





Diversity and mobility of conjugative elements in a population of

Streptomyces isolated from a rhizospheric micro-habitat

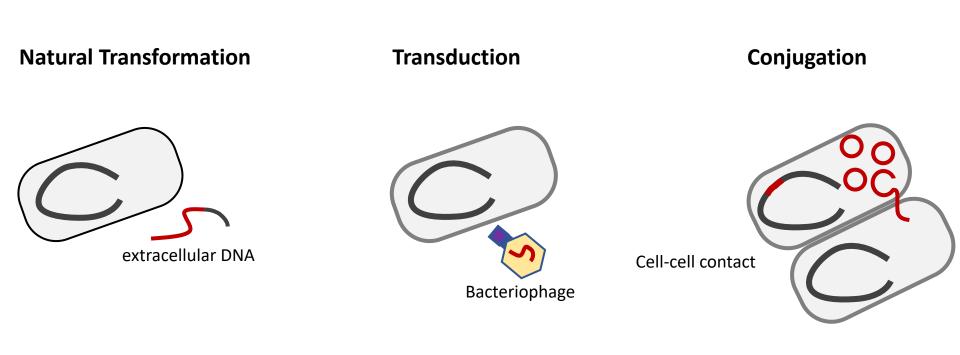
Caroline CHOUFA

UMR UL-INRAE 1128

Dynamique des Génomes et Adaptation Microbienne

Université de Lorraine

Mechanisms of horizontal gene transfer

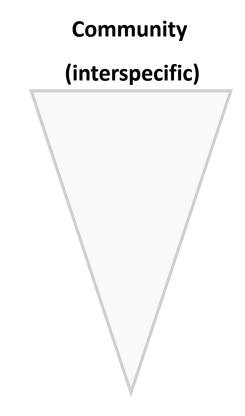


Mechanisms that promote bacterial diversity

Genetic diversity and evolution

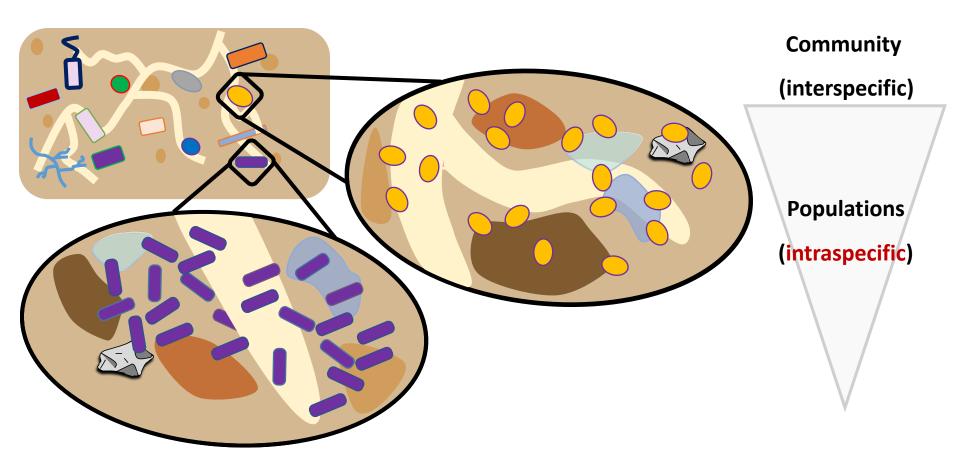
of natural populations of bacterial at micro-scale





Genetic diversity and evolution

of natural populations of bacterial at micro-scale

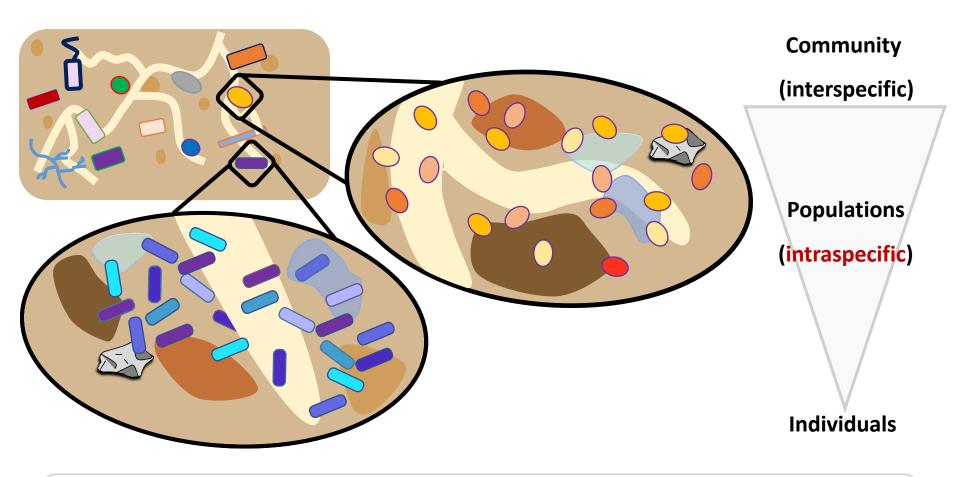


Bacterial population

Sympatric individuals of the same species

Genetic diversity and evolution

of natural populations of bacterial at micro-scale



What is the extent of horizontal transfer within a natural population?

Collection of a natural rhizospheric population of *Streptomyces*

Isolation from a grain of soil

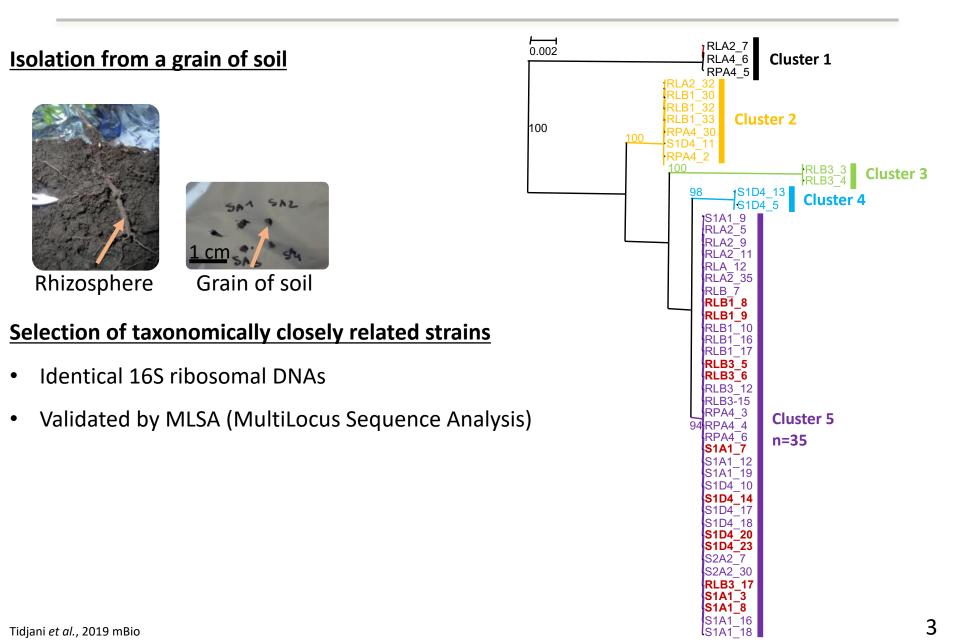




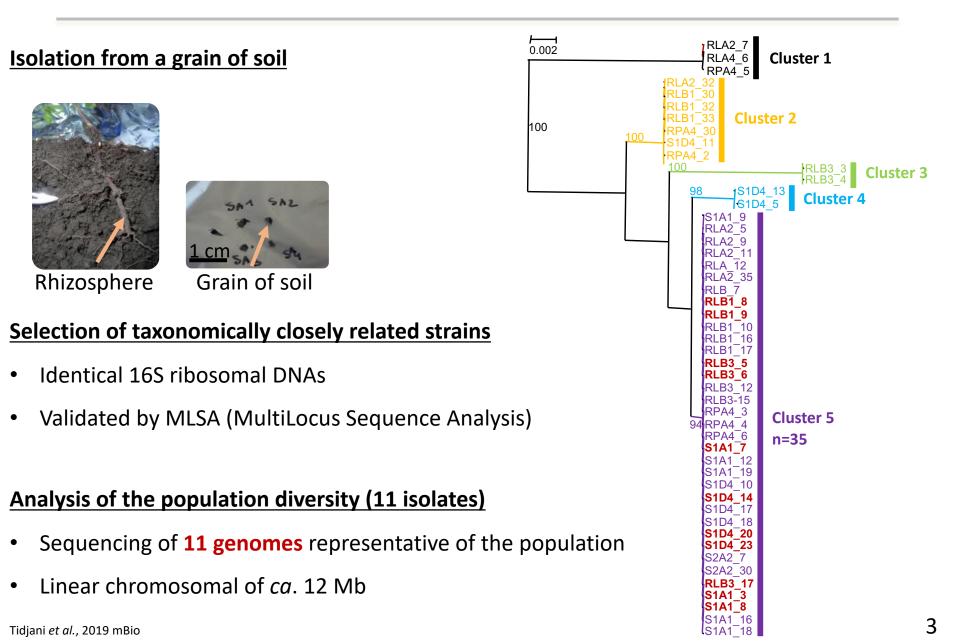
Rhizosphere

Grain of soil

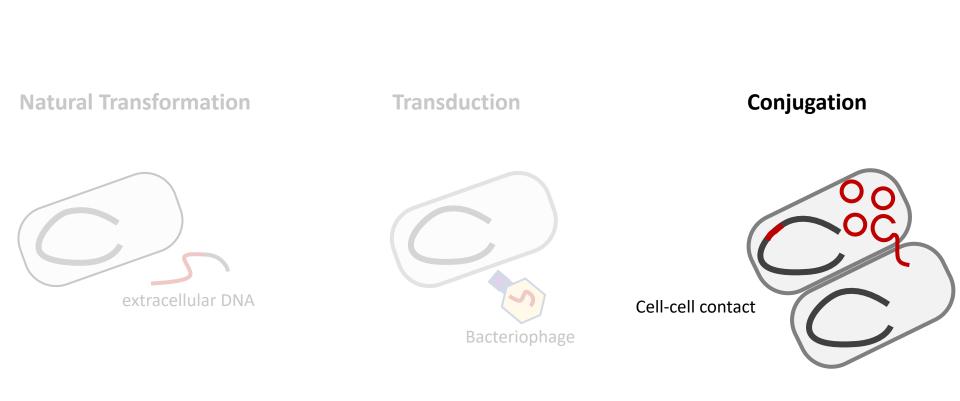
Collection of a natural rhizospheric population of Streptomyces

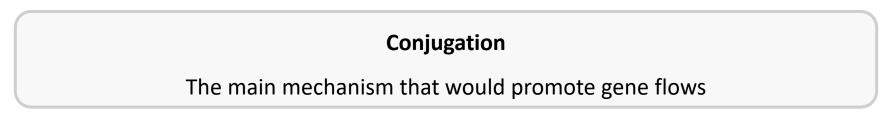


Collection of a natural rhizospheric population of Streptomyces



Mechanisms of horizontal gene transfer in *Streptomyces*

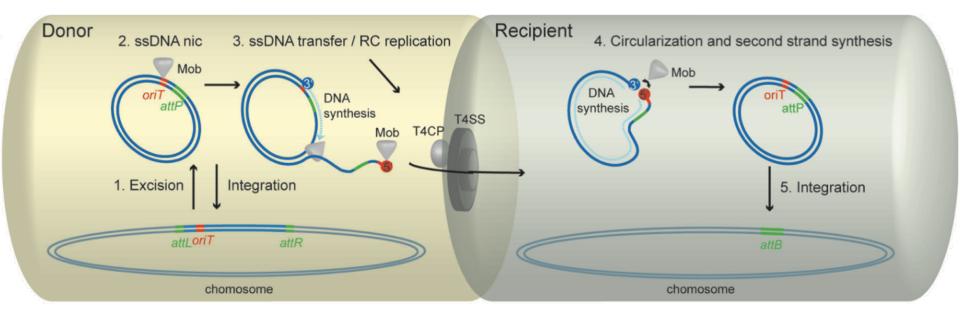




Two mechanisms of conjugation in *Streptomyces*

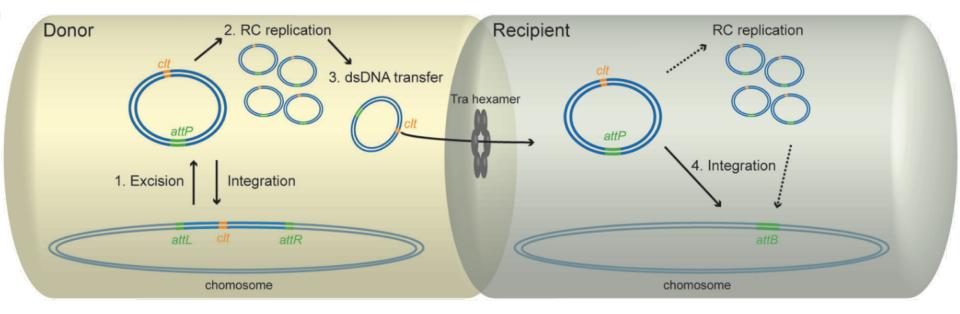
Type IV secretion system dependent (T4SS)

=> ICE (Integrative and Conjugative Elements)

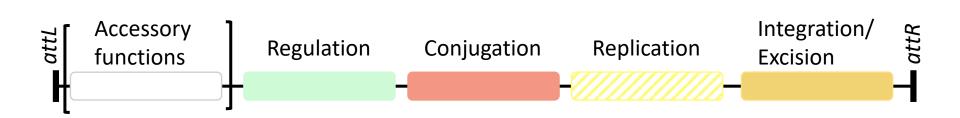


Two mechanisms of conjugation in *Streptomyces*

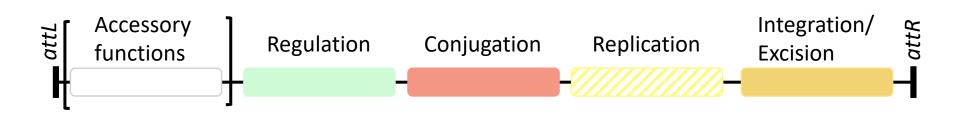
TraB protein dependent (cell division and sporulation FtsK/SpoIIIE-homolog protein)
=> AICE (Actinomycete Integrative and Conjugative Elements)



Search for conjugative elements in *Streptomyces* genome

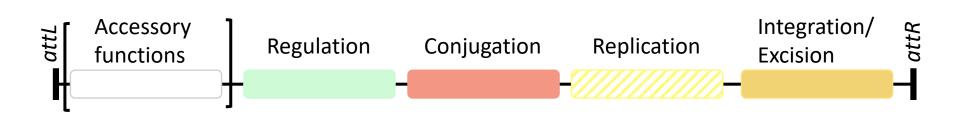


Search for conjugative elements in *Streptomyces* genome



AICE (e.g. pSAM2 S. ambofaciens)

Search for conjugative elements in Streptomyces genome



Ftsk-Spolle (TraB) AICE (e.g. pSAM2 S. ambofaciens) l ^Vor_{S integrase} Replicase Ю or Thp. DDE transposes ICE (e.g. ICESt3 Streptococcus thermophilus) Co_{upling} protein Relaxase (Mob) ^ror_{S inte}grase (Tcoaninoa) Vir_{B4}

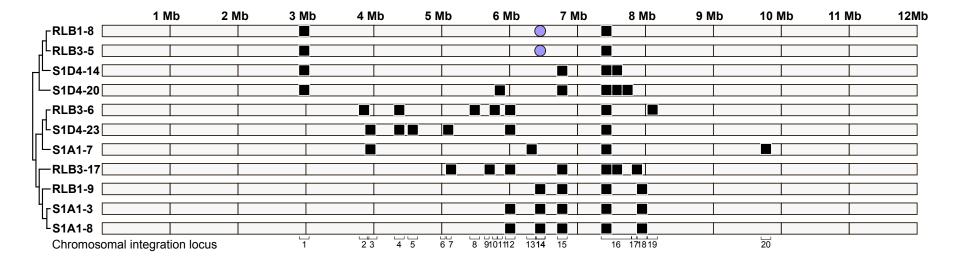
Bordeleau et al., 2012; Hagège et al., 1994; Pavlovic et al.; 2004, Guglielmini et al., 2014; Abby et al, 2016

	1 Mb	2 Mb	3 Mb	4 Mb	5 Mb	6 Mb	7 Mb	8 Mb	9 Mb	10 Mb	11 Mb	12Mb
_┌ RLB1-8												
_LRLB3-5												
S1D4-14												
S1D4-20												
_ RLB3-6												
S1D4-23												
S1A1-7												
RLB3-17												
 S1A1-3												
LS1A1-8												

• 2 ICEs 100% identical in sequence

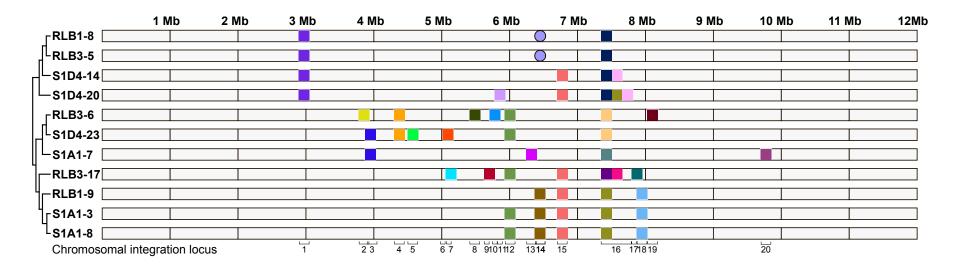
	1 Mb	2 Mb	3 Mb	4 Mb	5 Mb	6 Mb	7 Mb	8 Mb	9 Mb	10 Mb	11 Mb	12Mb
_RLB1-8						(
RLB3-5						(
S1D4-14												
S1D4-20												
_RLB3-6												
S1D4-23												
S1A1-7												
RLB3-17												
_□ S1A1-3												
¹ S1A1-8												
						L	14					

- 2 ICEs 100% identical in sequence
- 52 AICEs



- 2 ICEs 100% identical in sequence
- 52 AICEs divided into 26 families

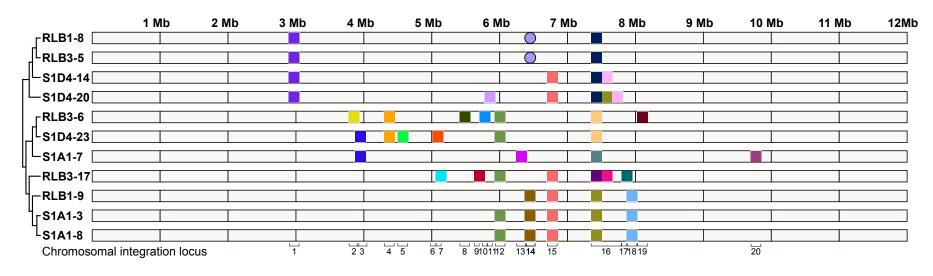
AICEs diverge in size (11.9 to 44.1 kb) and in gene content (15 to 41 CDSs)



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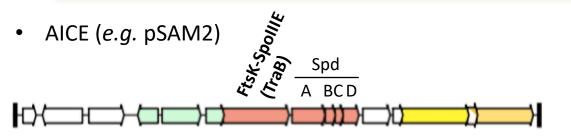
• 54 conjugative elements contain the signature genes



There is considerable diversity in AICEs.

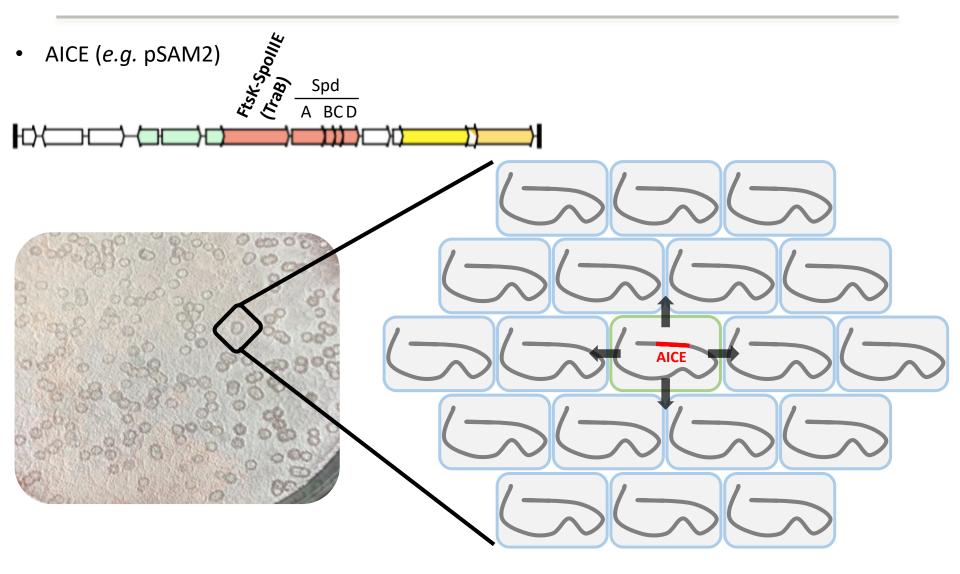
Their distribution suggests an effective dynamic within the population.

AICE transfers are visible to the naked eye: pock phenomenon

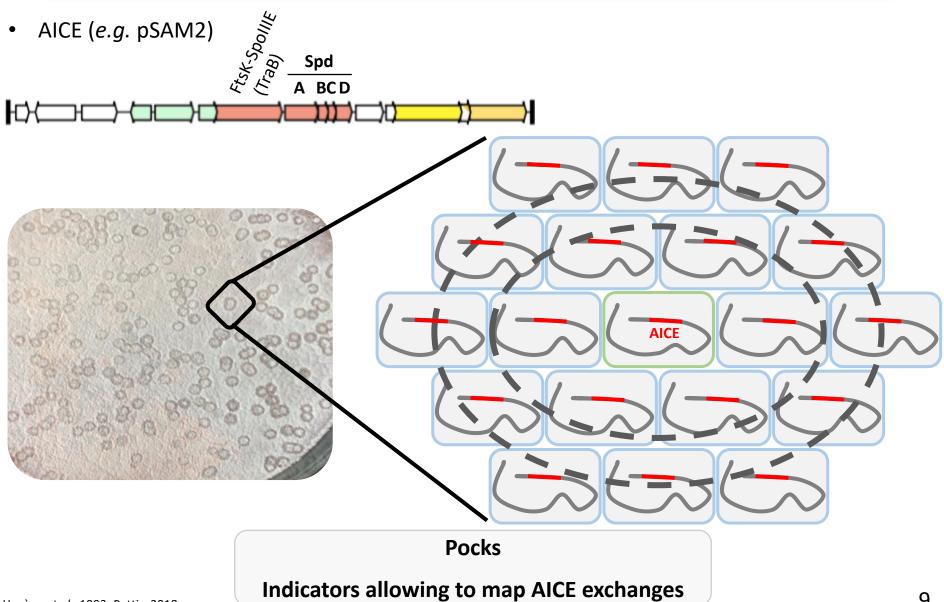




AICE transfers are visible to the naked eye: pock phenomenon



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• 13/110 mating pairs form pocks

AICEs can transfer between indivuals of the population S1D4-20 RLB3-17 A181.9 RLB3:5 RLB1-8 S1A1-3 SID4-14 SIA1-8 RLB3-6 SIAI-> S1D4-23

• 13/110 mating pairs form pocks

AICEs can transfer between indivuals of the population S1D4-20 RLB3-17 A1491.9 PLB3:5 RLB1-8 S1A1-3 D4-14 STAT-8 RLB3-6 SIAI-> S1D4-23

Transfers are not panmictic

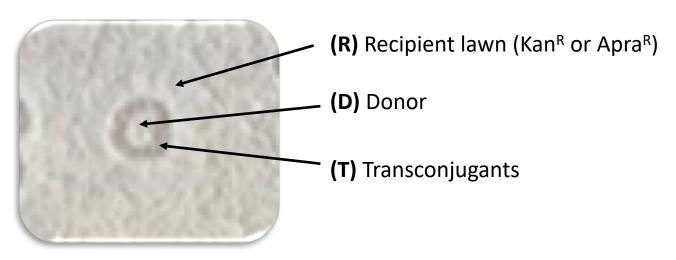
Spd proteins can be absent

Experimental conditions are not optimal

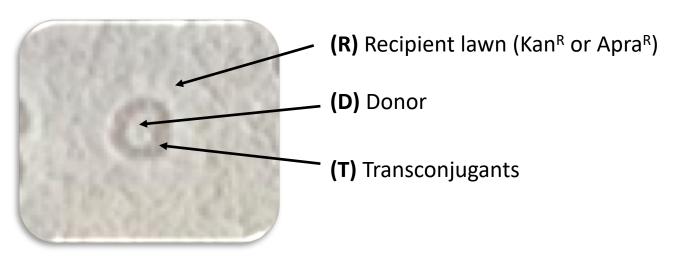
• The absence of pocks does not exclude the conjugation

Identify AICEs responsible for the formation of pocks

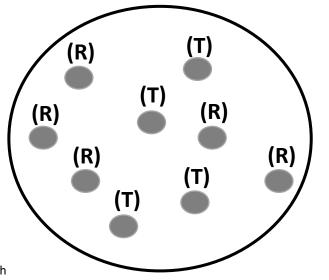
Validation of AICE transfers

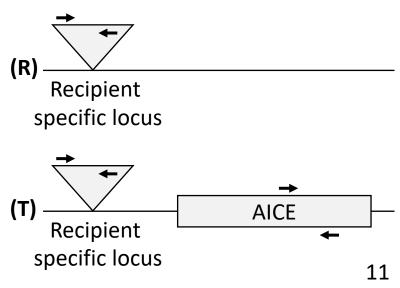


Validation of AICE transfers



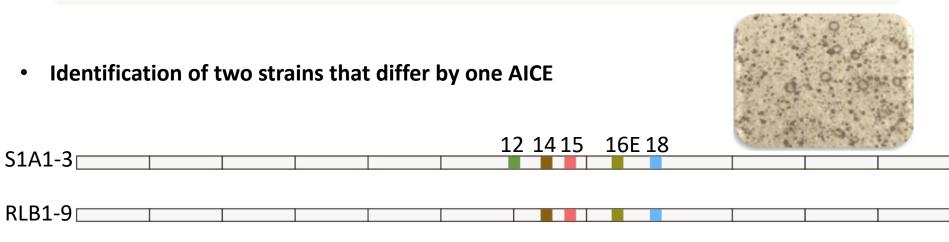
• Isolation and selection of transconjugants from pocks • PCR screening of transconjugants





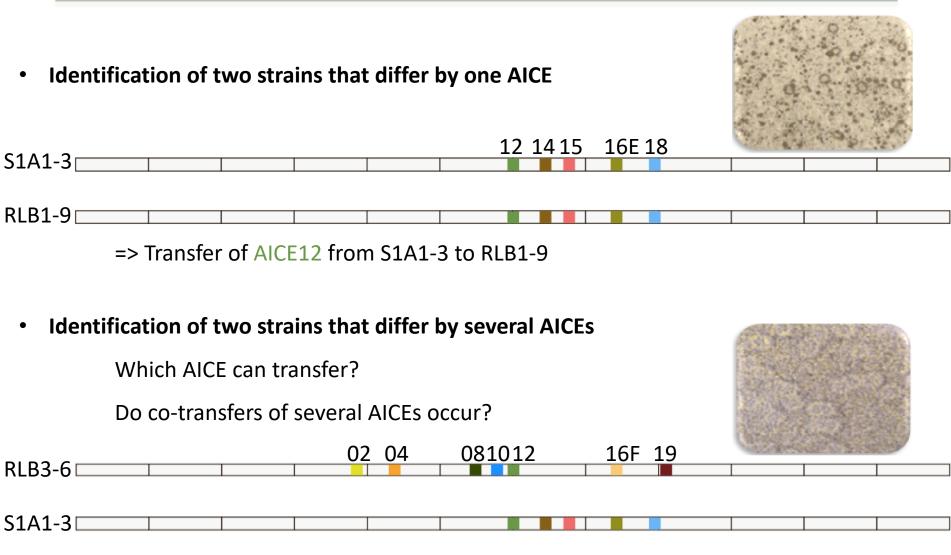
Choufa et al., unpublish

Confirmation of AICE transfer



=> Transfer of AICE12 from S1A1-3 to RLB1-9

Confirmation of AICE transfer

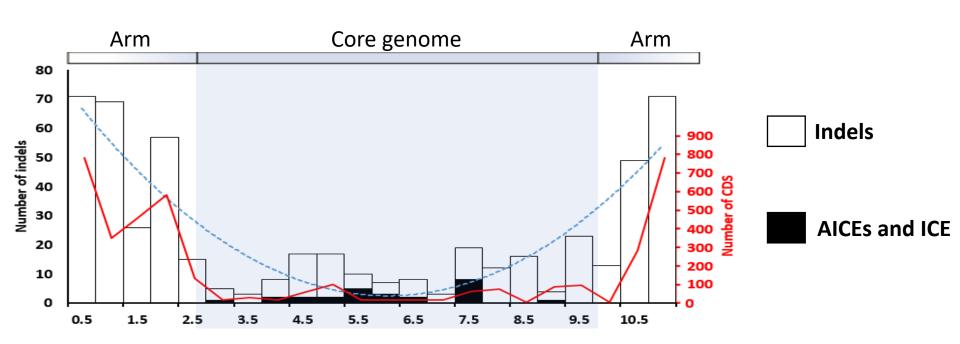


=> Transfer of AICE10 from RLB3-6 to S1A1-3

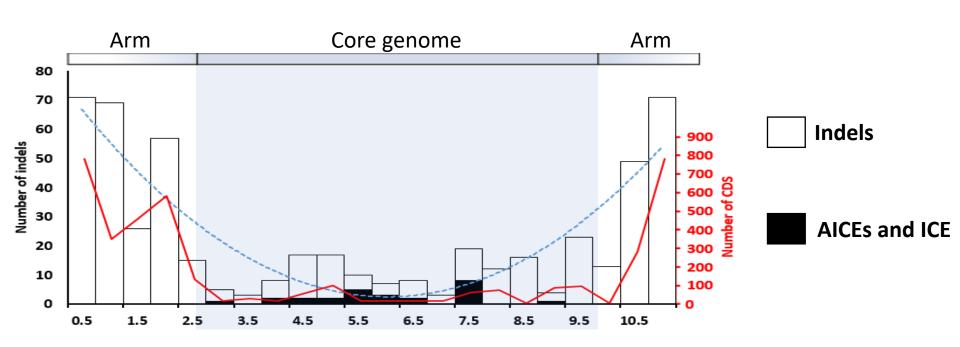
What is the impact of AICE transfers

in the population diversification ?

Distribution of Indels along a consensus genome



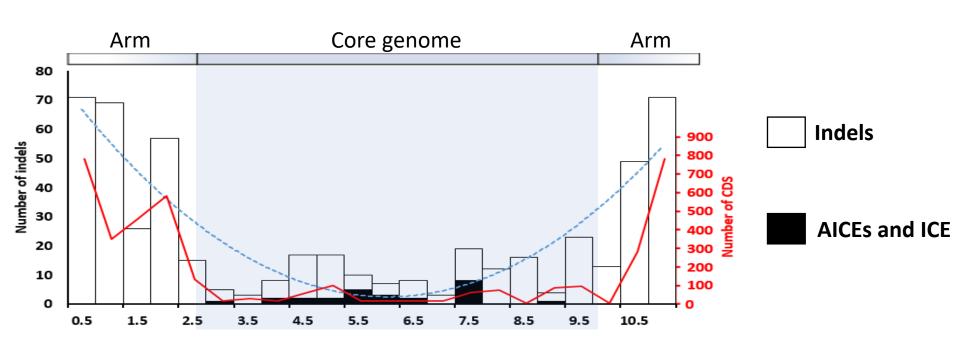
Distribution of Indels along a consensus genome



Indels are more located in chromosomal extremities

AICEs are more located in the chromosomal middle

Distribution of Indels along a consensus genome



Indels are more located in chromosomal extremities

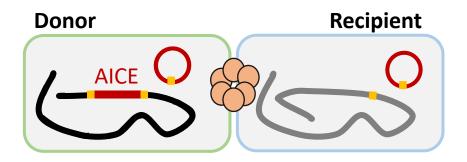
AICEs are more located in the chromosomal middle

Do AICEs promote genetic diversification of the population? Are they responsible for the gradient of gene flux towards the terminal ends ?

Mobilization of chromosomal DNA in *trans* by AICEs

The DNA translocase TraB is capable of mobilizing chromosomal DNA in trans,

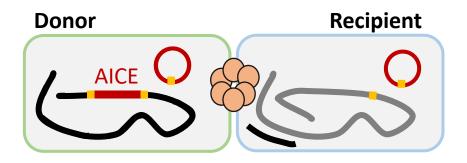
i.e., **not physically linked** to the element.



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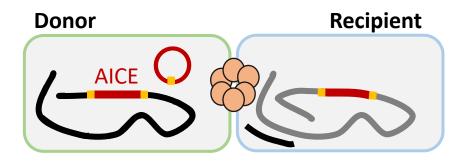
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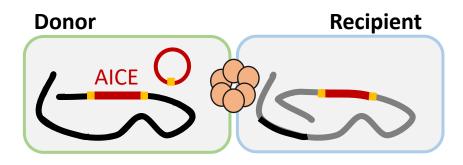
i.e., **not physically linked** to the element.



Extent of the mobilization of chromosomal DNA in *trans* by AICEs

The trans mobilization of DNA fragments during mating

could contribute to the genetic diversification of a population

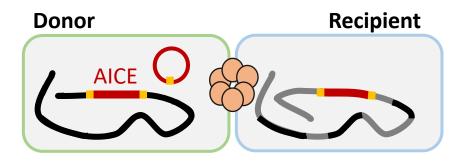




Extent of the mobilization of chromosomal DNA in trans by AICEs

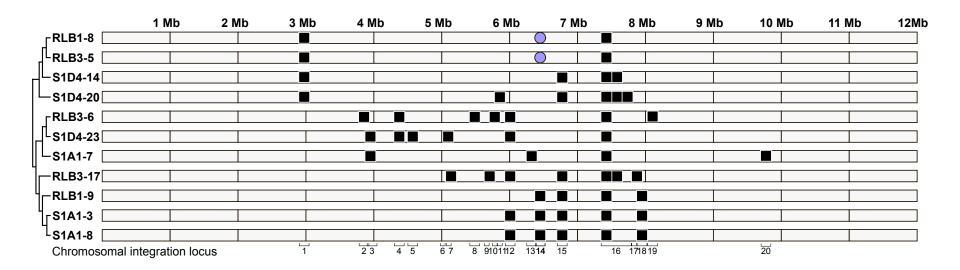
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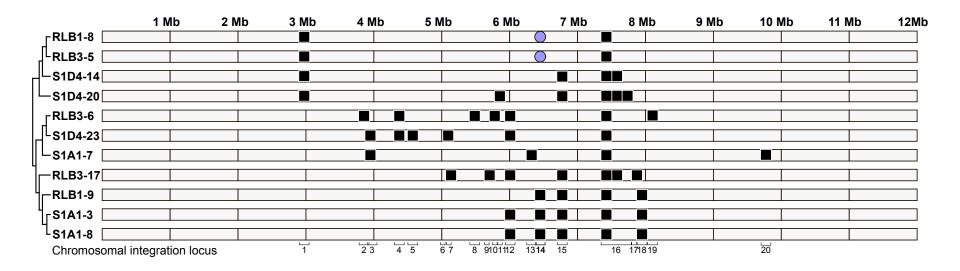


Is the T4SS-conjugation dependent effective in *Streptomyces*



Is the T4SS-conjugation dependent effective in *Streptomyces*

- Some scarce T4SS-ICEs have been predicted in silico in actinobacteria.
- The excision of two ICEs from *Frankia sp.* Cci3 was confirmed by PCR
- The T4SS tranfer has never been experimentally demonstrated



Is the transfer of the ICE14 of *Streptomyces sp.* RLB1-8

or Streptomyces sp. RLB3-5 effective?

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Dynamique des Génomes et Adaptation Microbienne

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Frédéric Bourgaud, Vice-Président de Recherche

Merci pour votre attention

Dynamique des Génomes et Adaptation Microbienne

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