

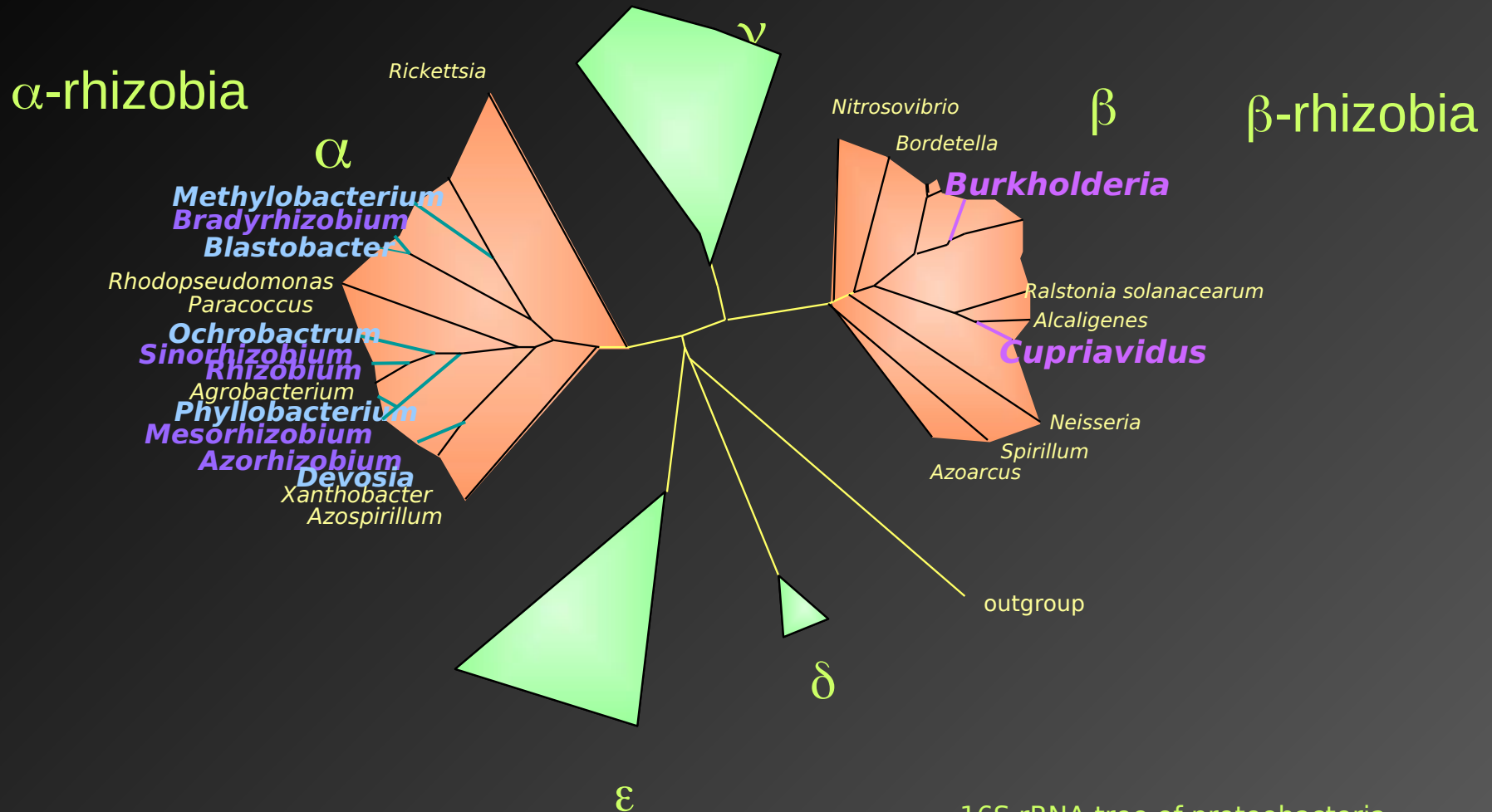
Betaproteobacterial symbioses with legumes

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diversity of rhizobia

“rhizobia” can be a functional term to define bacteria able to fix nitrogen in symbiosis with legumes



16S rRNA tree of proteobacteria
(adapted from Moulin et al. 2001)

Beta-rhizobia: facts & questions

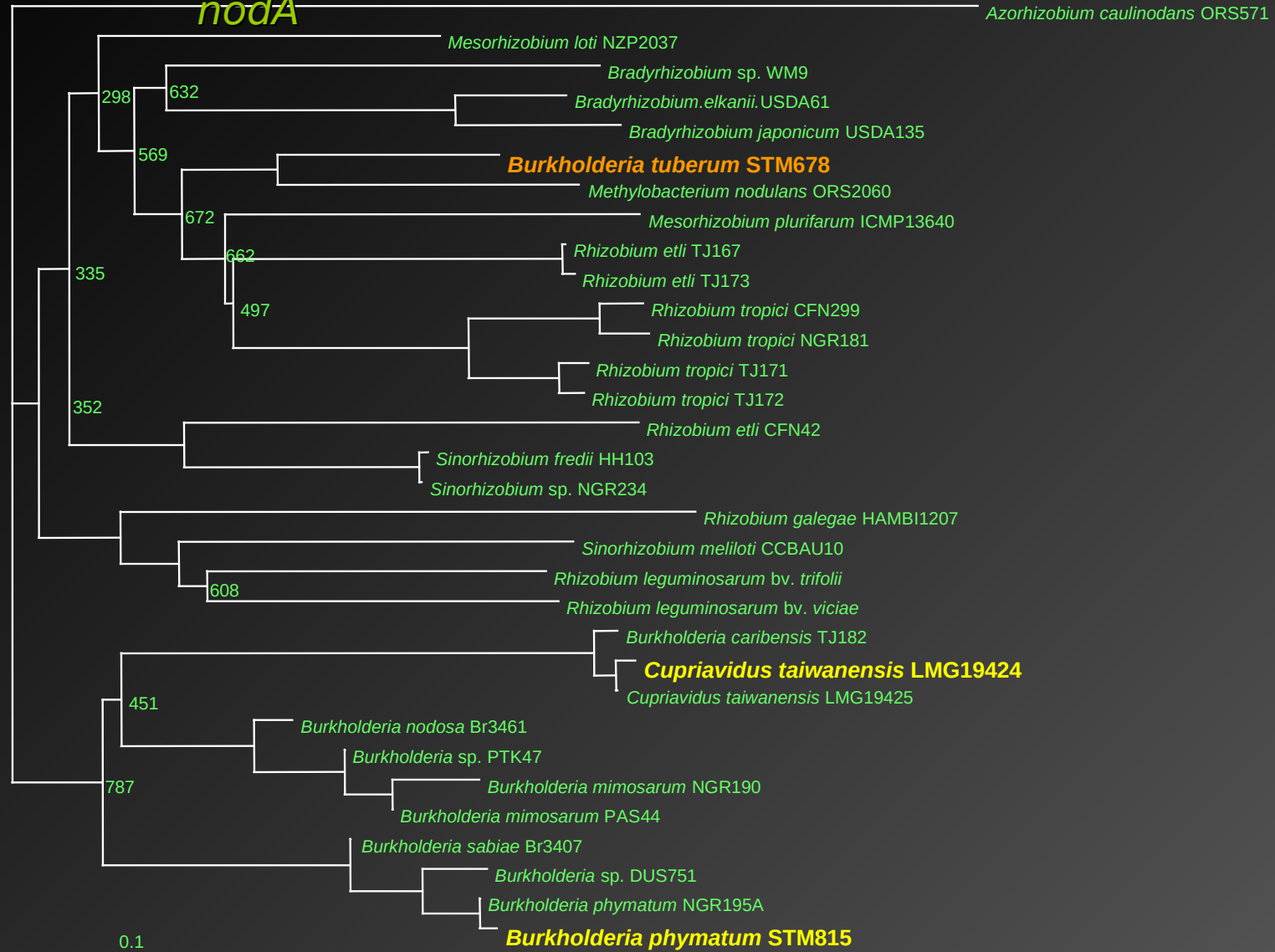
- FACTS:
- So far comprise *Cupriavidus* (*Ralstonia*) and *Burkholderia* spp.
- Not related to *Rhizobium*, but are closely related to plant-associated and endophytic Betaproteobacteria, such as other N₂-fixing *Burkholderia* spp. and to plant growth-promoting bacteria, such as *Herbaspirillum*.
- QUESTIONS:
- Are they genuine symbionts?
- From where do they originate?
- Are they widespread and important to ecology?
- Is there any taxonomic relationship between them and their legume hosts?

Are they genuine symbionts?

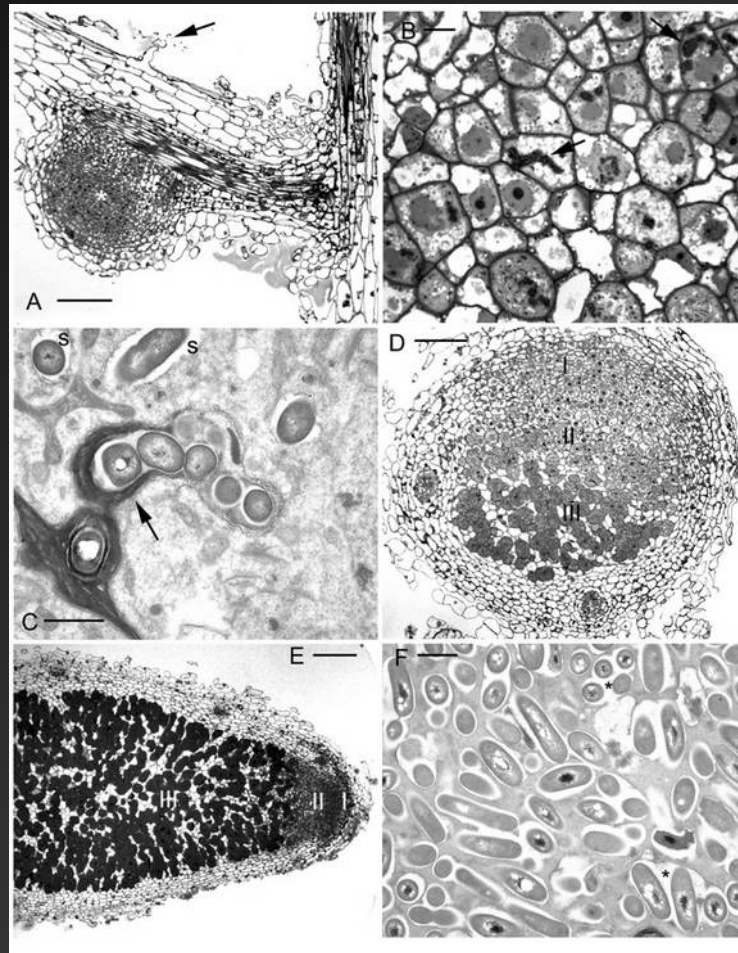
- Need to demonstrate a genetical predisposition to nodulate and fix N_2 (i.e. *nod* and *nif* genes)
- Need to demonstrate an actual ability to nodulate and fix N_2 to the benefit of their legume hosts

Beta-rhizobia from *Mimosa* spp. have *nod* and *nif* genes that cluster together e.g.

nodA

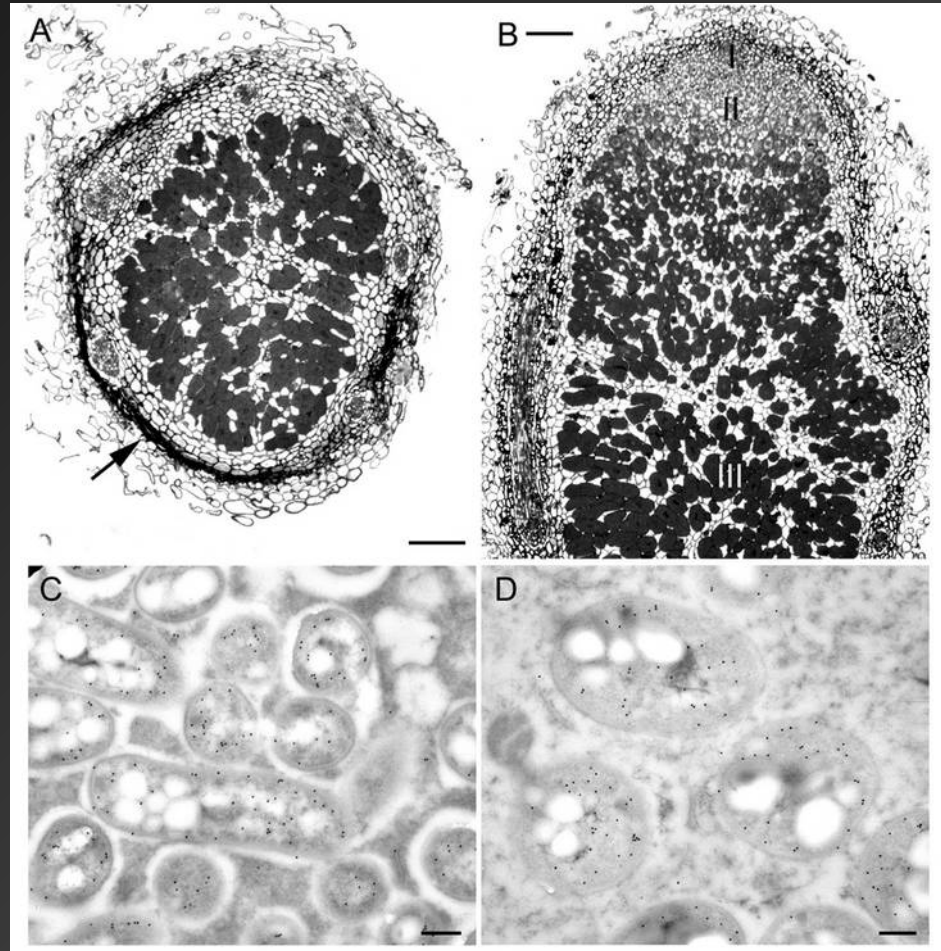
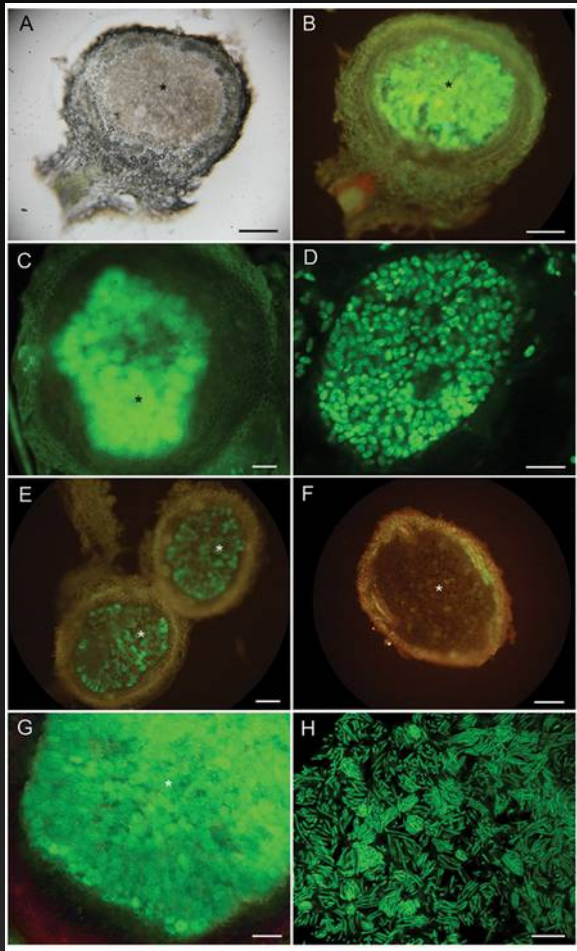


Mimosa pudica nodules formed by *C. taiwanensis* are fully functional symbiotic organs



First confirmation of symbiotic nodulation by Beta-rhizobia: Chen et al. 2003 MPMI **16**:1051-1061.

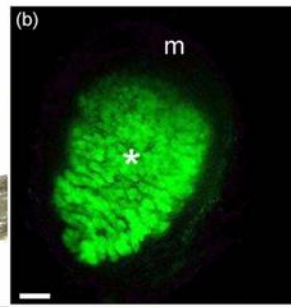
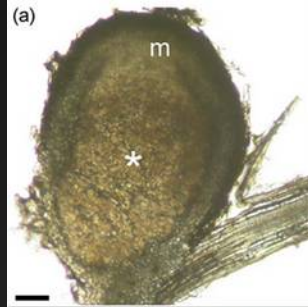
Mimosa nodules formed by *Burkholderia mimosarum* and *B. nodosa* are also fully functional



Chen et al. 2005a *AEM* **71**: 7461-7471; Chen et al. 2005b *New Phytol.* **168**: 661-675.

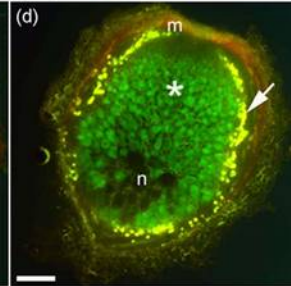
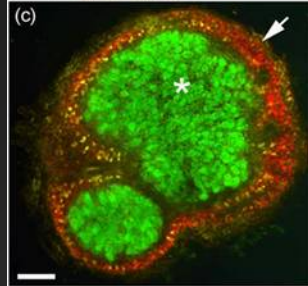
Burkholderia phymatum STM815 is also an effective symbiont of *Mimosa* spp.

M. pudica
+ Bp-GFP



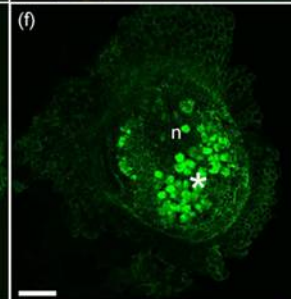
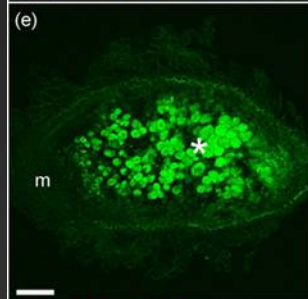
M. pudica
+ Bp-GFP

M. himalayana
+ Bp-GFP



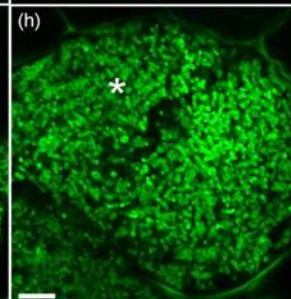
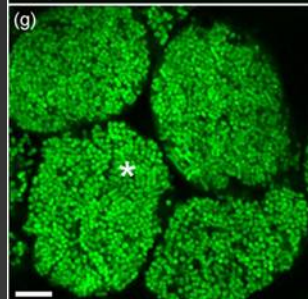
M. himalayana
+ Ct-GFP

M. latispinosa
+ Bp-GFP



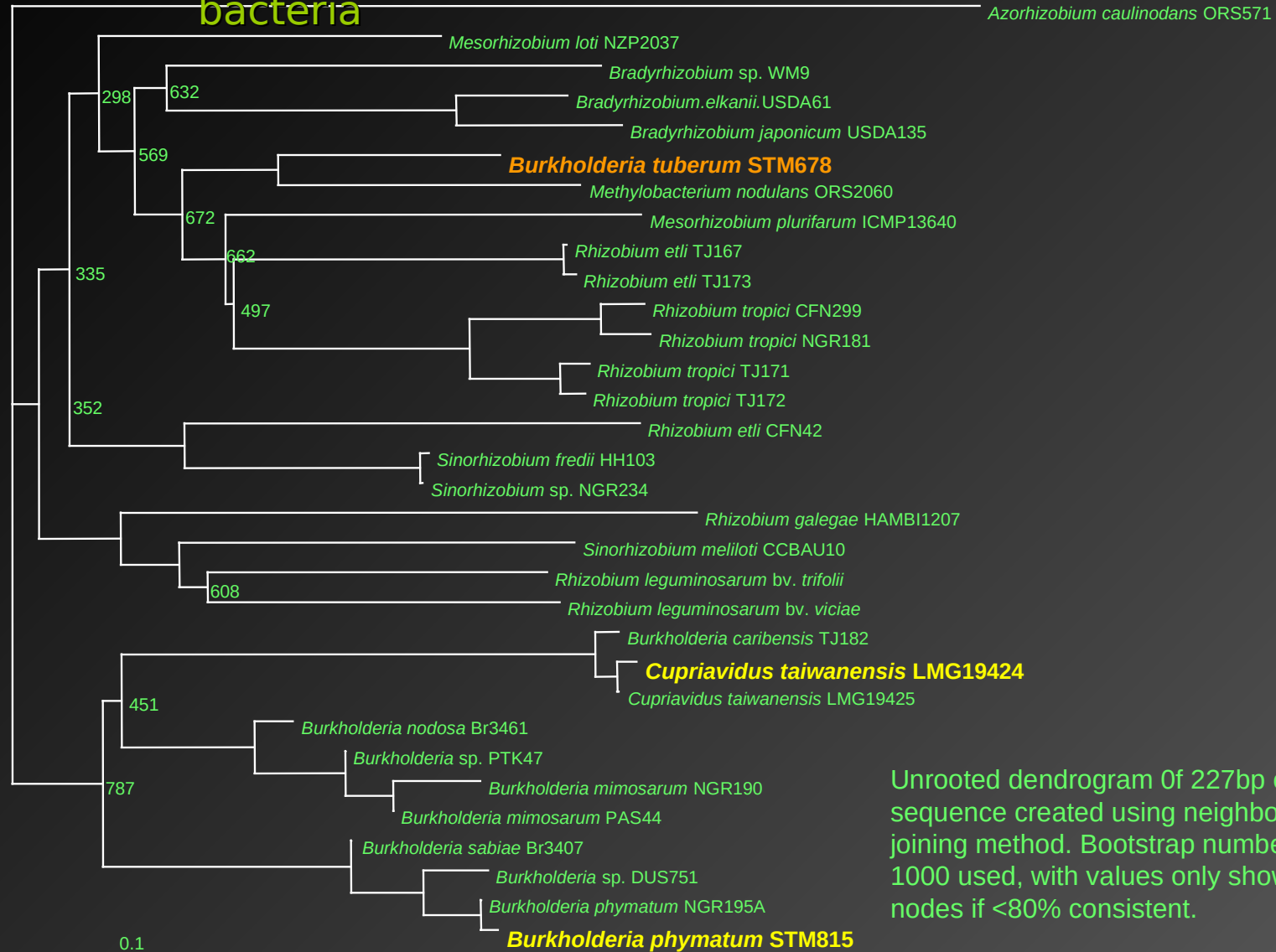
M. latispinosa
+ Ct-GFP

M. latispinosa
+ Bp-GFP



M. latispinosa
+ Ct-GFP

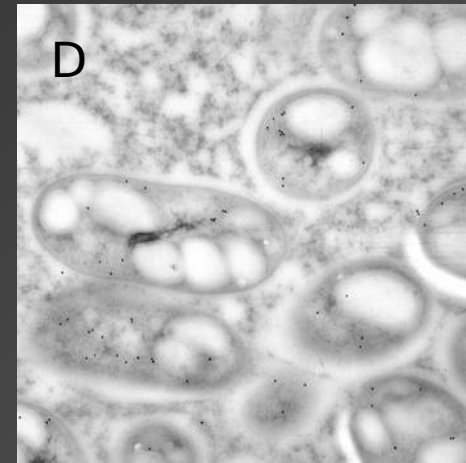
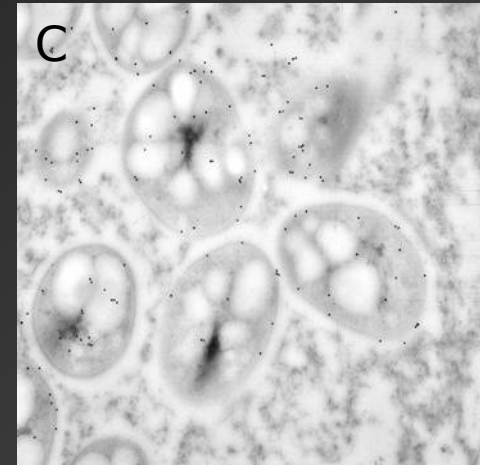
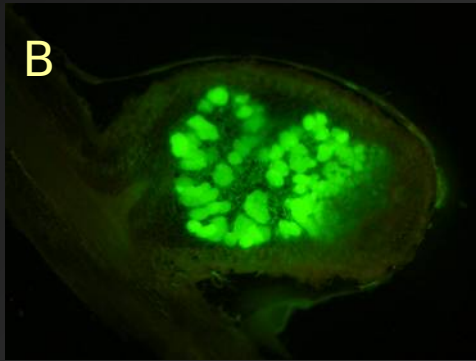
The *nodA* gene of *B. tuberum* is distant from *Mimosa*-nodulating bacteria



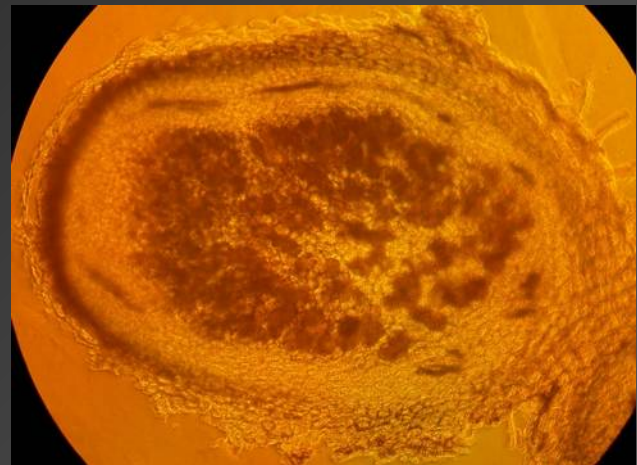
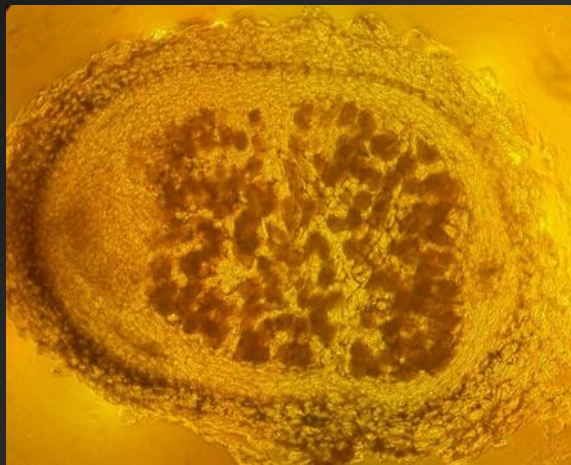
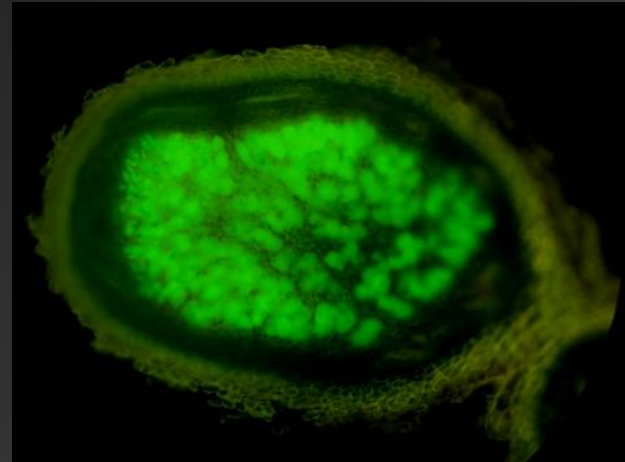
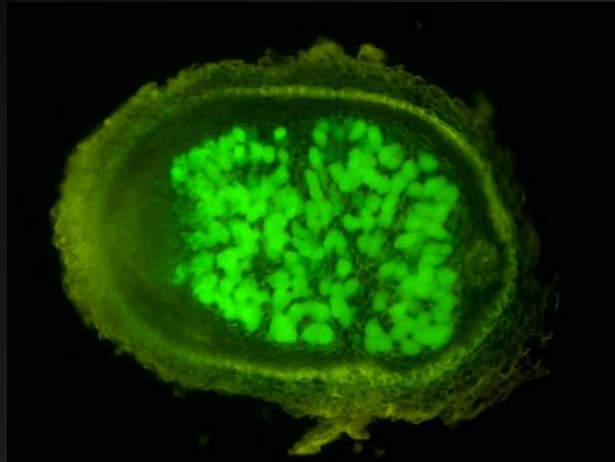
B. tuberum STM678 effectively nodulates the South African Papilionoid legume *Cyclopia*

Cyclopia genistoides
+ *B. tuberum* STM678GFP (A, B)

Cyclopia bacteroids labelled with antibodies against *Burkholderia* (C) and *nifH* protein (D)



B. tuberosum STM678 nodulates other
legumes in the tribe Podalyreae
(Papilionoideae)



Podalyria canescens

Virgilia oropoides

Does NOT nodulate *Mimosa* spp.

Beta-rhizobia are genuine symbionts:

- They possess *nod* and *nif* genes
- They can form functional nodules on their legume hosts, as demonstrated under sterile conditions with reporter gene (GFP)-tagged strains and/or specific antibodies

Where do they come from?

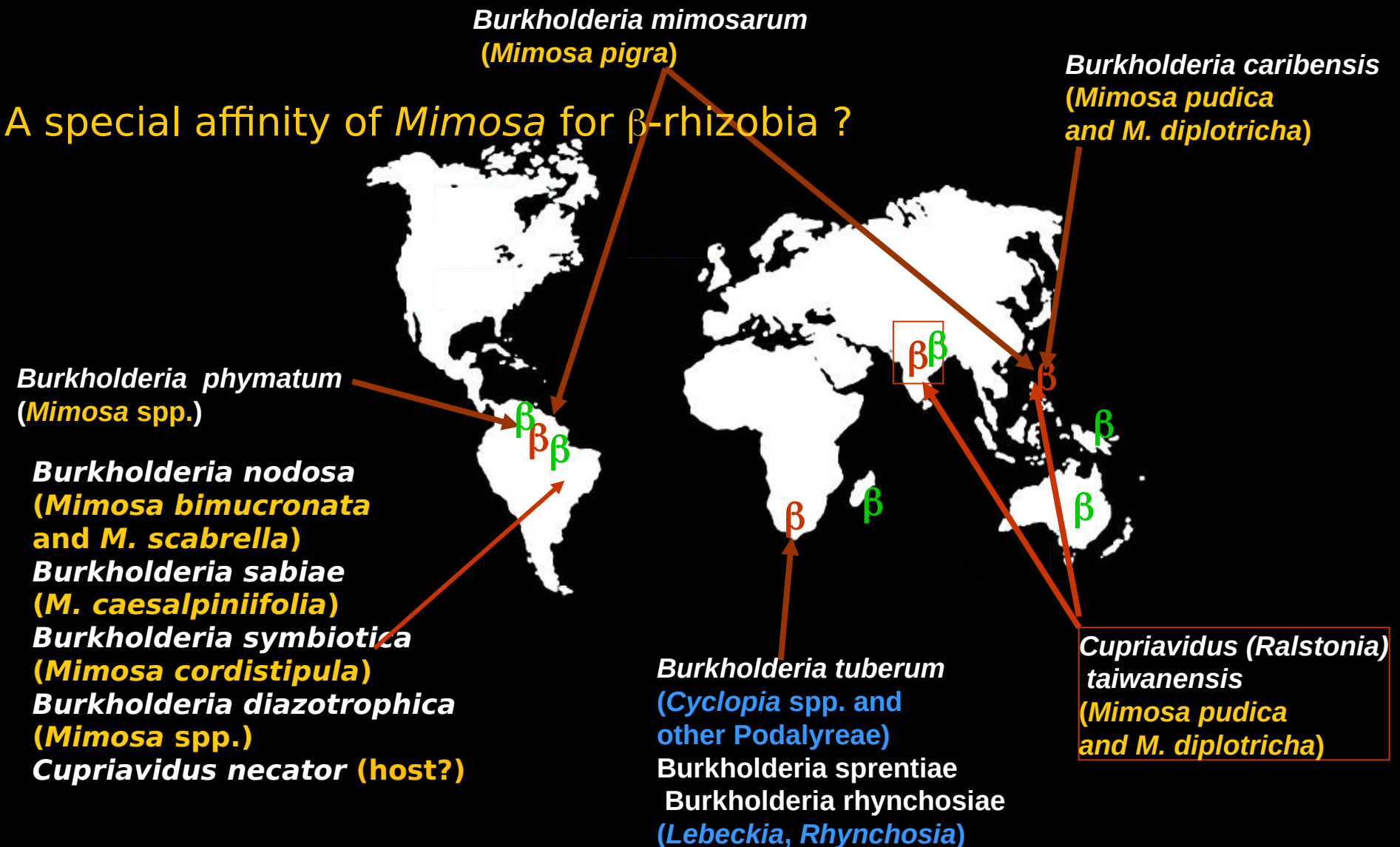
- In terms of geography
- In terms of evolution

Are they widespread and important to ecology?

-rhizobia are widespread

13 characterised species of β -rhizobia and many other isolates of β -rhizobia

A special affinity of *Mimosa* for β -rhizobia ?



The large legume genus *Mimosa*

Mimosa species can be herbs but most are shrubs or trees from tropical or subtropical regions

Mimosa is one of the largest genera of Mimosoid legumes, with approximately 500 species

About 450 *Mimosa* species are native to Central and South America



Cerrado and Caatinga regions of Central Brazil are the major centres of diversification



Collecting Brazilian *Mimosa* symbionts



Chapada dos Guimaraes



Wetland - Pantanal

The Cerrado and Caatinga environments



Chapada de Diamantina



Pantanal wetlands



Flooded Caatinga



Chapada de Diamantina sands

Collecting Brazilian *Mimosa* symbionts

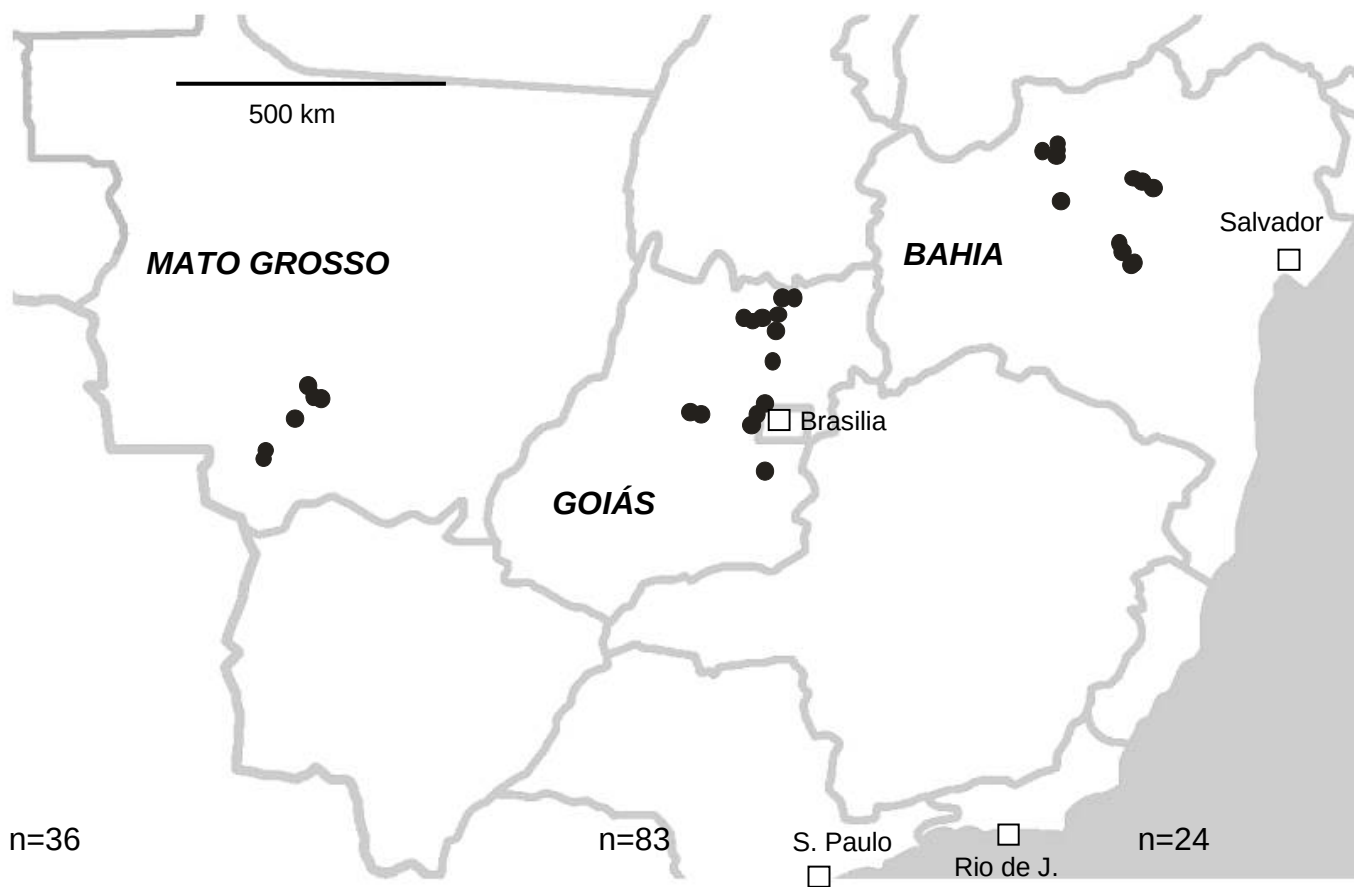
143 isolates from 49
different *Mimosa*
species

M. pyreneia (endemic)

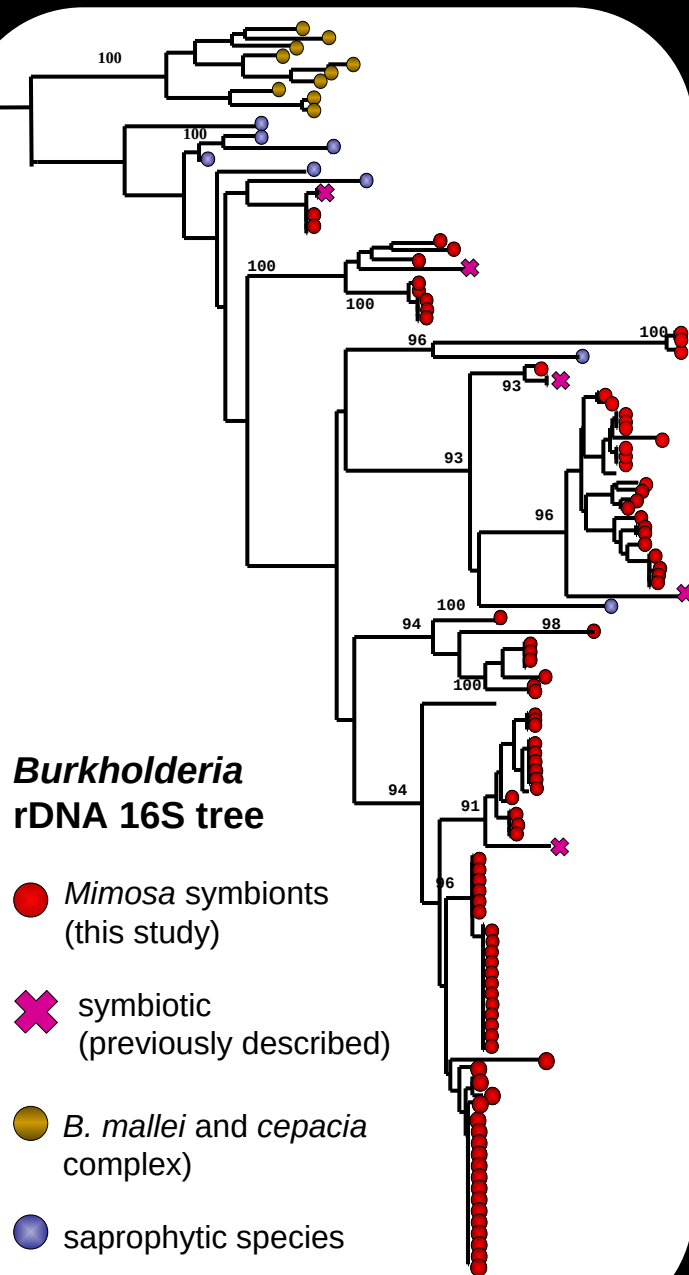
M. albolanata (restricted to Cerrado)

M. pigra (worldwide distribution)

M. hirsutissima
(widespread in S. America)



What do housekeeping genes tell us?



97% of *Mimosa* symbionts in Brazil belong to the *Burkholderia* genus

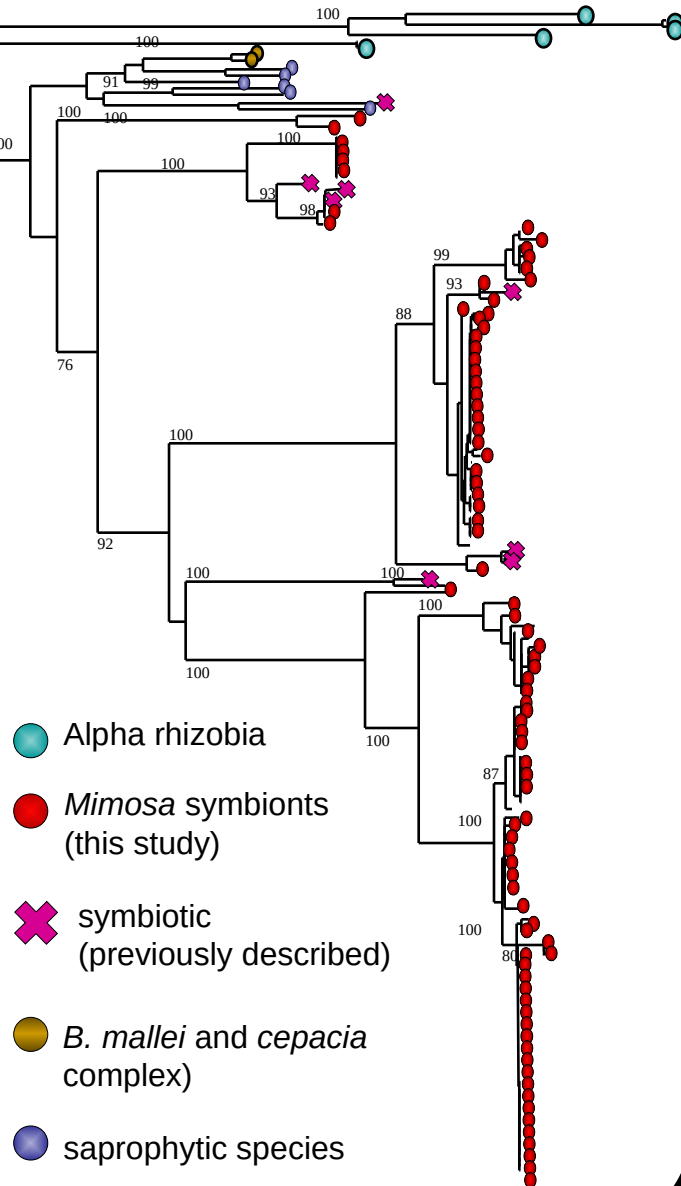
Burkholderia are the main symbionts of *Mimosa*

16S and *recA* tree topologies are congruent and indicate that Brazilian *Burkholderia*

- are close to already described symbiotic *Burkholderia*
- may constitute several new species
- are closer to saprophytic and generally diazotrophic *Burkholderia* rather than to pathogenic *Burkholderia*

What do nitrogen fixation genes tell us?

nifH tree



The diazotrophic character is present in many non-symbiotic *Burkholderia*

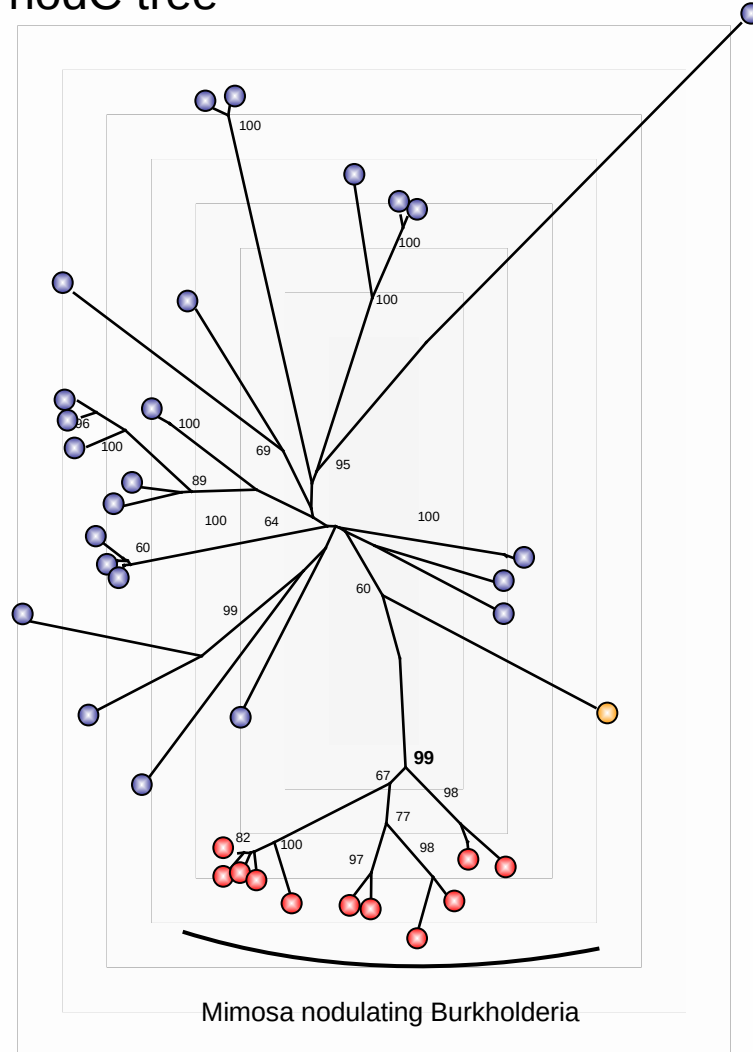
Symbiotic *Burkholderia nifH* genes are related to those of free-living burkholderias

Suggests that symbiotic *Burkholderia* didn't acquire their nitrogen fixing capabilities from α -rhizobia, but added nodulation to pre-existing N-fixing capabilities

In contrast to most α -rhizobia, some of them are known to be free living diazotrophs

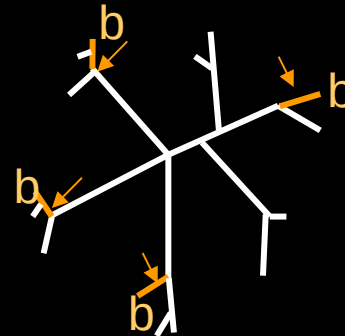
What do nodulation genes tell us?

nodC tree

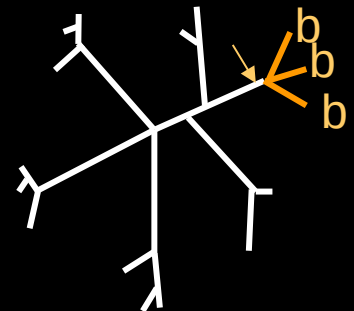


- *nodC* genes of *Burkholderia* associated with *Mimosa* form a monophyletic group

Suggests a unique acquisition of symbiotic genes



Several transfers
from different sources



unique transfer

Cerrado & Caatinga spp. grow in low pH soils (>3.0) and prefer acid-tolerant *Burkholderia*



Nodules from 67 *Mimosa* spp. gave a positive signal with an antibody against *Burkholderia*, and no signal with an antibody against *Cupriavidus*



M. ursina inoculated with *Burkholderia* (left) or *Cupriavidus* (right)

Summary – The Betas from Brazil

- *Burkholderia* species are the natural symbionts of native *Mimosa* species in Brazil.
- Symbionts are found in 7 distinct species complexes.
- Symbiosis-related genes have the same phylogeny as core genes – HGT is rare.
- Ancient symbionts (>50 million years)
- Species differ in environmental preference, not host specificity.
- Endemic Cerrado/Caatinga spp. cannot form effective symbioses with other bacteria.

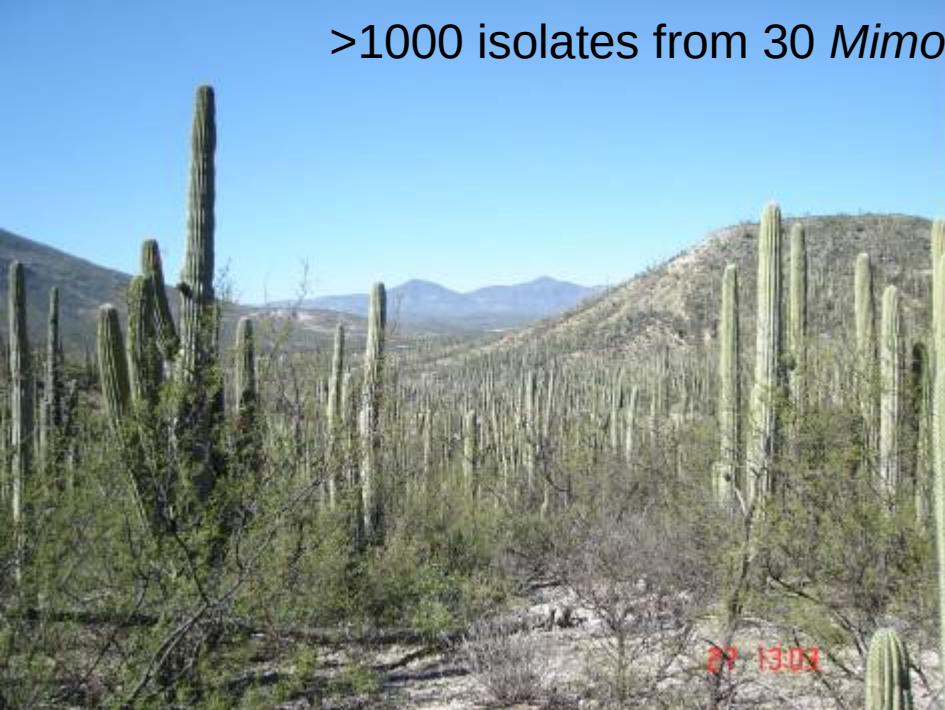
Mimosa spp. are nodulated
exclusively by Beta-rhizobia

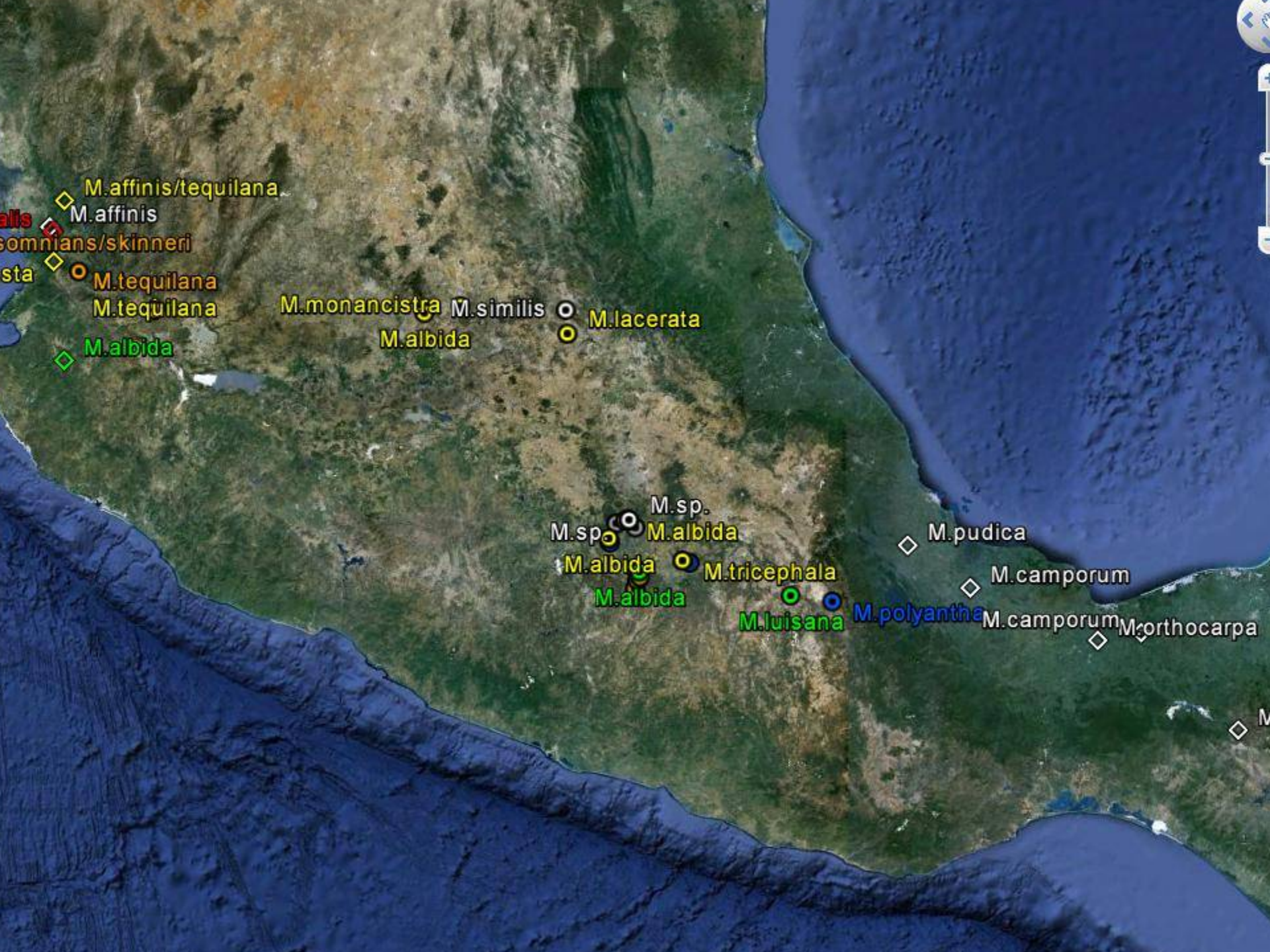
Or are they?

Outside Brazil.....

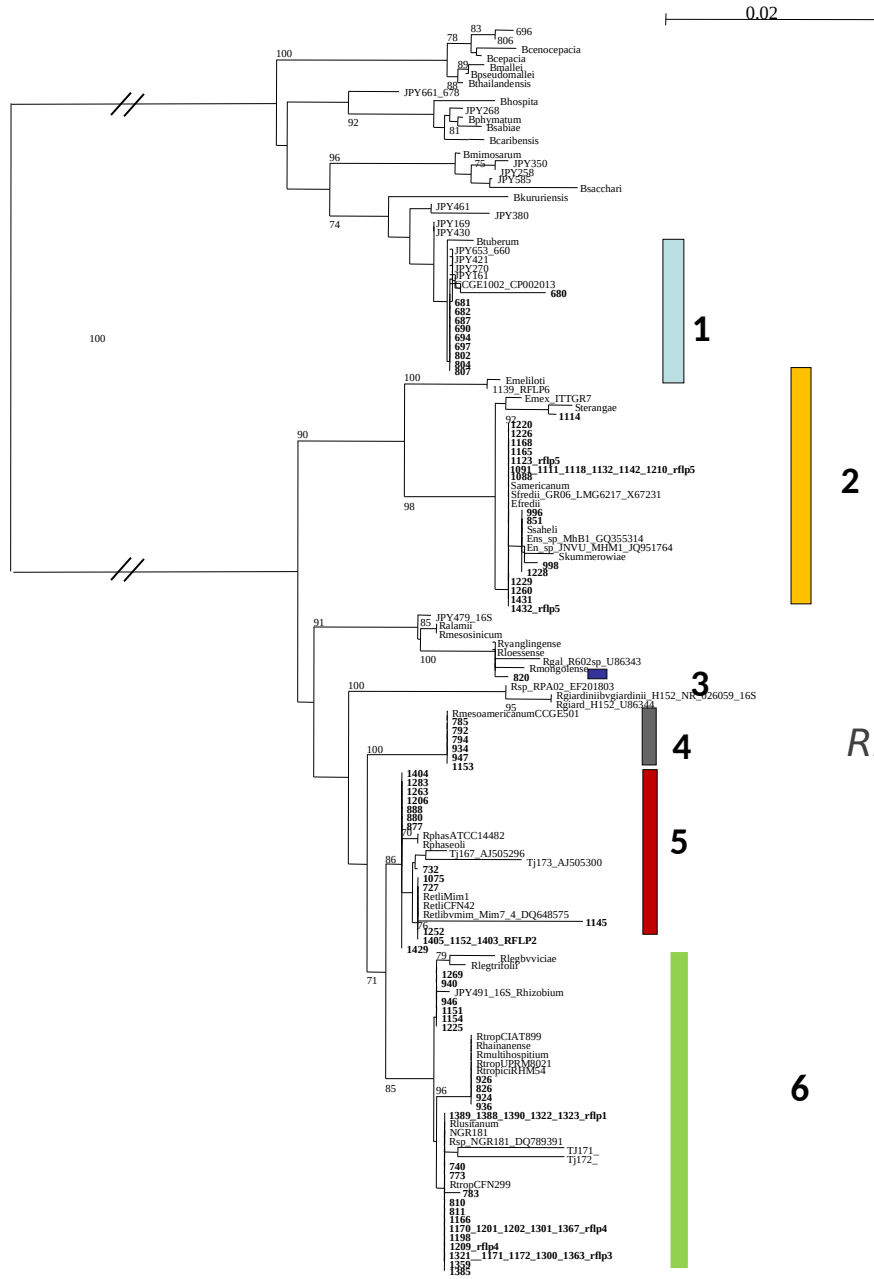
Mexico is the most important centre of *Mimosa* diversity in the Northern hemisphere (>100 spp.), so what are the symbionts of Mexican *Mimosa* spp.?

>1000 isolates from 30 *Mimosa* spp.





16S rRNA



Burkholderia

Ensifer (Sinorhizobium)

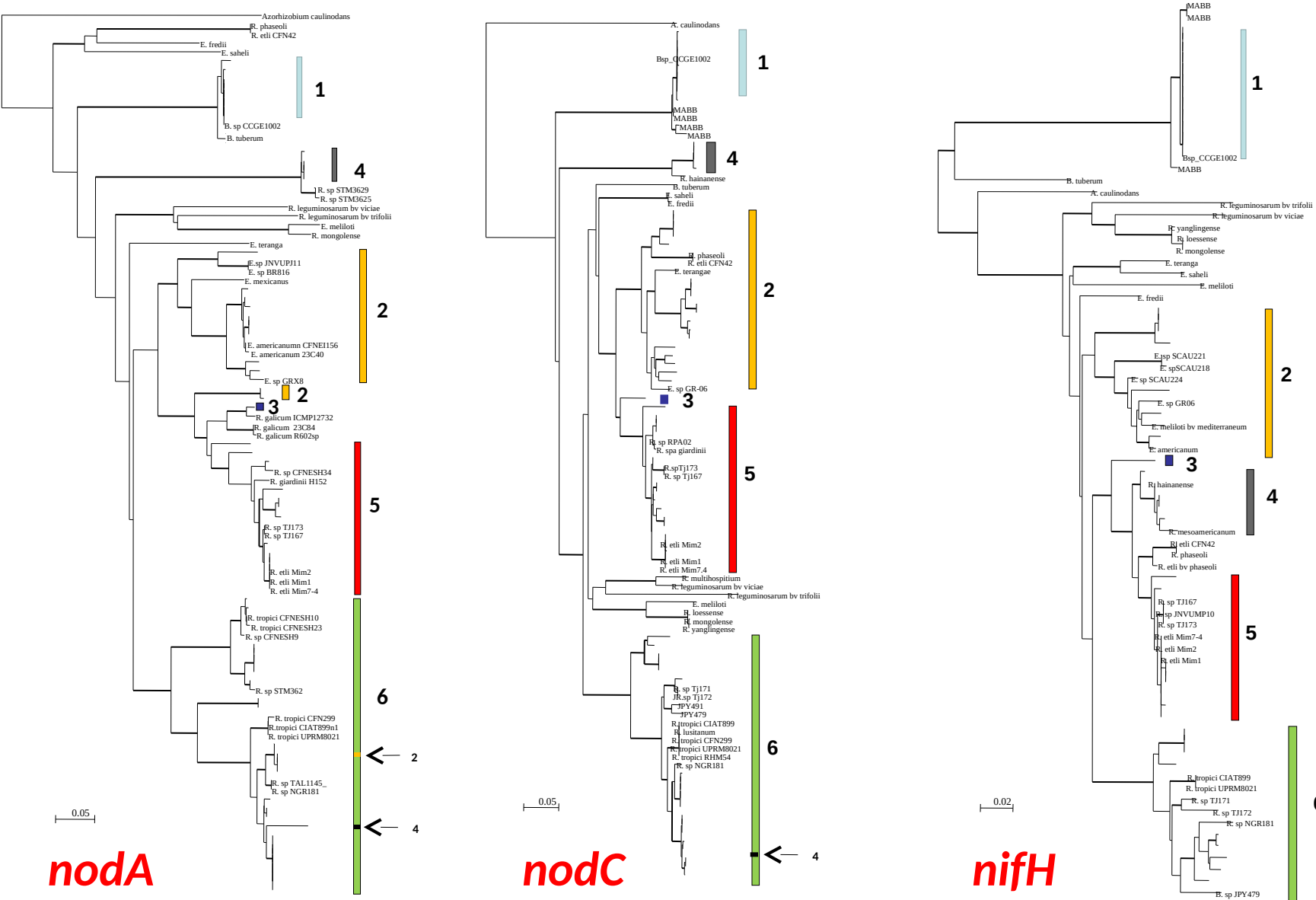
Rhizobium mesoamericanum

Rhizobium etli

Rhizobium tropici

16S

Symbiosis-related genes have the same phylogeny as the house-keeping genes



Mexican *Mimosa* symbionts

- 25 *Mimosa* spp. have been sampled from central Mexico as far south as Tehuacan (Puebla), and as far west as Tequila (Jalisco).
- The majority of the symbionts are either *Rhizobium* or *Sinorhizobium*; these are closely related to species already present in central Mexico, particularly in common bean (*Phaseolus vulgaris*).
- Mexican endemic *Mimosa* spp. cannot nodulate effectively with Beta-rhizobia, but will nodulate with native *Rhizobium* strains.
- “Widespread” Mexican species can nodulate with *Burkholderia*
- **Soils are neutral - alkaline.**

Other centres of *Mimosa* diversity

- Madagascar and East Africa: 30 spp.
- India: 4 – 6 spp.
- Uruguay: 12 spp.

“Old World” *Mimosa* spp. are not nodulated by Beta-rhizobia



M. hamata = *Sinorhizobium*

Does NOT nodulate effectively with Beta-rhizobia

Gehlot et al. (2013) Ann Bot **112**: 179-196



M. himalayana = *Sinorhizobium*

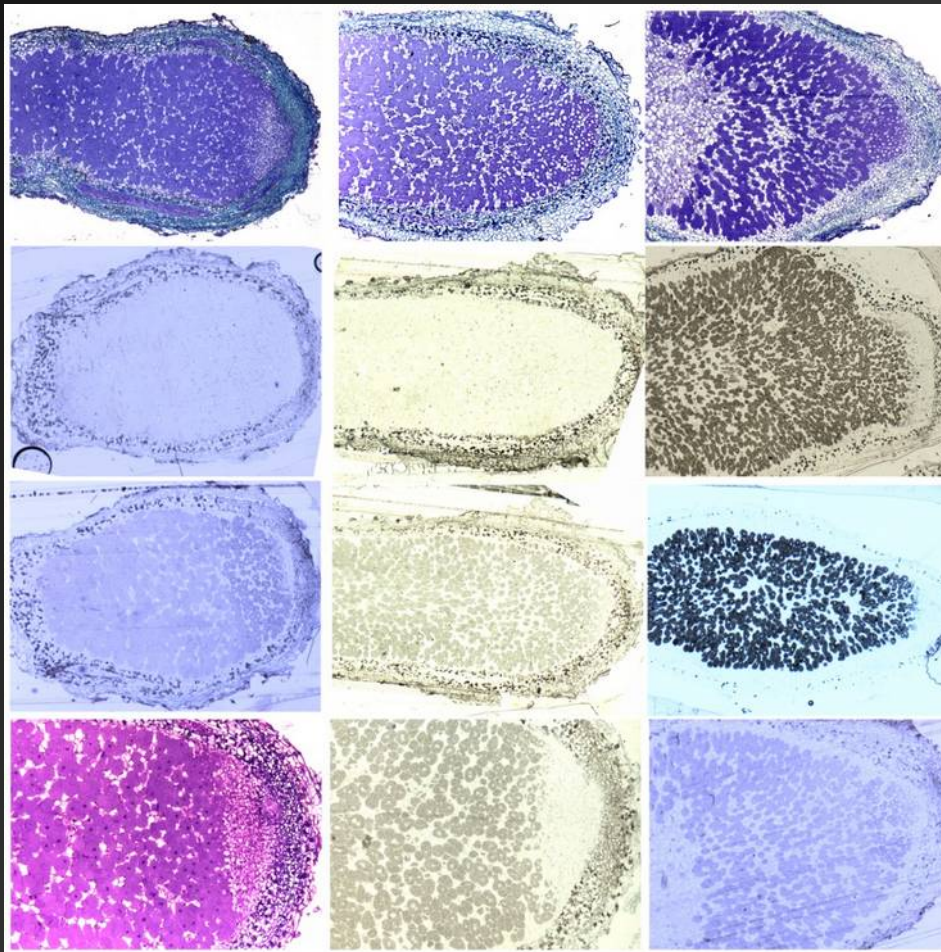
India *Mimosa* symbionts are (Old World) *Sinorhizobium* spp., but we know most nothing about Madagascan endemics

Southern *Mimosa* spp. are nodulated by *Cupriavidus*?

M. magentea

M. ramulosa

M. schledeneii

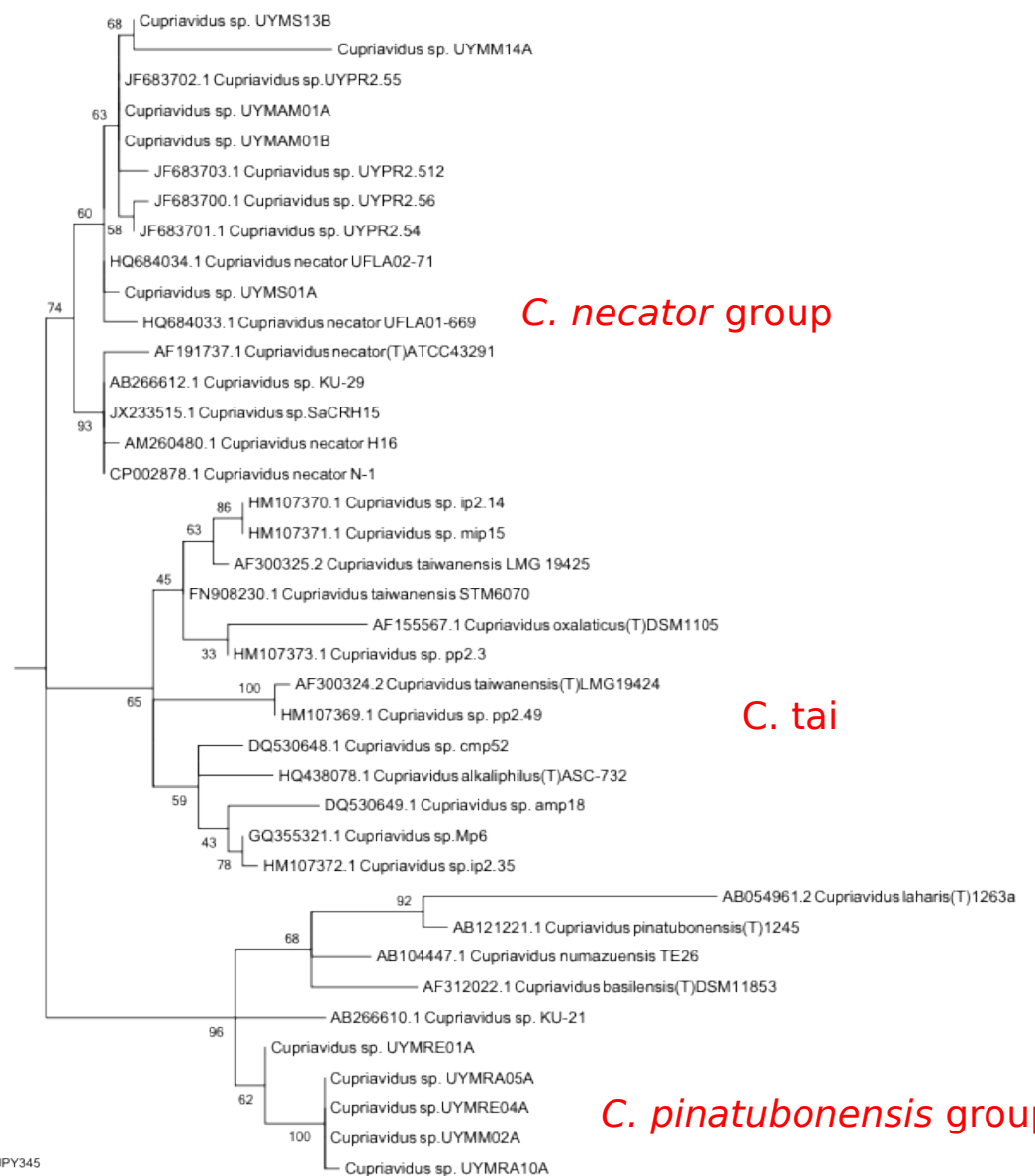
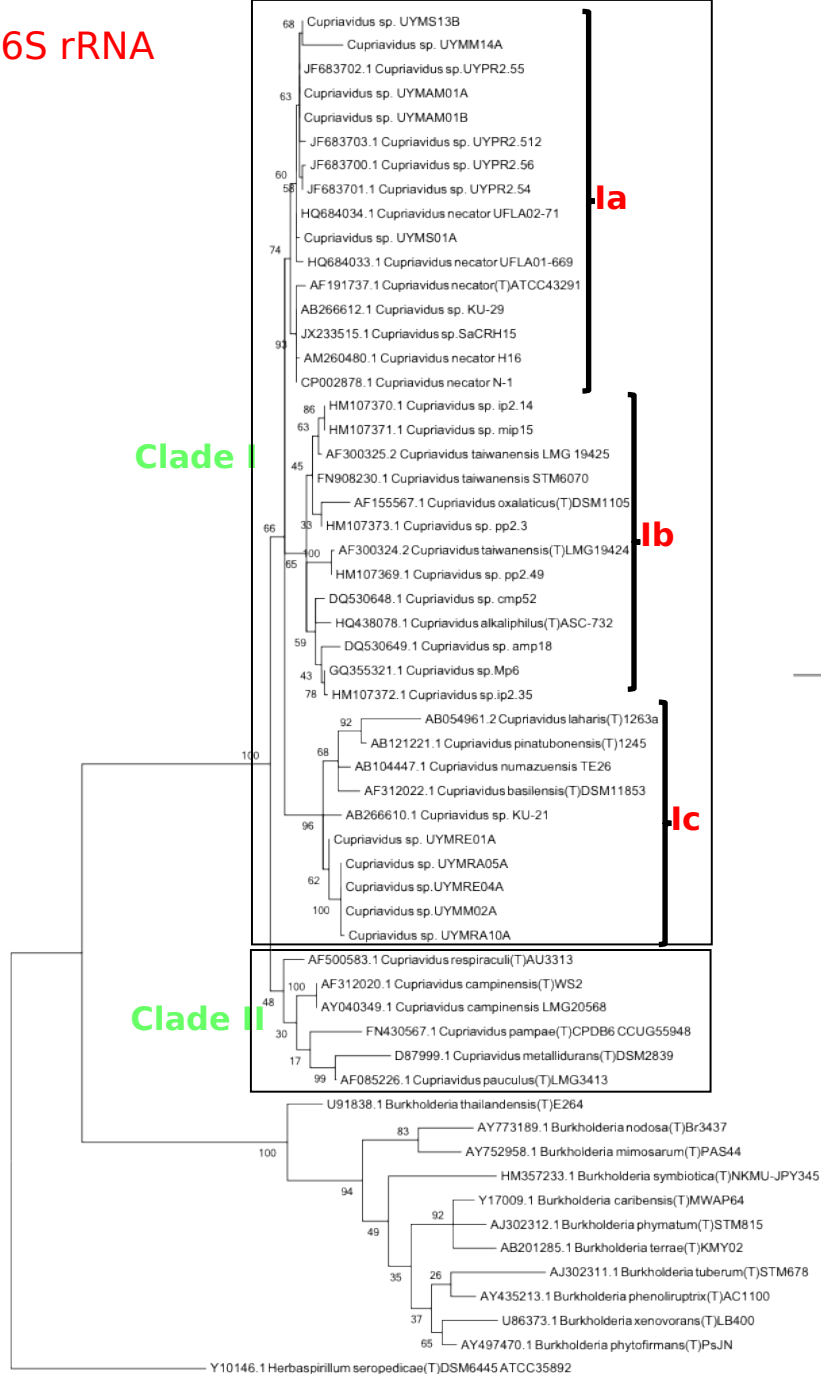


Burkholderia

Cupriavidus

But NOT *C. taiwanensis*

ML 16S rRNA



0.002

Zoom of clade I for more detail

Rufo Cold et al. in prep.

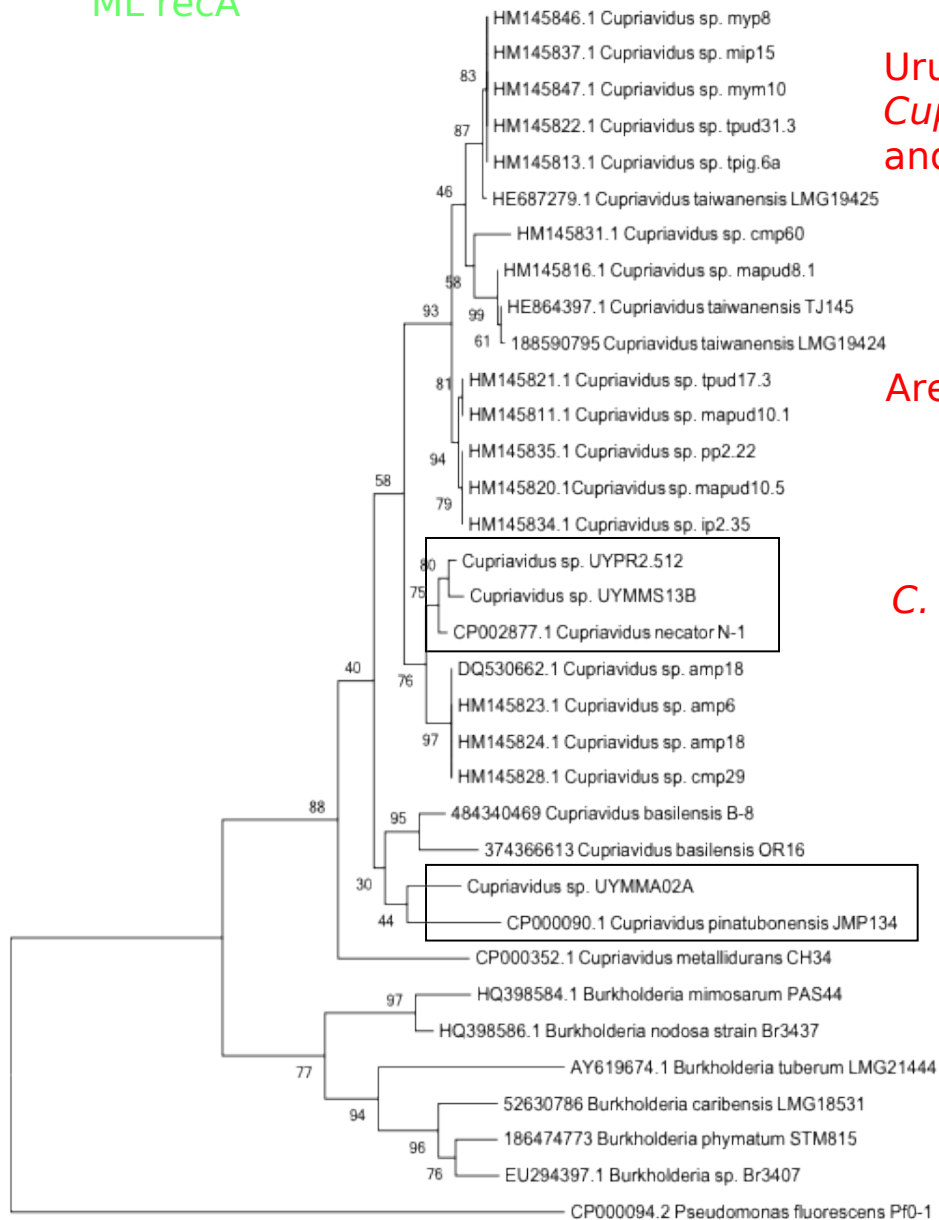
ML recA

Uruguayan *Mimosa* spp. are nodulated by novel *Cupriavidus* symbionts close to *C. necator* and *C. pinatubonensis*

Are the heavy metals in the soils (mining areas) a factor?

C. necator

C. pinatubonensis

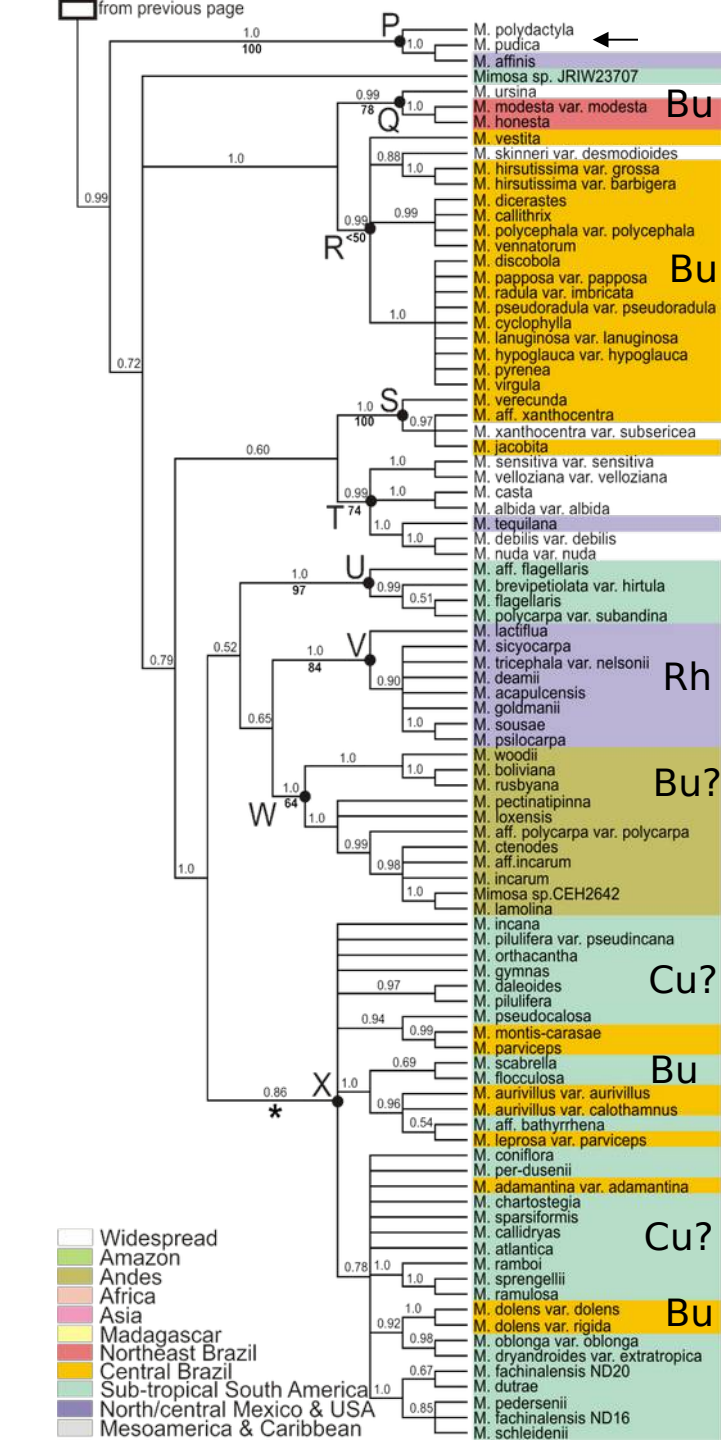
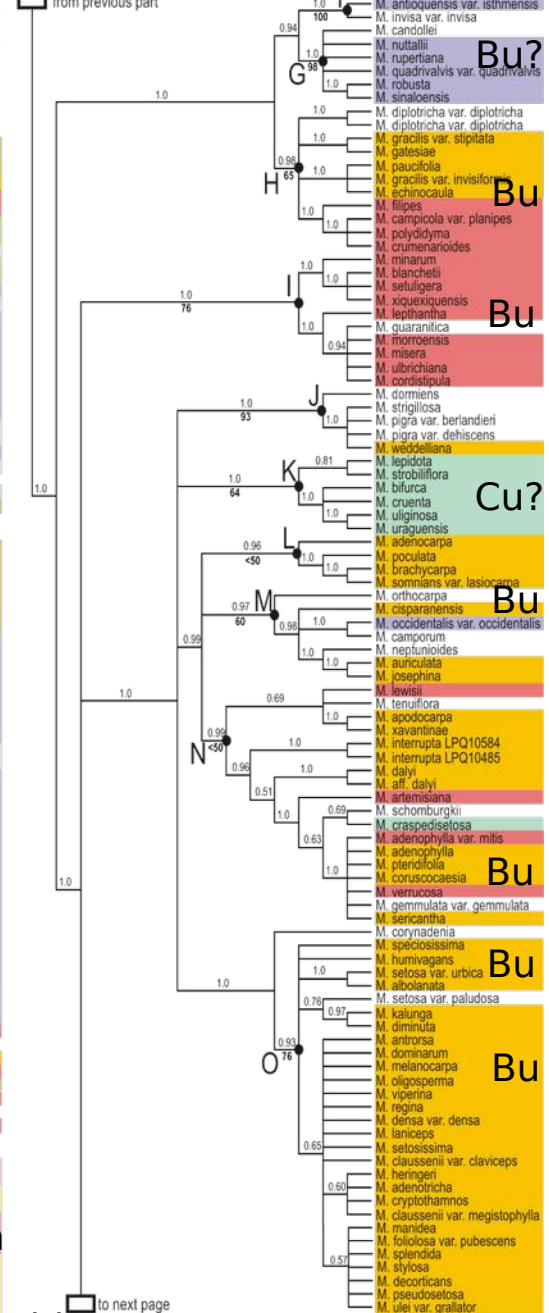
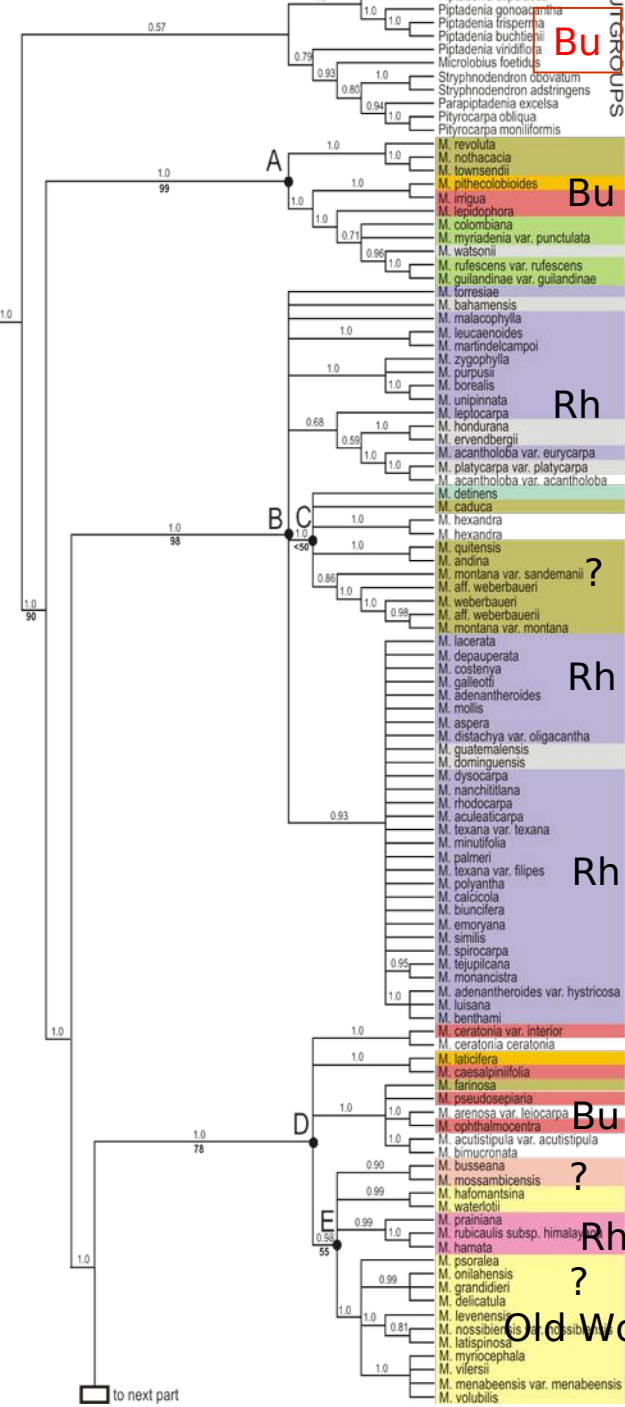


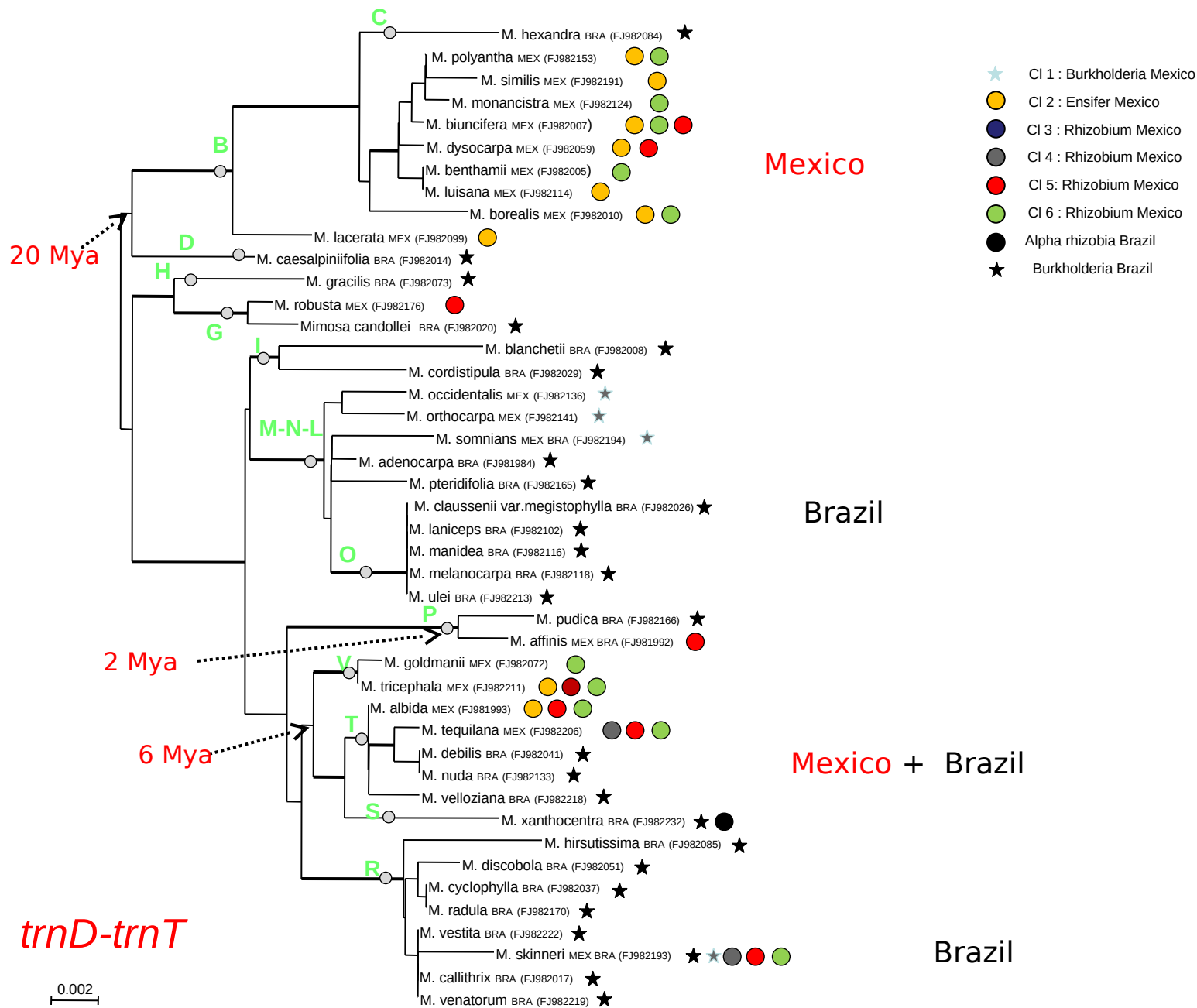
0.05

Rufo Cold et al. in prep.

Summary of endemic *Mimosa* symbionts by biome

- Cerrado (Brazil) = *Burkholderia*
- Caatinga (Brazil) = *Burkholderia*
- Mata Atlantica (Brazil) = *Burkholderia*
- Central highlands (Mexico) = *Rhizobium/Sinorhizobium*
- Uruguay (heavy metal soils) = *Cupriavidus*
- Madagascar/E. Africa = ?
- India = *Sinorhizobium*
- Does *Mimosa* phylogeny/taxonomy have a role in this distribution or is it soil/environment that drives it?





Nodulation of *Mimosa*: conclusions

- *Mimosa* has a propensity for endemism (particularly at high altitudes), and this has resulted in large clades that have strict preferences for particular symbiont genotypes: *Burkholderia* in the case of Brazil and *Rhizobium/Ensifer* in the case of Mexico.
- **Soil characteristics, particularly pH, appear to be the main determining factors in the selection of symbionts.**
- Geographical separation has resulted in these older clades no longer being able to nodulate with other symbiont types.
- BUT, other (much younger) clades are “split” between Brazil and Mexico; these have retained their ancestral ability to nodulate with *Burkholderia*.
- The “switch” from a basal *Burkholderia* preference to the “local” microsymbionts can occur remarkably quickly (<2 my).
- **CONCLUSION 1: Beta-rhizobia are very widespread.**
- **CONCLUSION 2: *Mimosa* is very plastic in terms of adapting to nodulate with a wide range of symbionts, and this has assisted its spread into diverse environments.**

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