

Interplay between lateral gene transfer and rhizosphere functioning

Caroline Choufa, Abdoul-Razak Tidjani, Claire Veneault-Fourrey, Vos Michiel, Cyril Bontemps, Pierre Leblond

▶ To cite this version:

Caroline Choufa, Abdoul-Razak Tidjani, Claire Veneault-Fourrey, Vos Michiel, Cyril Bontemps, et al.. Interplay between lateral gene transfer and rhizosphere functioning. Functional ecology conference 2018, Dec 2018, Nancy, France. hal-03060574

HAL Id: hal-03060574

https://hal.science/hal-03060574

Submitted on 14 Dec 2020

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.









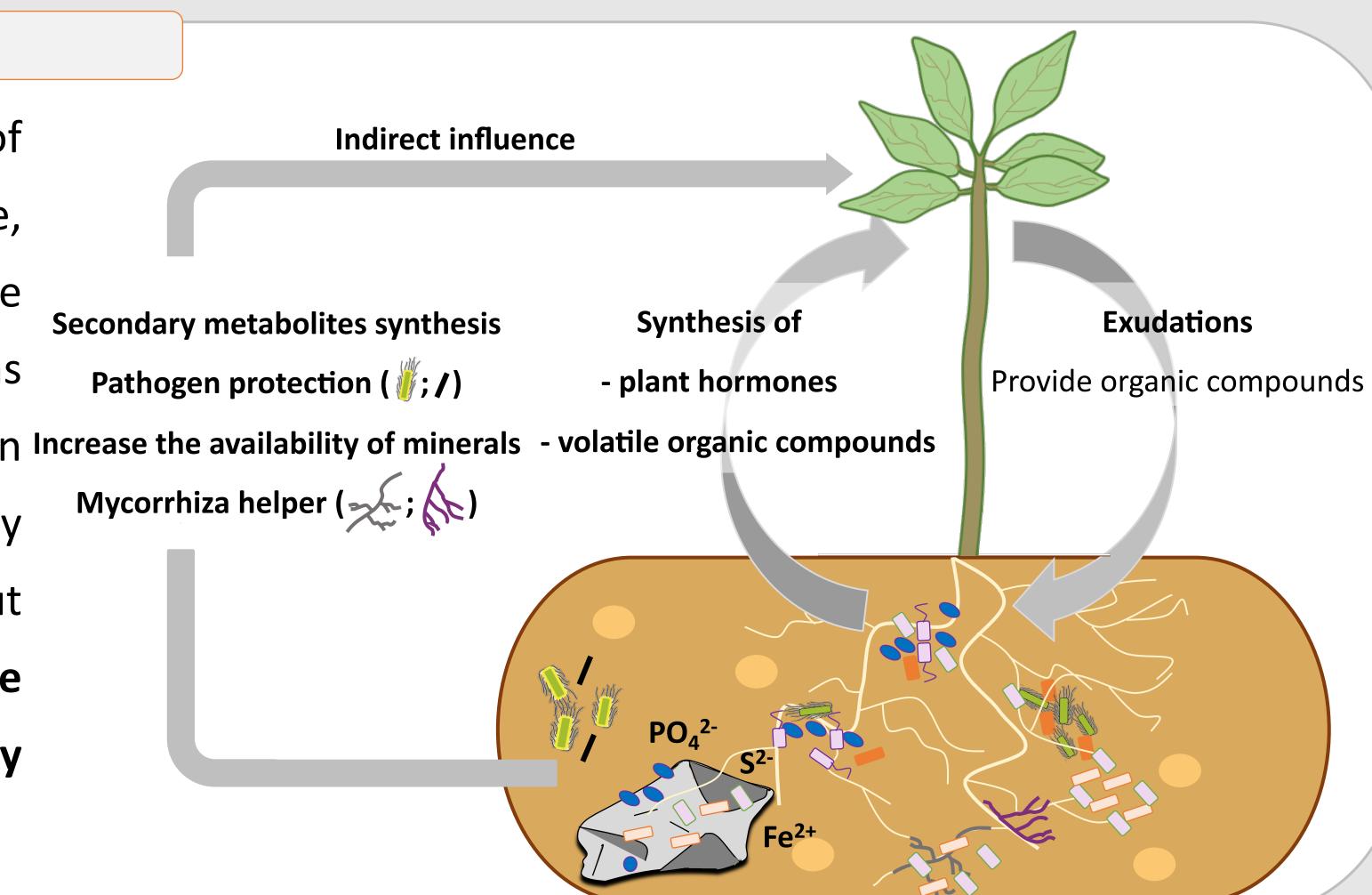
Interplay between lateral gene transfer and rhizosphere functioning

Caroline CHOUFA¹, Abdoul-Razak TIDJANI¹, Claire VENEAULT-FOURREY², Michiel VOS³, Cyril BONTEMPS¹, Pierre LEBLOND¹

¹Université de Lorraine, Inra, DynAMic, F-54000 Nancy, (Fr); ²Université de Lorraine, Inra, IAM, F-54000 Nancy, (Fr); ³European Centre for Environment and Human Health, University of Exeter, (UK)

The rhizosphere and its microbiota

The rhizosphere microbiota contributes to the growth, health of plants and resilience of forest ecosystem [1]. In the rhizosphere, bacteria can influence the plant directly or indirectly and take advantage from **root exudates** in return [2;3]. Until now, it was considered that a bacterial species isolated from a given Increase the availability of minerals environment is a homogenous related cells characterized by identical functions. Yet, bacteria in soil are not single clones, but are structured in populations (sympatric strains of the same species but with different functions) that might act in synergy to achieve a function.



Genus Streptomyces: a model of rhizospheric bacterium

Genome:

Plasticy: chromosomal rearrangements,

linear and circular plasmids [4]

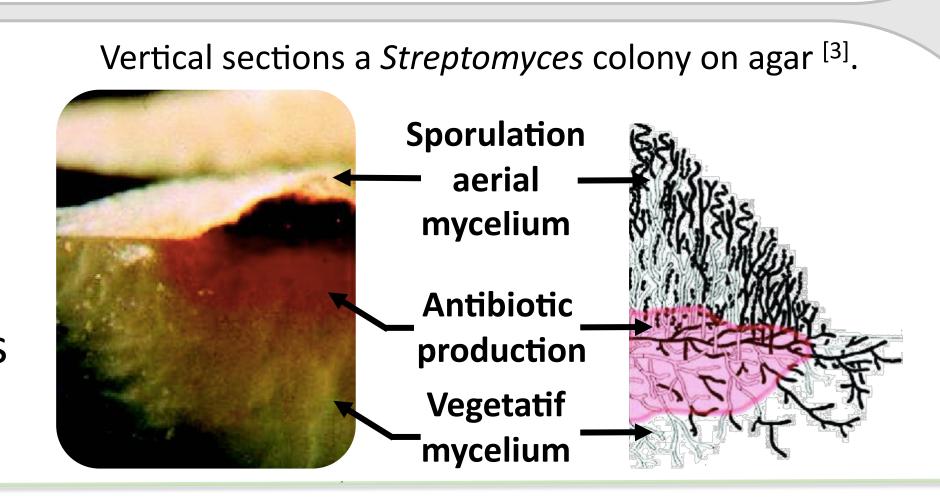
Phylum: *Actinobacteria*

Habitat: soil and rhizosphere

In the lab: collection of natural populations

(rhizospheric and bulk soil)

Early steps of conjugaison



End of conjugaison

Horizontal gene transfer (HGT) promotes functional diversity.

- **AICE** = *Actinobacteria* integrating conjugative element
- Mobilization of chromosomal locus (/)

Conjugative DNA transfer visible to the mere eye (pocks)



† pocks

pocks

Donor strain spore

Recipient strain growth delay

Lawn of recipient cells

The transfer of AICE triggers a growth and sporulation delay (pock) in the recipient cell [4].

What are the impacts of HGT on the sharing of

Donor

Recipient

What are the drivers of HGT in rhizospheric Streptomyces populations?

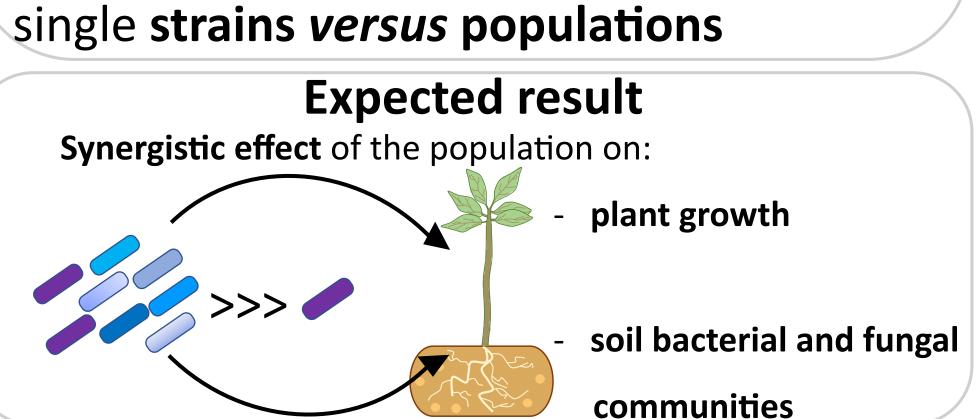
Effect on the HGT in presence of:

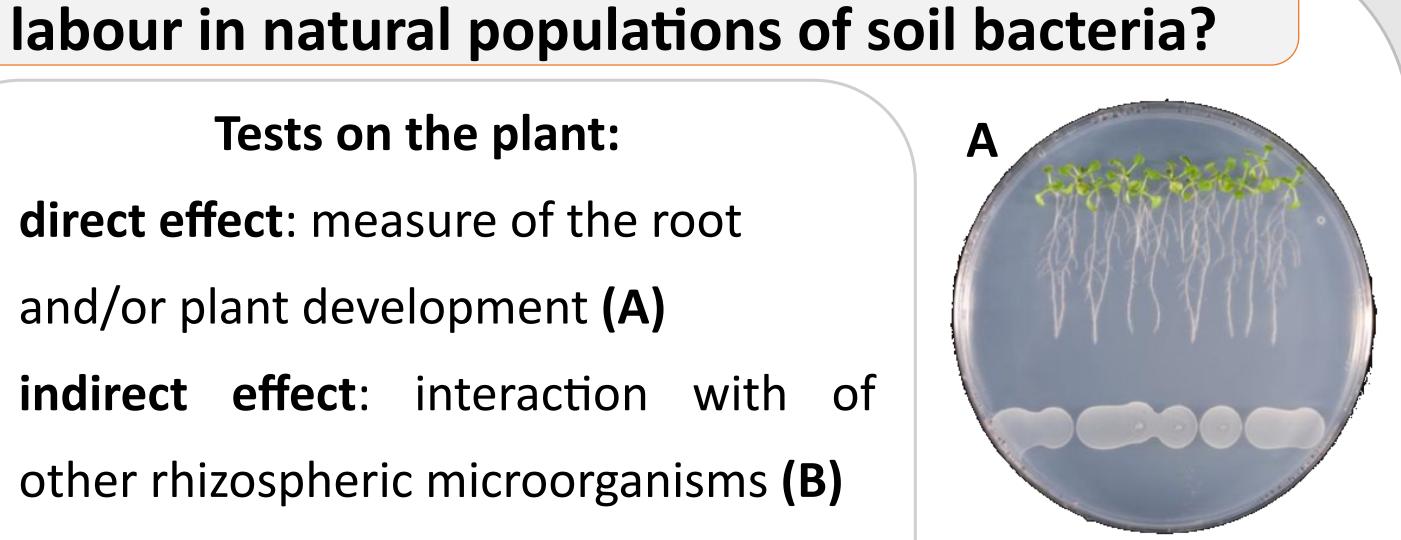
- root system
- exudates
- specific compounds (trehalose, flavonoids, chitin, cellulose ...)

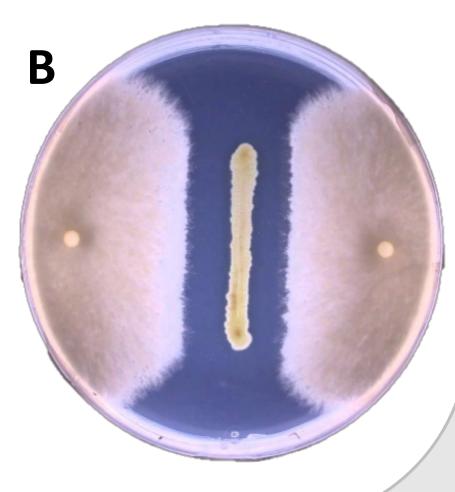
Expected result

direct effect: measure of the root and/or plant development (A) indirect effect: interaction with of other rhizospheric microorganisms (B) Comparison between the effect of natural

Tests on the plant:







Natural strains

Prospects

diversifies

This project will provide an overview of the effects of plant/rhizospheric microbiota interactions with two main objectives:

Natural Population

- understanding if a bacterial population constitutes an efficient ecological unit for plant interaction
- explaining how the plant can affect bacterial genomic diversity.

Furthermore, this project will reveal if gene transfer allows rapid microbiota and plant adaptation to environment changes.



[1] Backer R, Rokem JS, Ilangumaran G, Lamont J, Praslickova D, Ricci E, Subramarian S and Smith DL (2018) Plant growth-promoting rhizobacteria: context, mechanisms of action;

