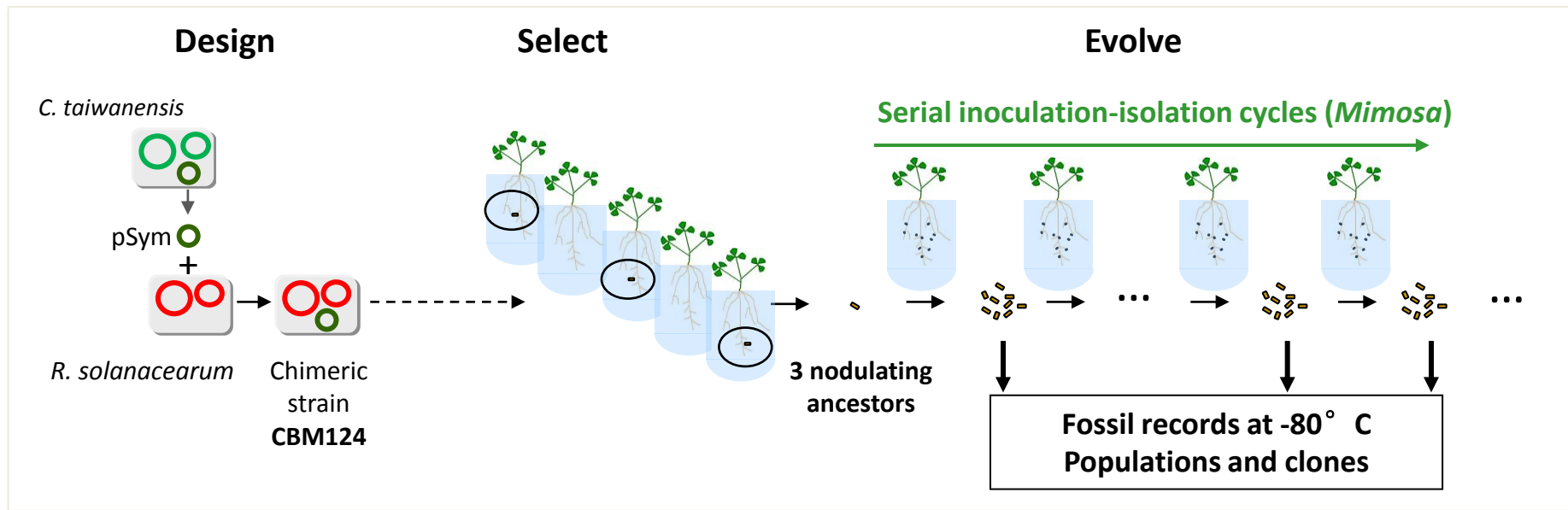
R. solanacearum

Chimeric
strain
CBM124

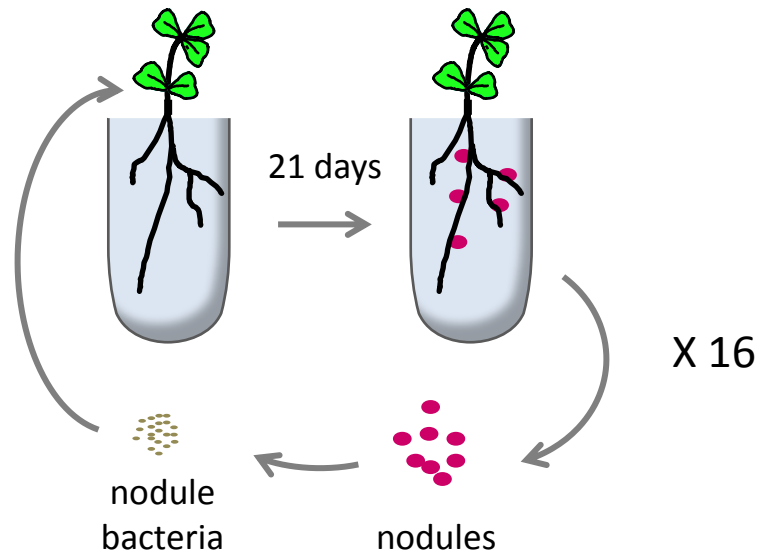
Obtaining nodulating chimeric clones



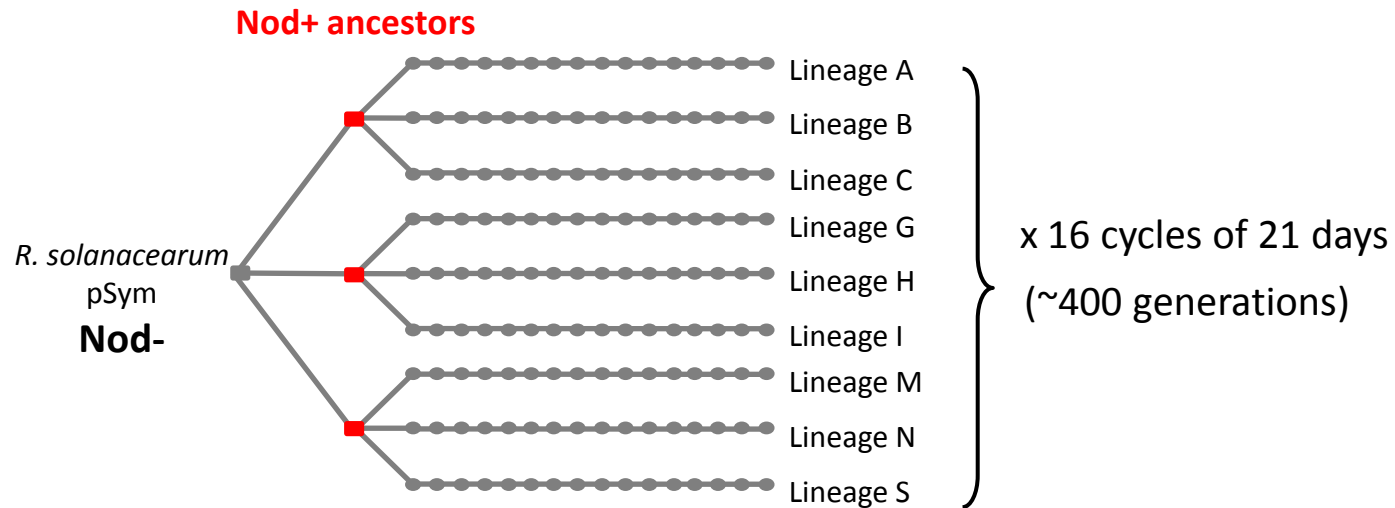
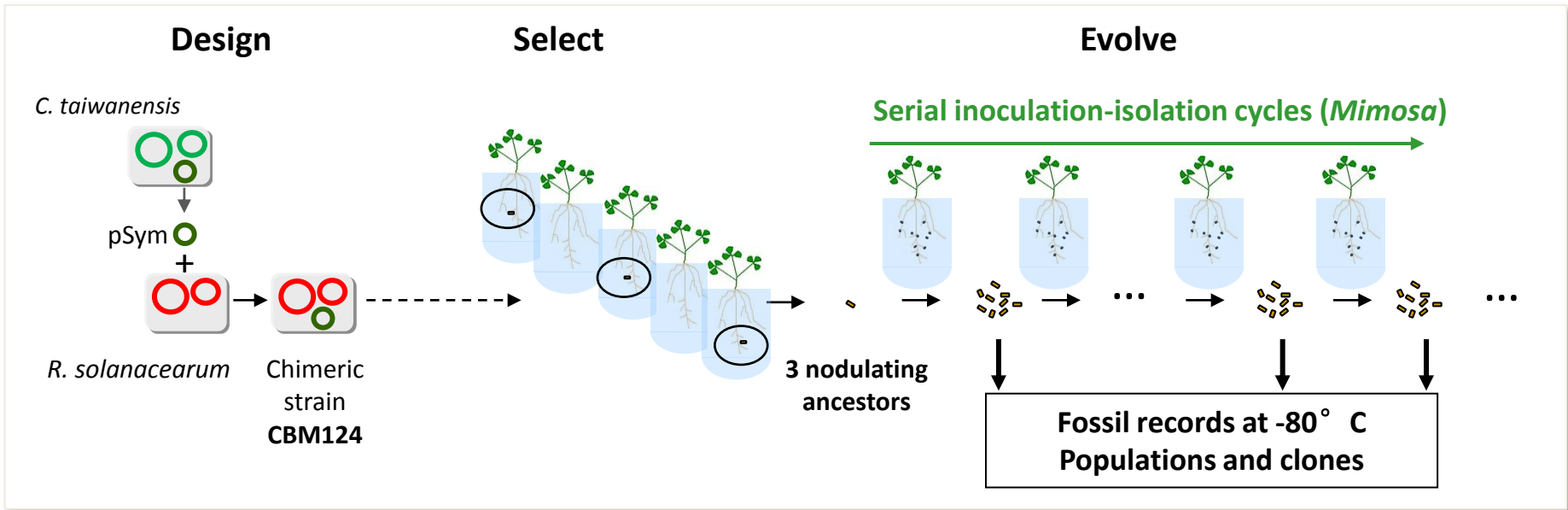
The evolution experiment



Experimental evolution cycles

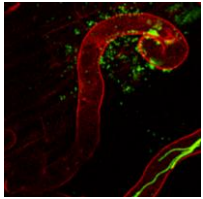


The evolution experiment

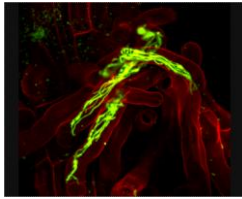


Evolution of symbiotic properties of evolved clones

C. taiwanensis / *M. pudica*



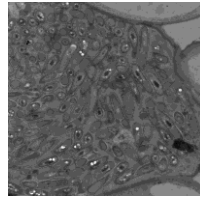
Infection site
formation



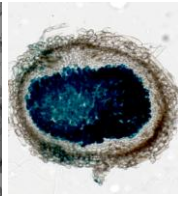
Infection thread
formation



Nodule
formation



Intracellular
infection



Bacteroid
persistence



Nitrogen
fixation

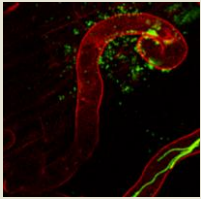
Root hair entry and nodulation

Nodule cell infection

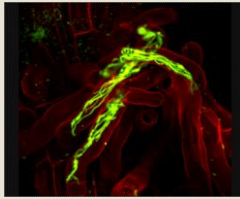
N₂ fixation

Evolution of symbiotic properties of evolved clones

C. taiwanensis / *M. pudica*



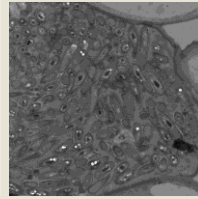
Infection site formation



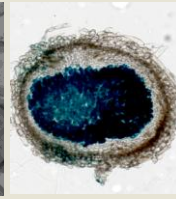
Infection thread formation



Nodule formation



Intracellular infection



Bacteroid persistence

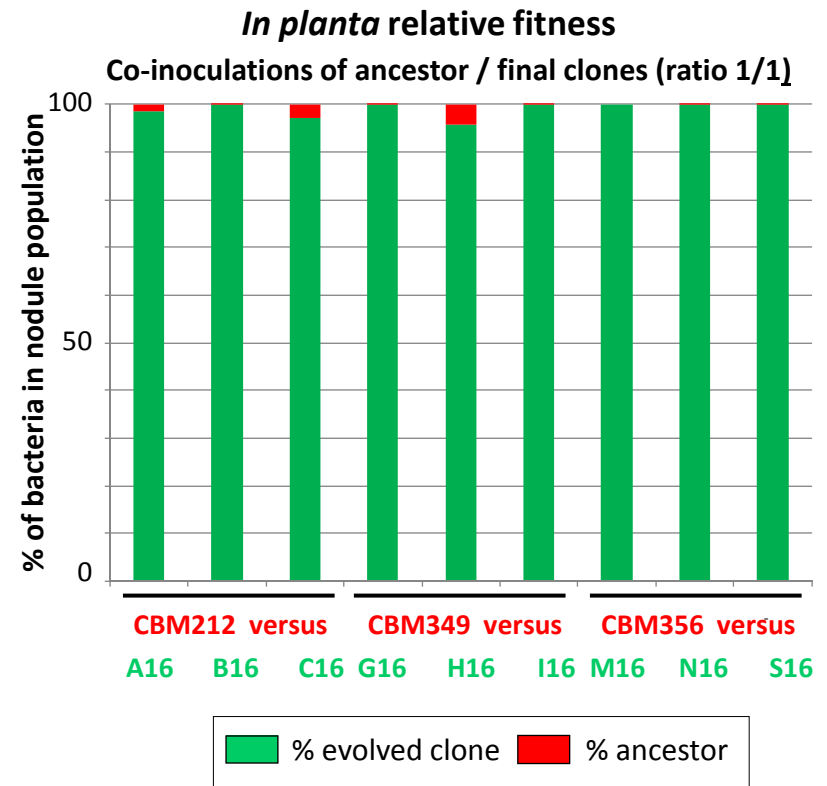
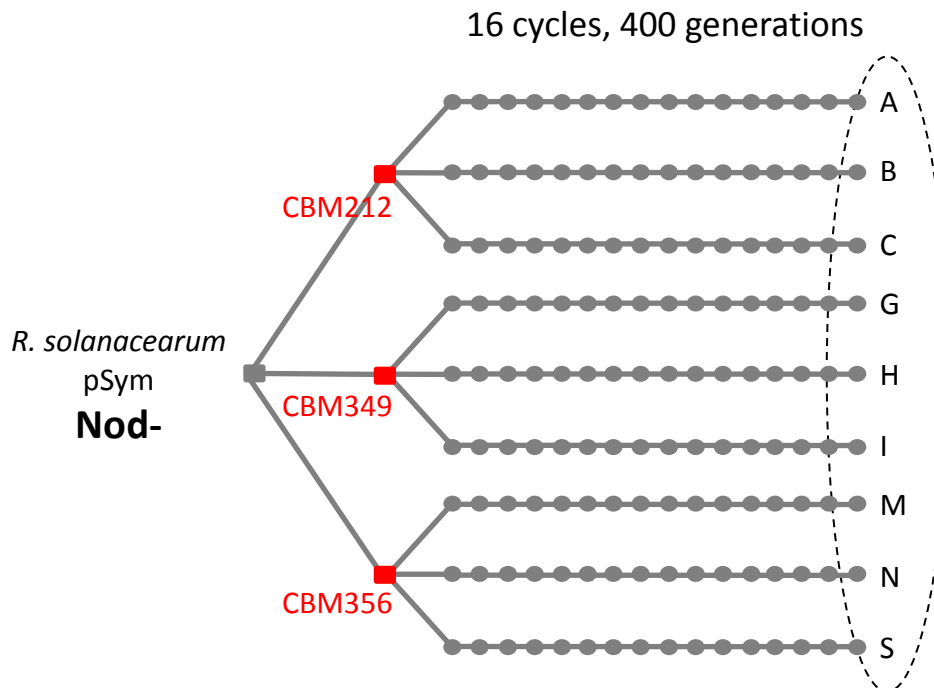


Nitrogen fixation

Root hair entry and nodulation

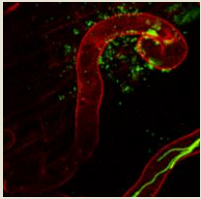
Nodule cell infection

N₂ fixation

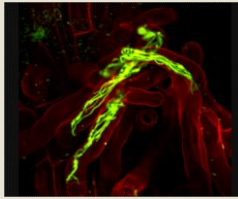


Evolution of symbiotic properties of evolved clones

C. taiwanensis / *M. pudica*



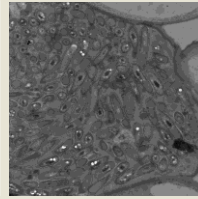
Infection site formation



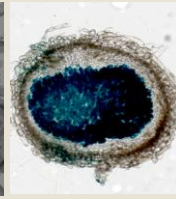
Infection thread formation



Nodule formation



Intracellular infection



Bacteroid persistence

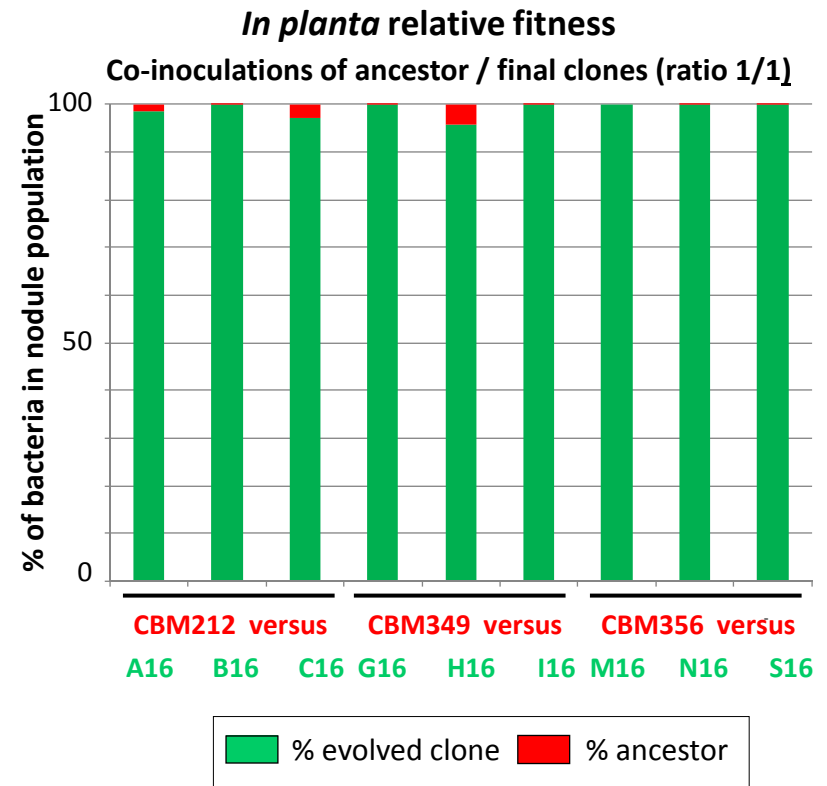
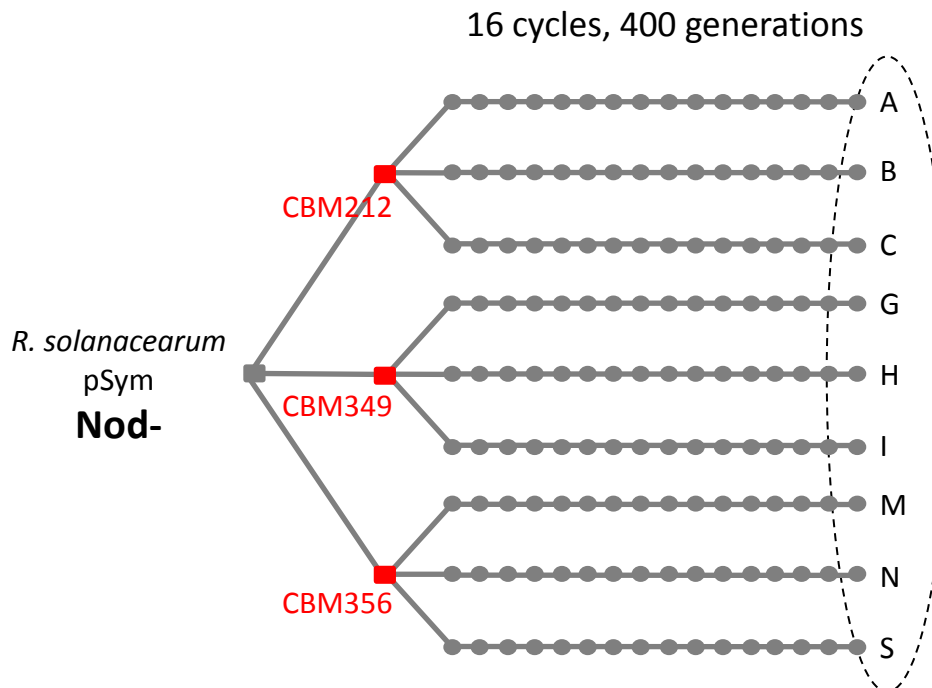


Nitrogen fixation

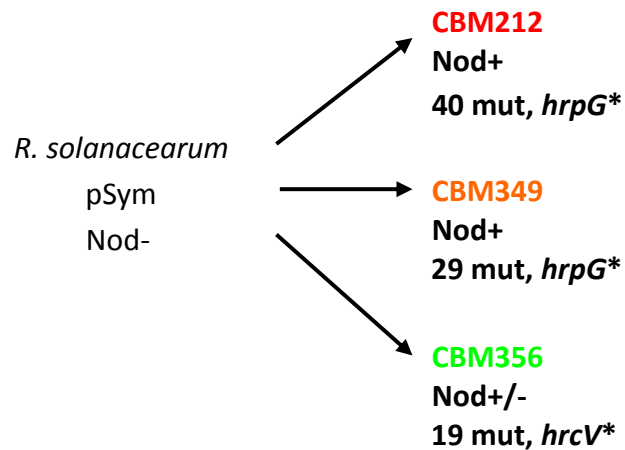
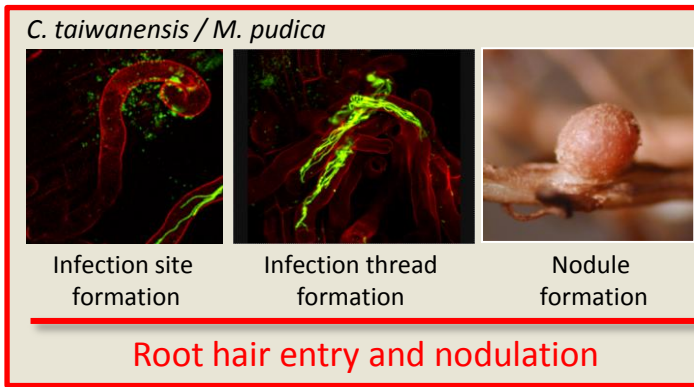
Root hair entry and nodulation

Nodule cell infection

N₂ fixation

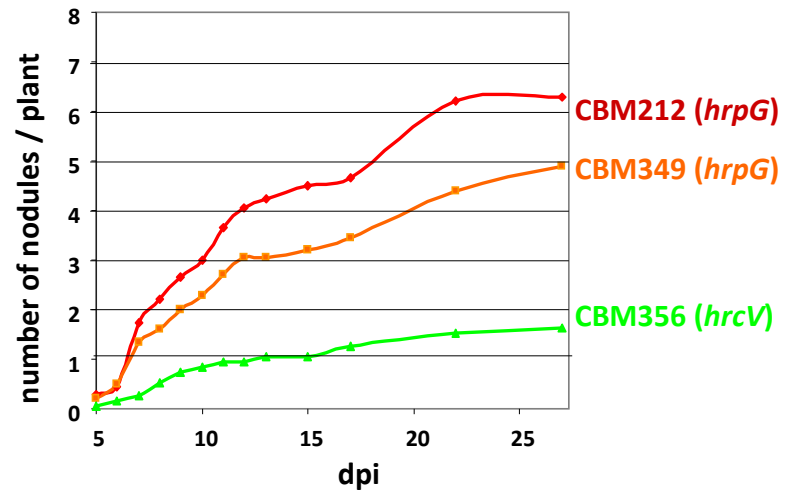


Activation of nodulation

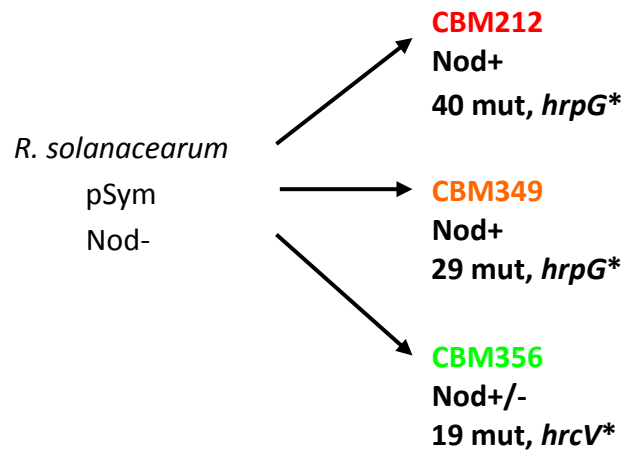
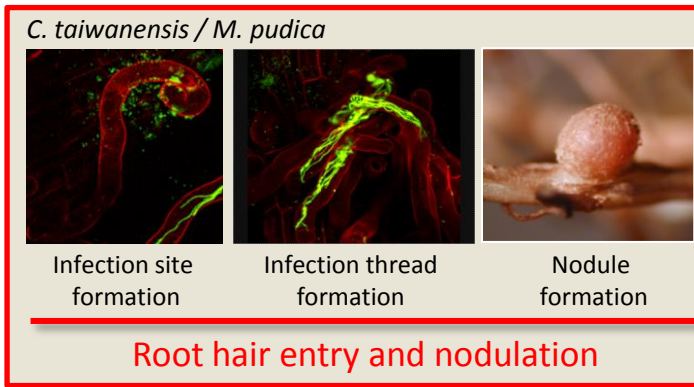


SNPs/Indels
(SNiPer - MaGe MicroScope platform)

Nodulation kinetics

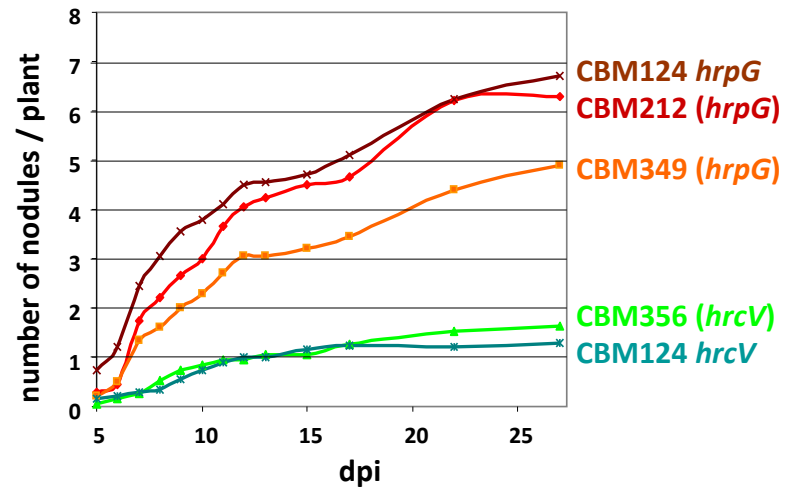


Activation of nodulation

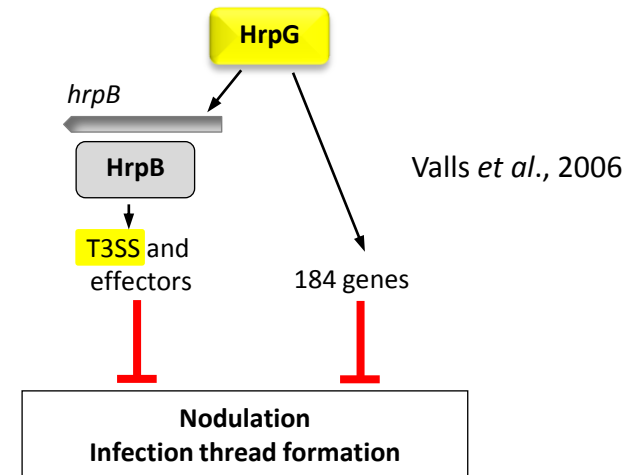
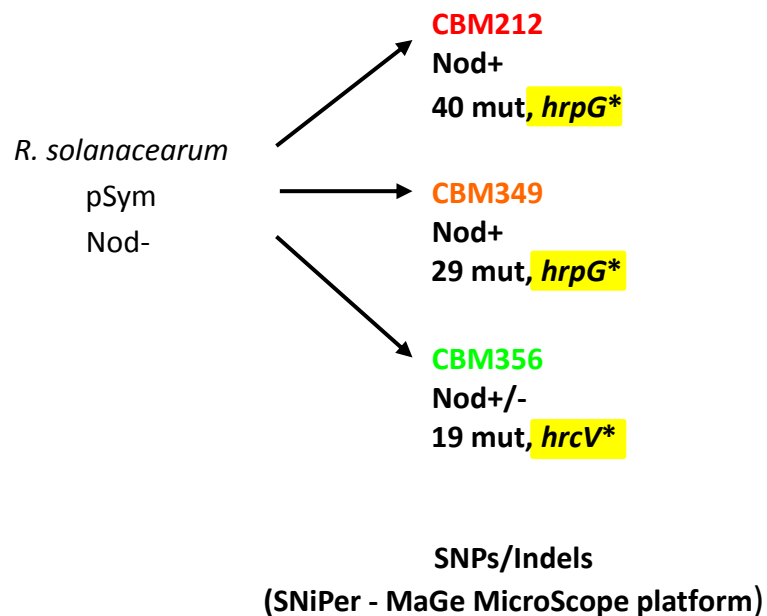
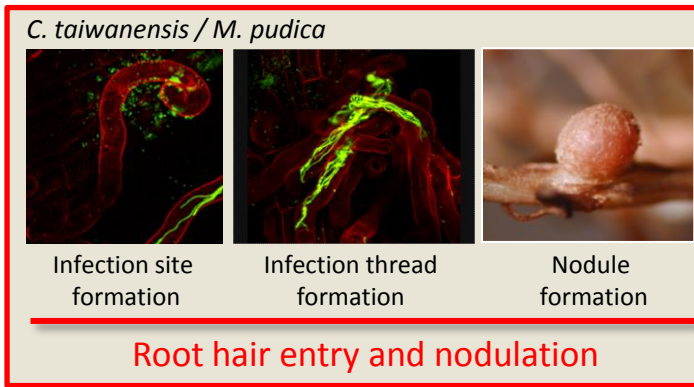


SNPs/Indels
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Nodulation kinetics

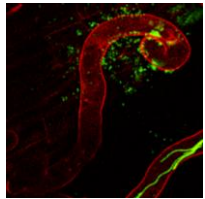


Activation of nodulation

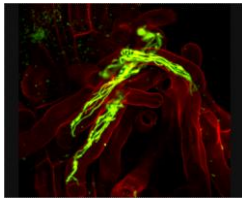


Activation of intracellular infection

C. taiwanensis / *M. pudica*



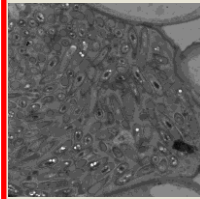
Infection site formation



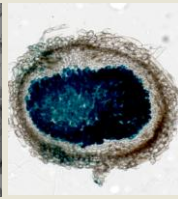
Infection thread formation



Nodule formation



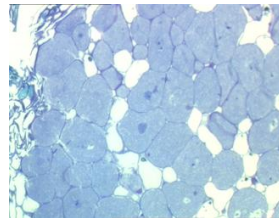
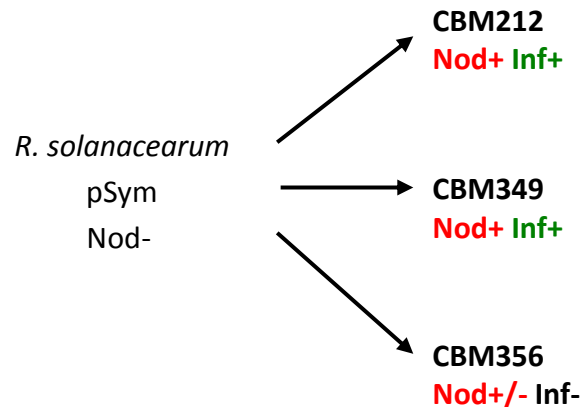
Intracellular infection



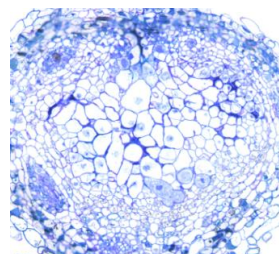
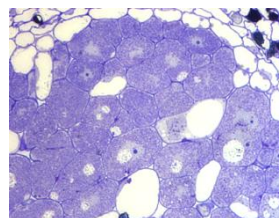
Bacteroid persistence

Root hair entry and nodulation

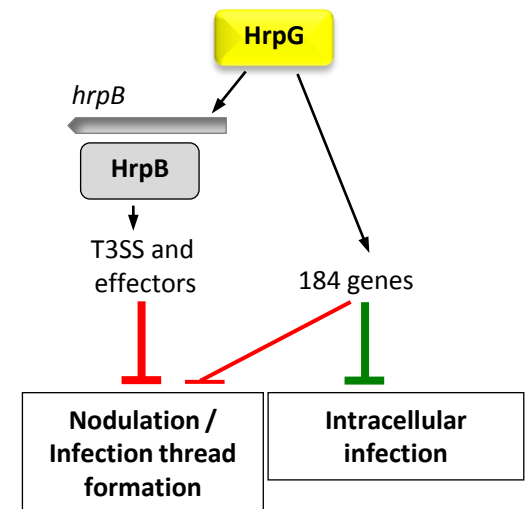
Nodule cell infection



*hrpG**

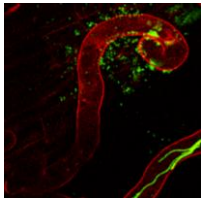


*hrcV**

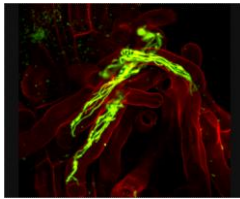


Activation of intracellular infection

C. taiwanensis / *M. pudica*



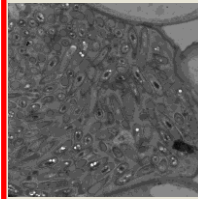
Infection site formation



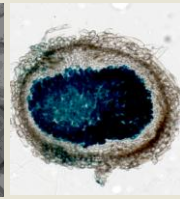
Infection thread formation



Nodule formation



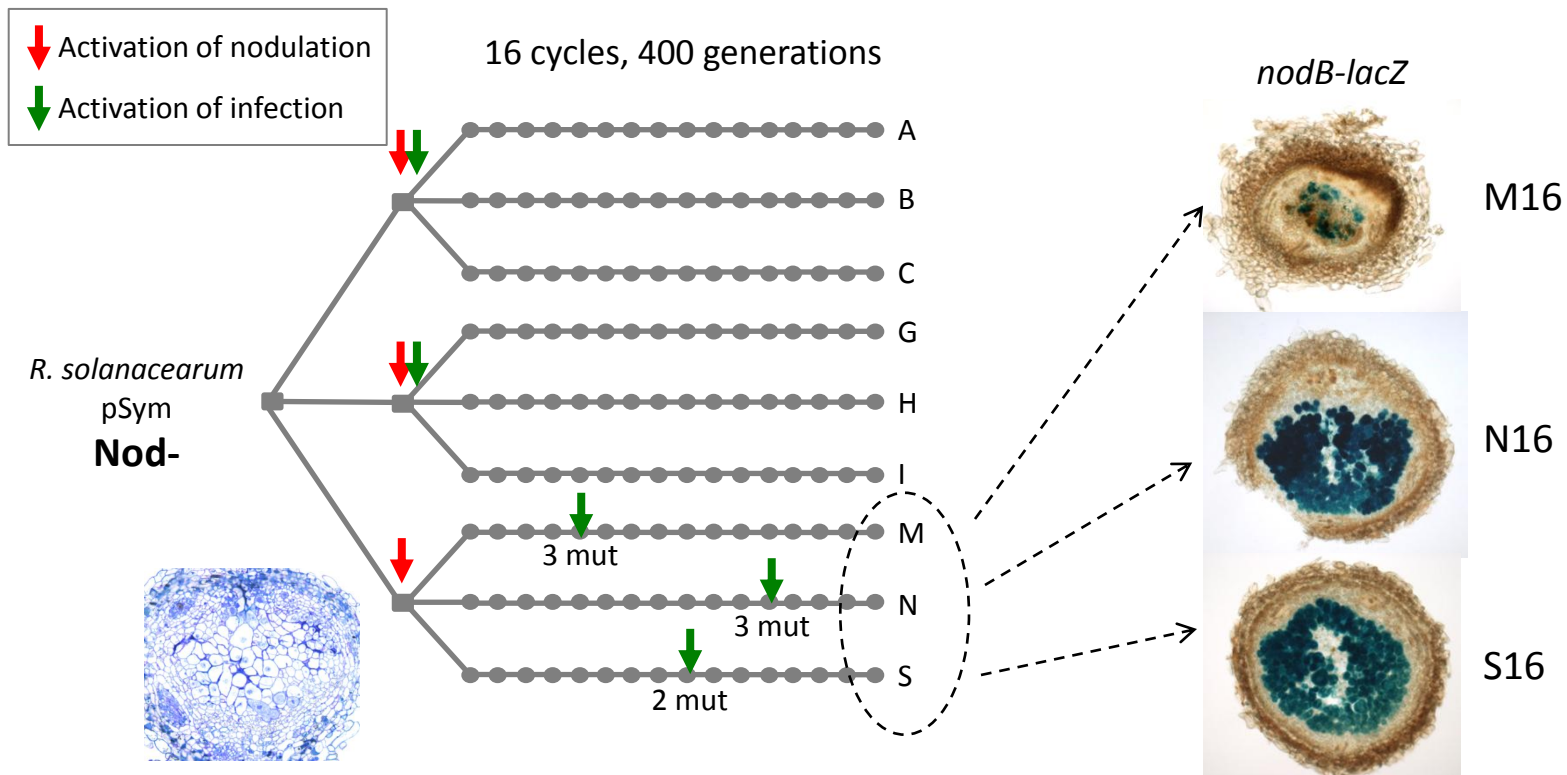
Intracellular infection



Bacteroid persistence

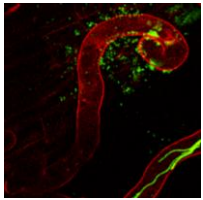
Root hair entry and nodulation

Nodule cell infection

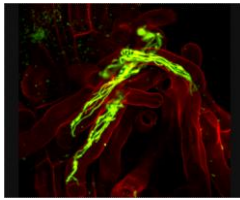


Activation of intracellular infection

C. taiwanensis / *M. pudica*



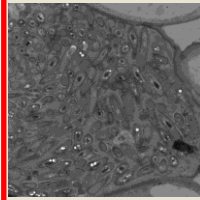
Infection site formation



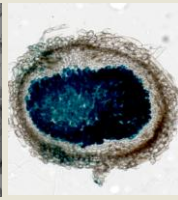
Infection thread formation



Nodule formation



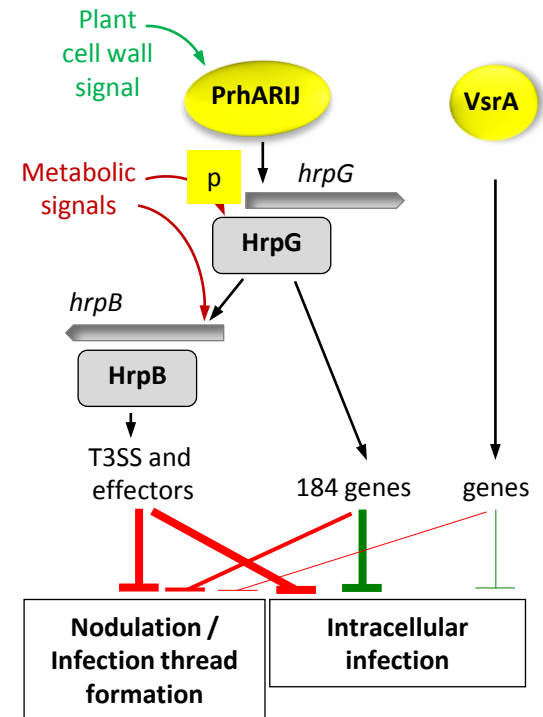
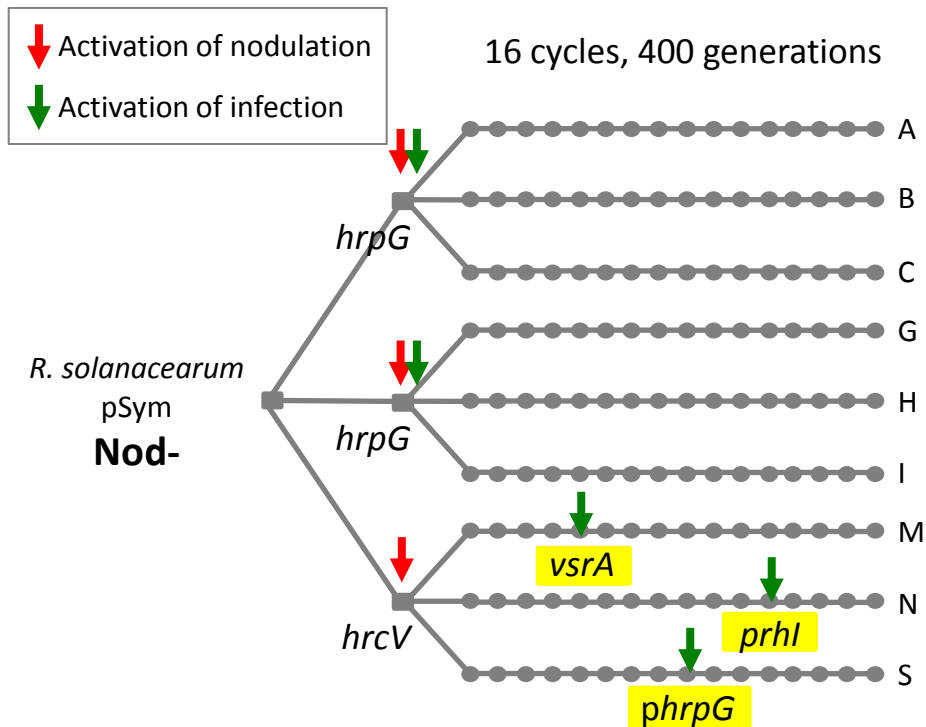
Intracellular infection



Bacteroid persistence

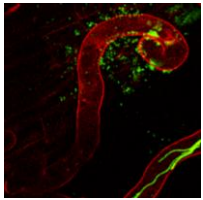
Root hair entry and nodulation

Nodule cell infection

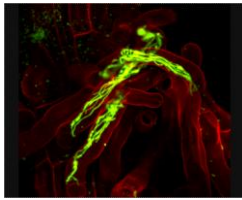


Improvement of nodulation and infection capacities along cycles

C. taiwanensis / *M. pudica*



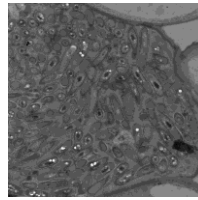
Infection site formation



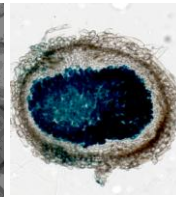
Infection thread formation



Nodule formation



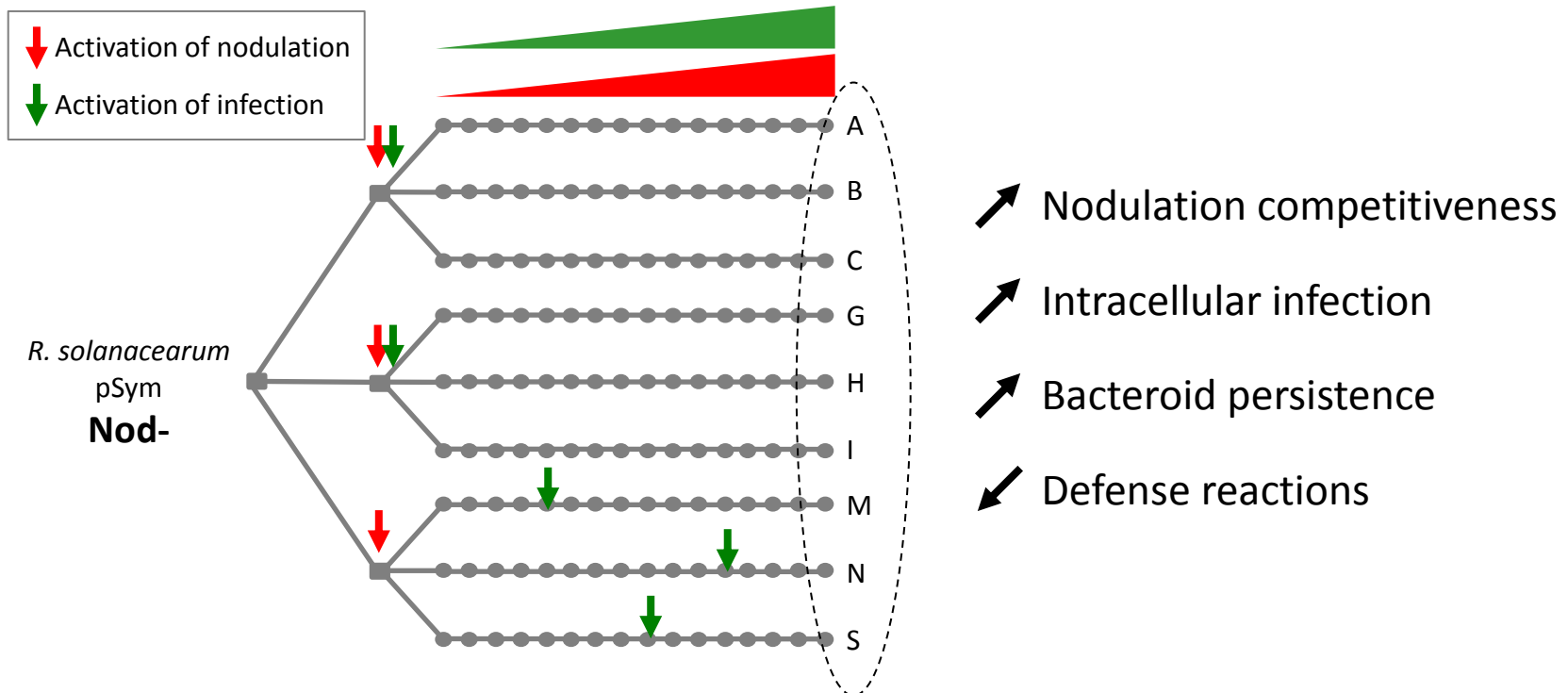
Intracellular infection



Bacteroid persistence

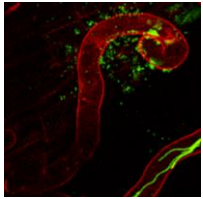
Root hair entry and nodulation

Nodule cell infection

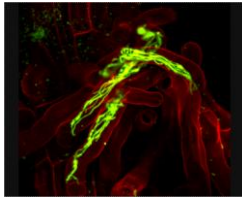


Improvements in symbiotic properties occurred by sharp shifts

C. taiwanensis / *M. pudica*



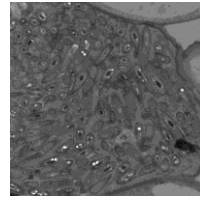
Infection site formation



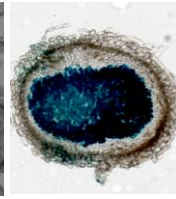
Infection thread formation



Nodule formation





Intracellular infection

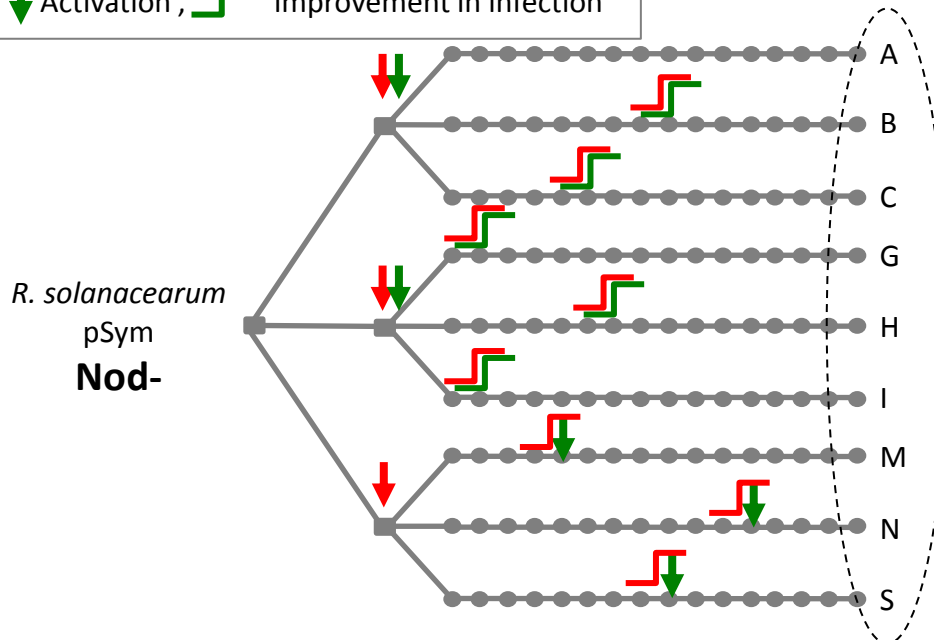


Bacteroid persistence

Root hair entry and nodulation

Nodule cell infection

↓ Activation,  improvement in nodulation
 ↓ Activation,  improvement in infection



↗ Nodulation competitiveness

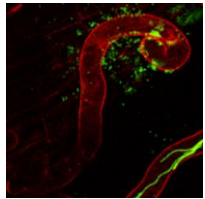
↗ Intracellular infection

↗ Bacteroid persistence

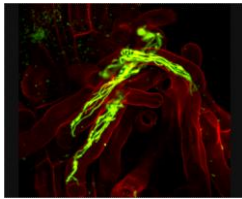
↘ Defense reactions

Evolution of symbiotic properties of evolved clones

C. taiwanensis / *M. pudica*



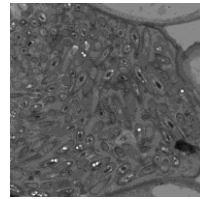
Infection site formation



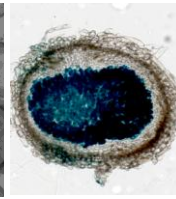
Infection thread formation



Nodule formation



Intracellular infection

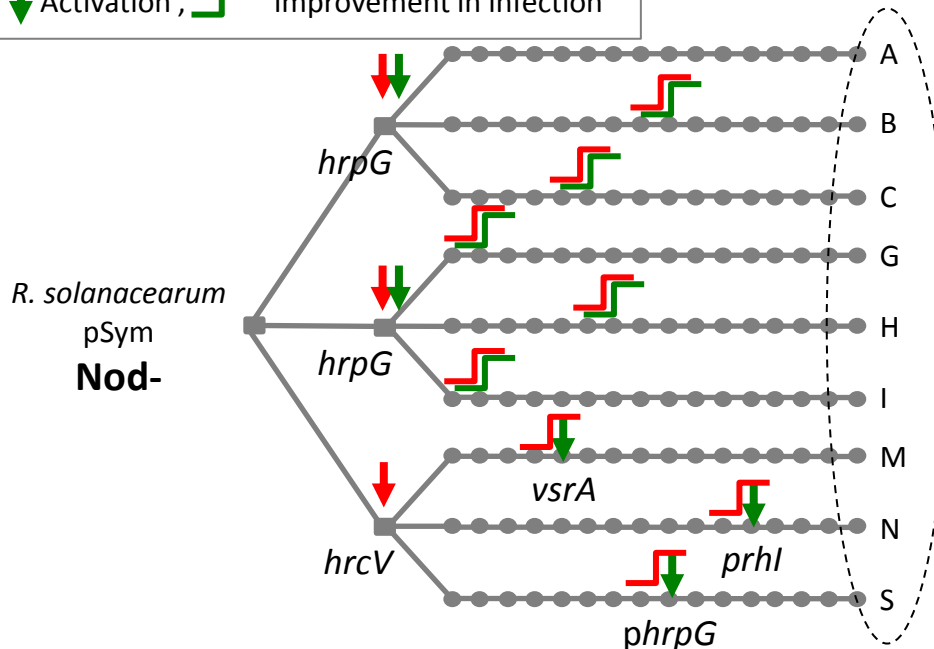


Bacteroid persistence

Root hair entry and nodulation

Nodule cell infection

↓ Activation, improvement in nodulation
 ↓ Activation, improvement in infection



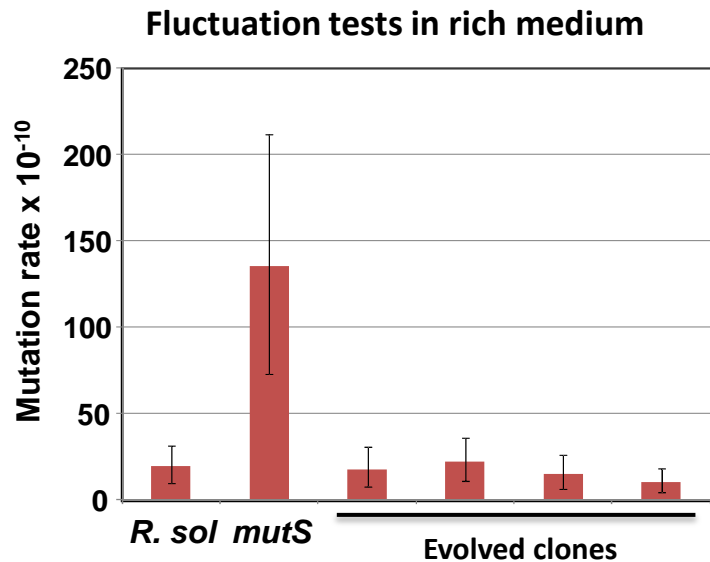
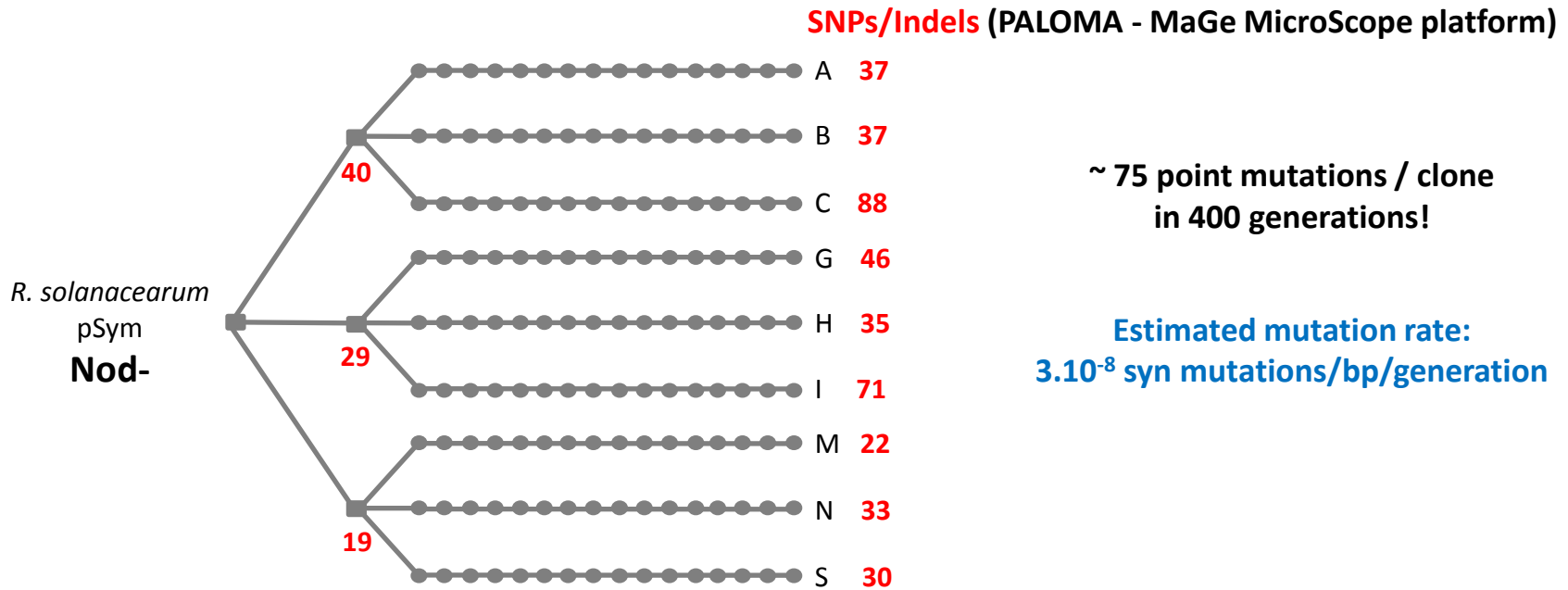
Evolution is sharp and fast: nodulation and infection were rapidly acquired and strongly improved

Evolution requires genome remodelling

Antagonism between symbiosis and elicitation of defense reactions

Regulators are preferred targets for evolution

Over-abundance of mutations in evolved clones

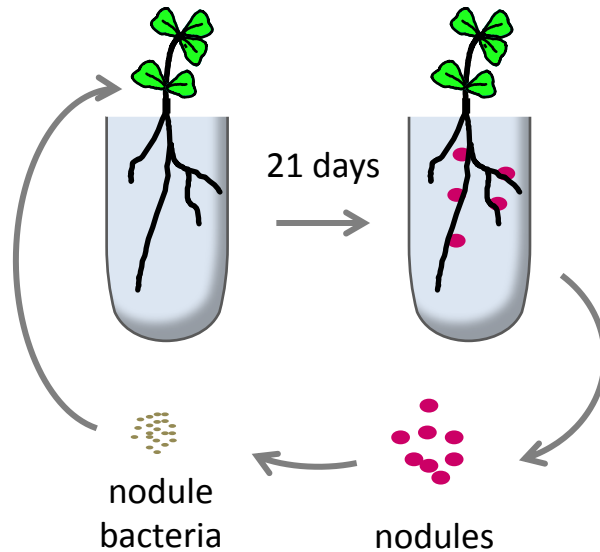


- ✓ Evolved clones have no mutation in DNA repair systems
- ✓ They are not constitutive hypermutators

Bacteria likely underwent transient hypermutagenesis during the experiment

Where do bacteria accumulate mutations?

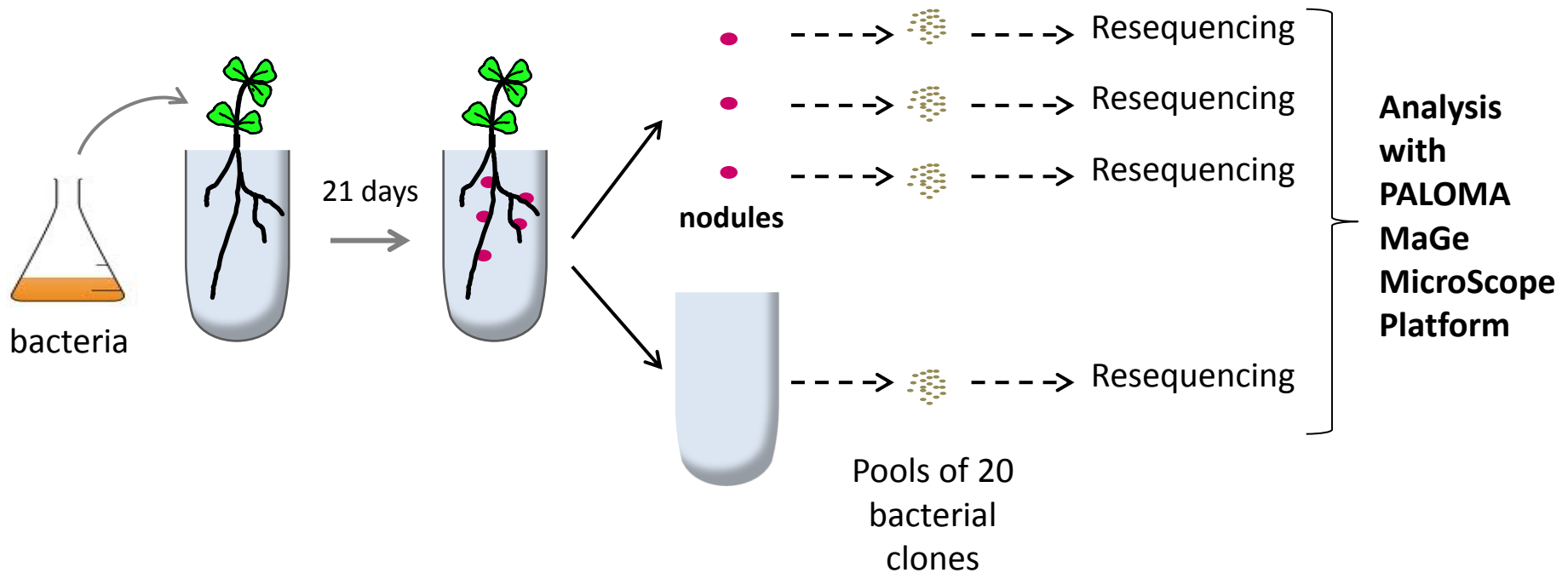
Bacteria were submitted to serial *ex planta* - *in planta* evolution cycle:



1. Colonization of the plant medium (C/N-free medium)
2. Multiplication within nodules

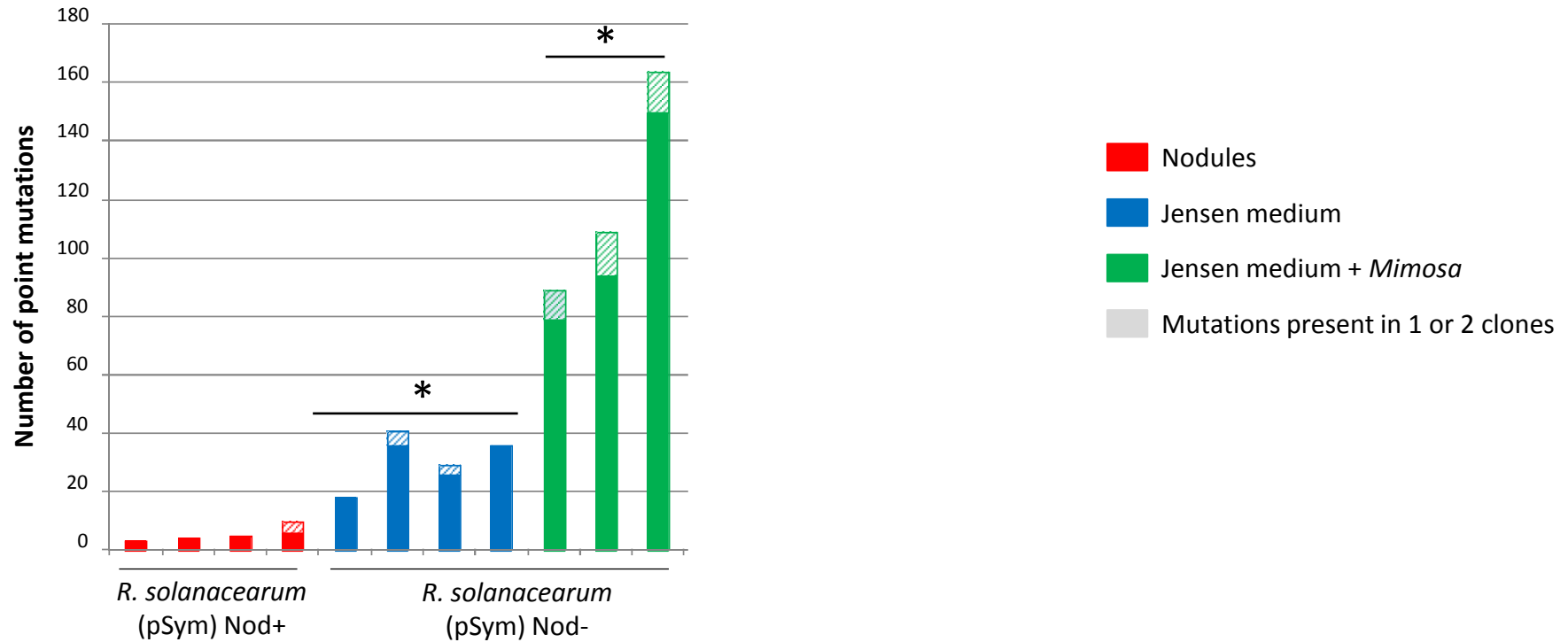
Where do bacteria accumulate mutations?

Replaying one cycle of evolution and resequencing populations to analyse the genomic diversity



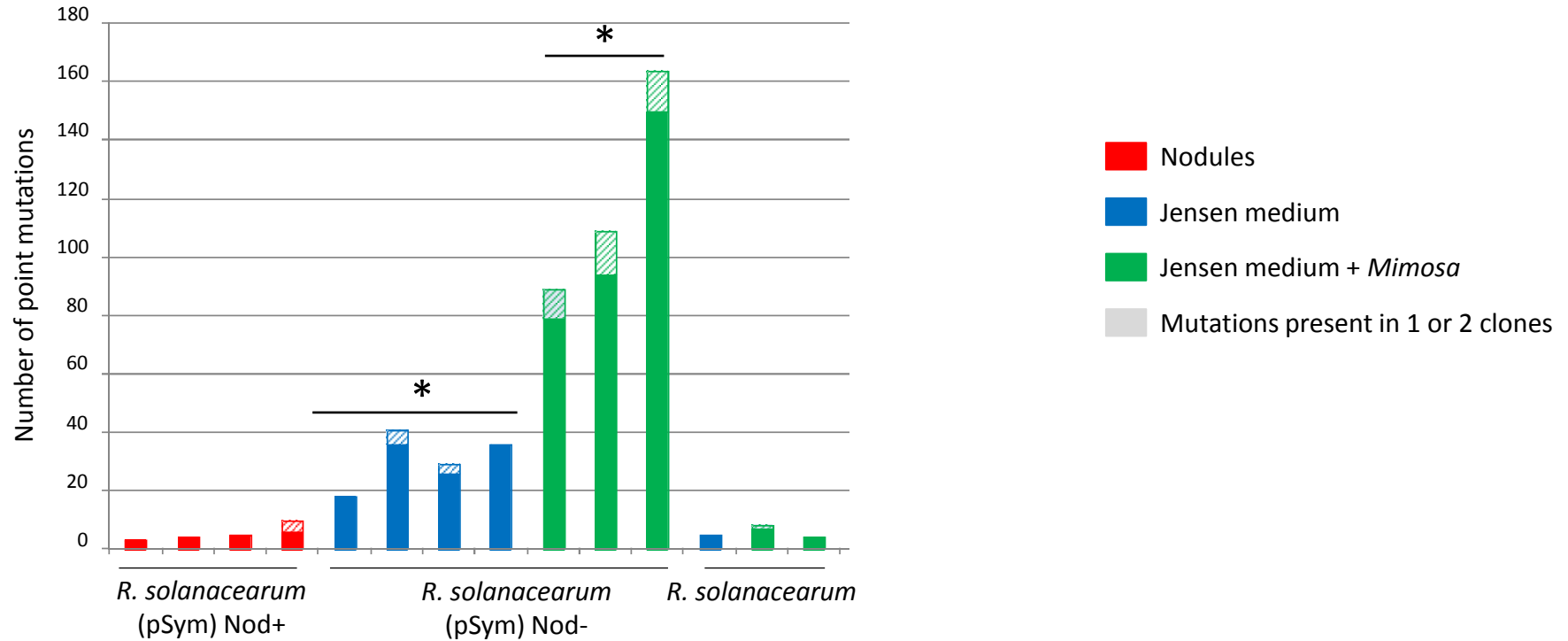
Transient mutagenesis occurs in the rhizosphere (*ex planta*)

Number of detected SNPs/Indel in pools of 20 clones by whole-genome sequencing



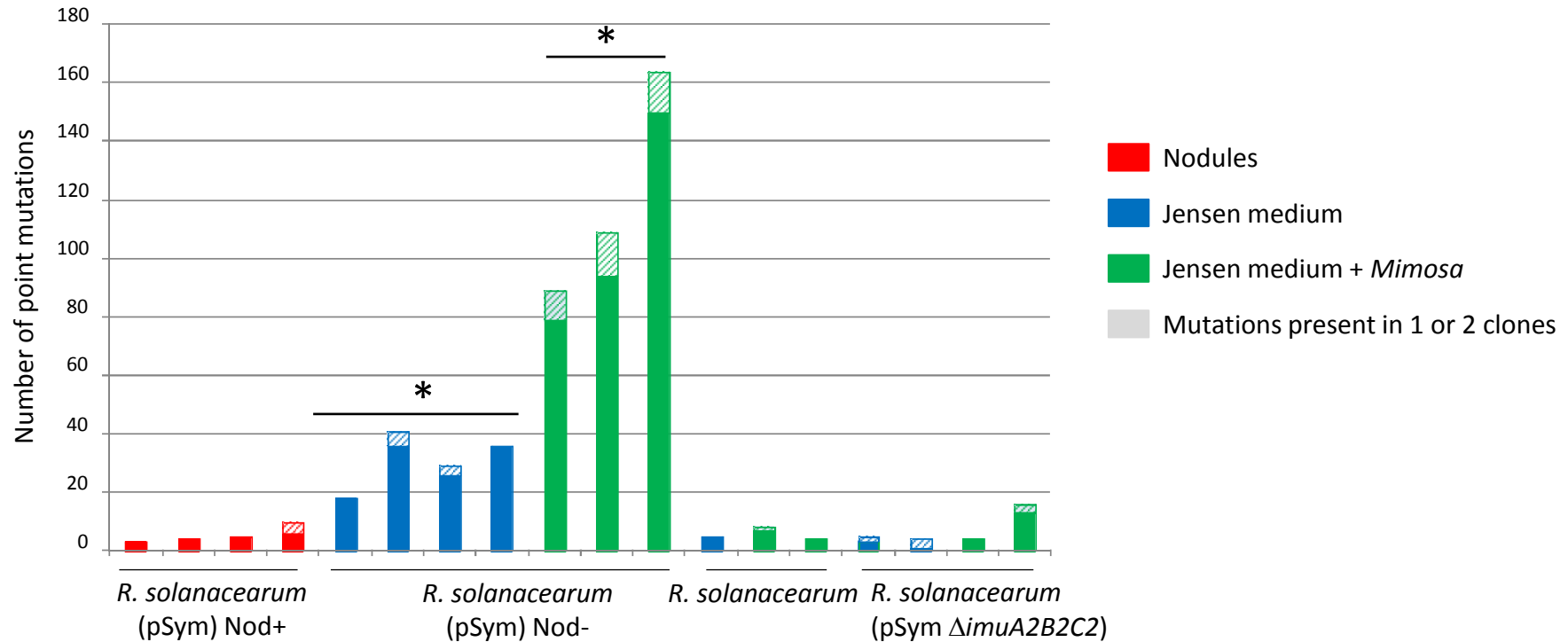
Transient mutagenesis is dependent on the symbiotic plasmid

Number of detected SNPs/Indel in pools of 20 clones by whole-genome sequencing



Plasmid error-prone DNA polymerases trigger hypermutability of the recipient genome

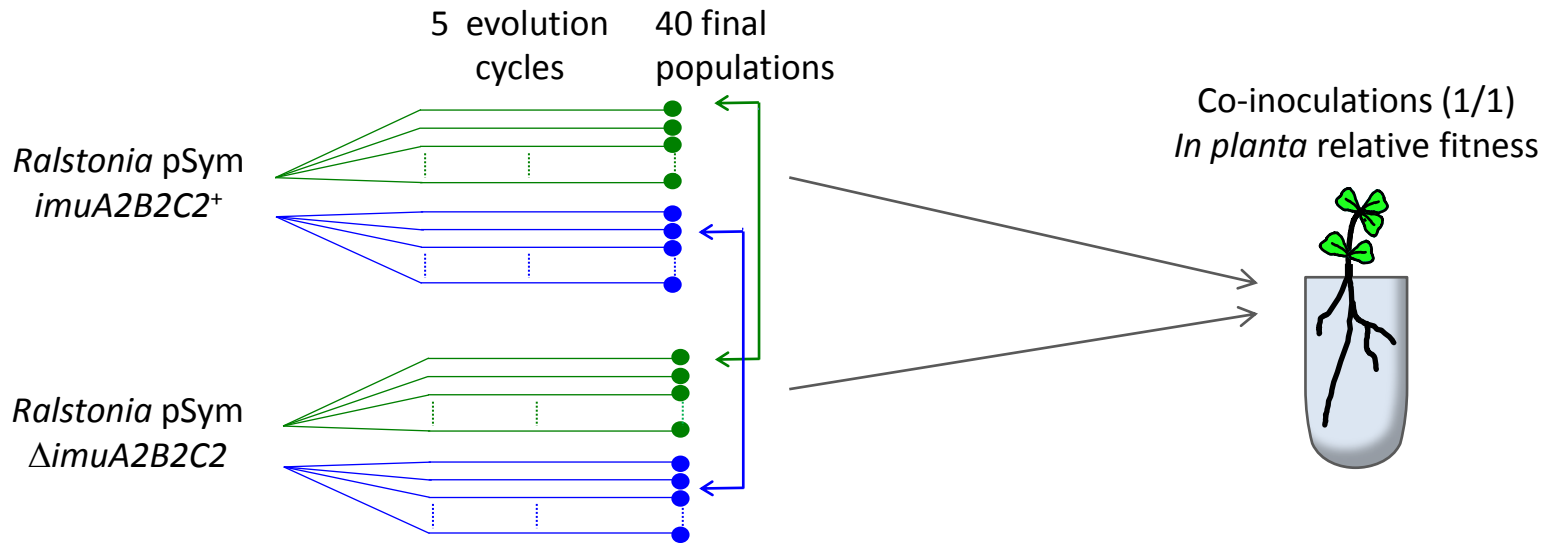
Number of detected SNPs/Indel in pools of 20 clones by whole-genome sequencing



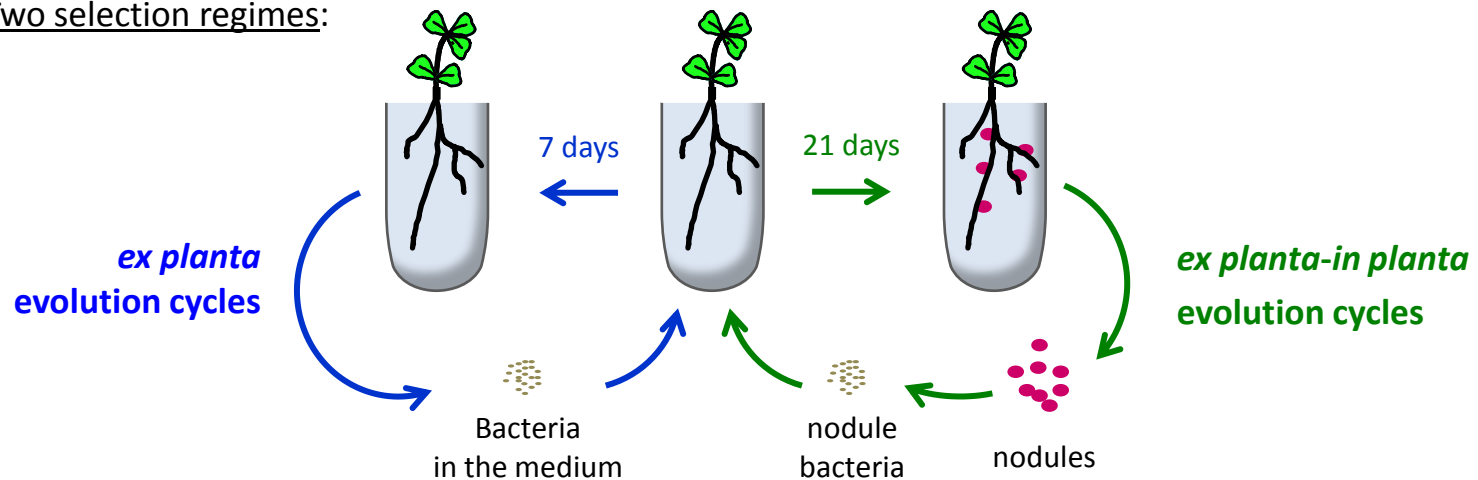
Mutagenesis cassette:



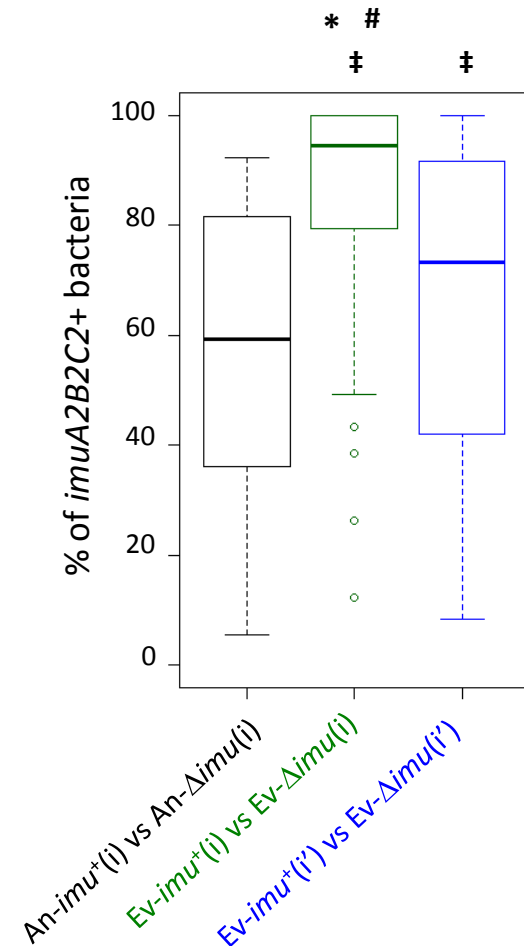
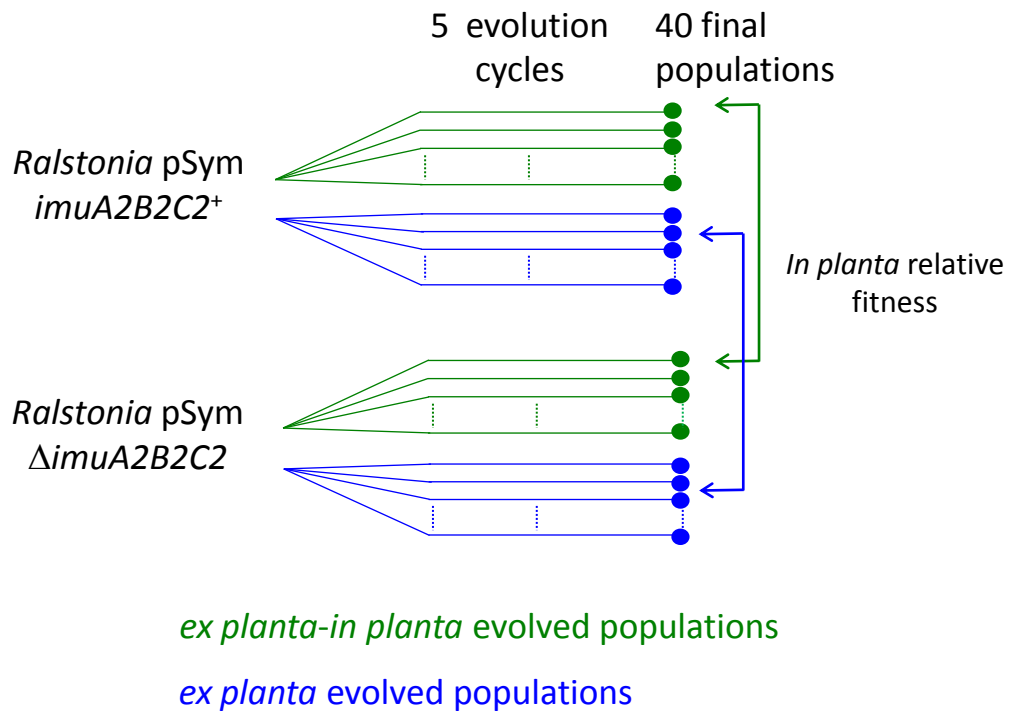
How hypermutability impact on symbiotic evolution?



Two selection regimes:



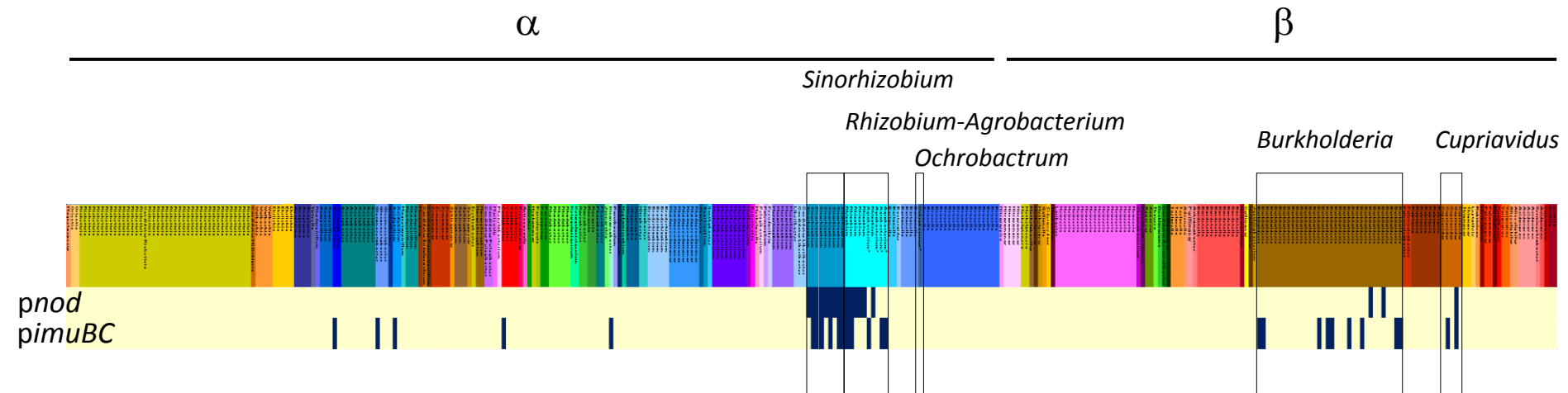
How hypermutability impact on symbiotic evolution?



***imuA2B2C2*-dependent mutagenesis accelerates the symbiotic evolution of *Ralstonia* under *Mimosa* selection pressure**

Phylogenetic distribution of the plasmid *imuBC* cassettes

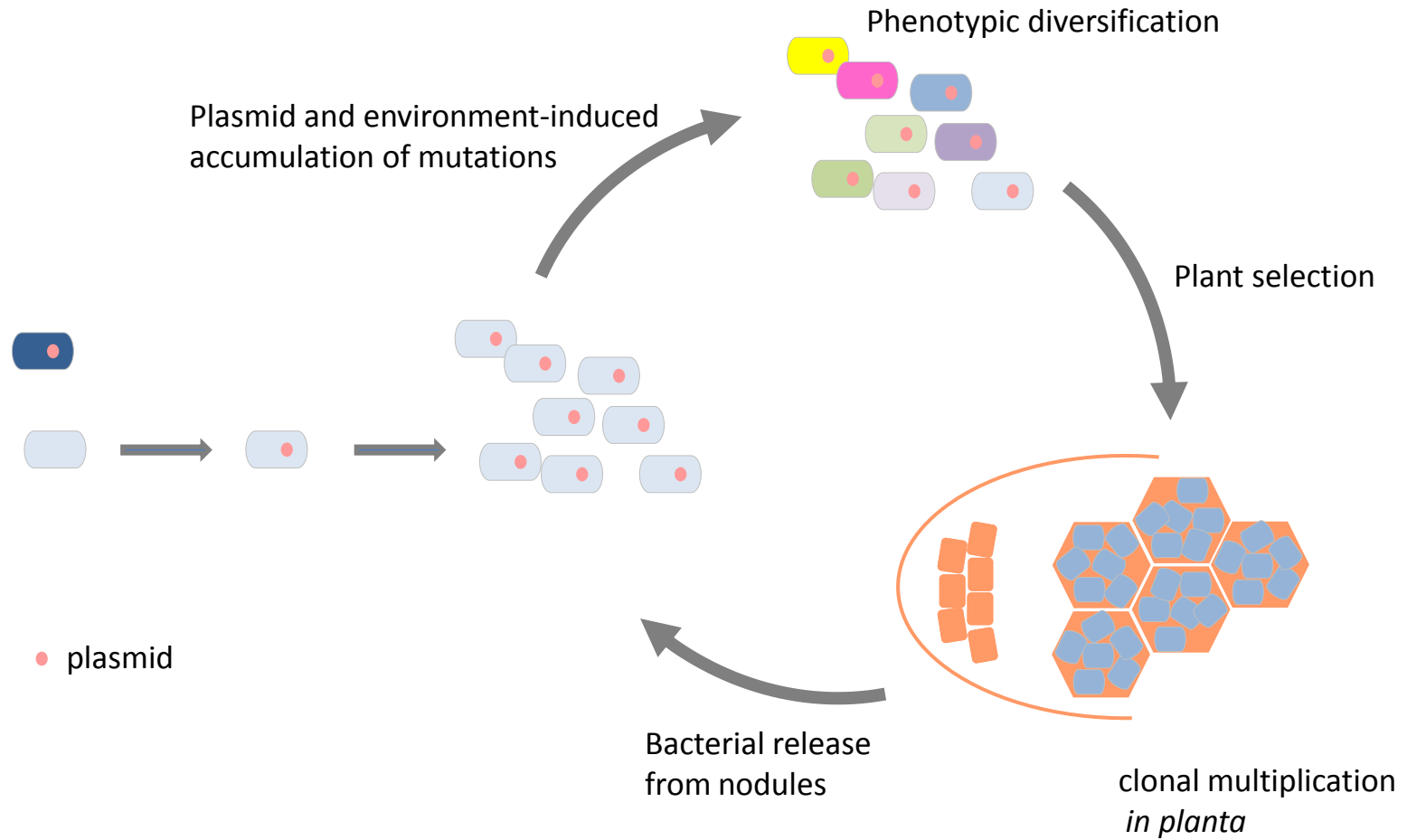
Among the 349 available genomes (109 genera) of α and β -proteobacteria, more than half possess a chromosomal *imuBC* cassette but only 28 have a plasmid cassette.



- 82% of all plasmid cassettes fall in the six genera that contain pSym-rhizobia
- 45% of the pSym possess a cassette

Plasmid *imuBC* cassettes may have favored dissemination of symbiotic genes within these taxa

Model for evolution of rhizobia



Conclusions

- We confirmed the remodelling of the recipient genome to express the symbiotic potential and adapt to legume symbiosis
- In some cases this process could have been accelerated by the co-transfer of mutagenic genes with symbiotic genes
- In our experiment, error-prone DNA polymerases encoded by the *C. taiwanensis* symbiotic plasmid trigger hypermutagenesis of the recipient genome, before bacteria enter the root
- This burst in genetic diversity has accelerated symbiotic evolution of *Ralstonia*
- ImuABC mutagenic cassettes are overrepresented in rhizobial lineages, supporting their role in rhizobium evolution

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