

A logic programming exploration of metabolic networks to explain microbial community models

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Ínría-

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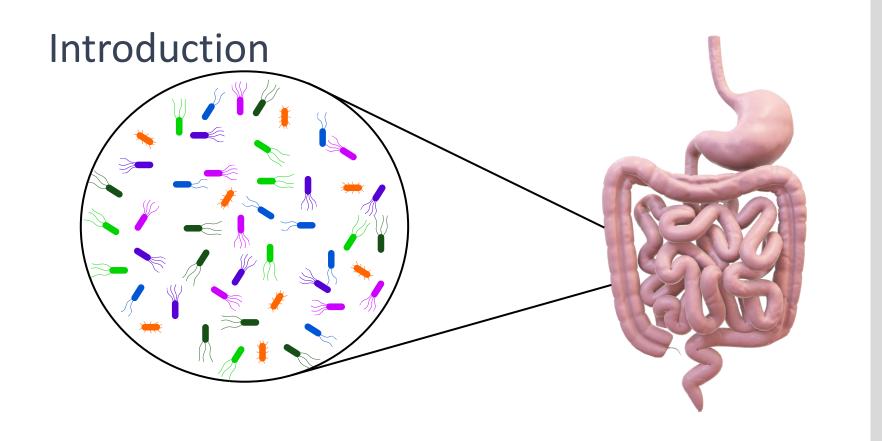
PhD student: Chabname GHASSEMI (LaBRI / Inria)

Co-directors : Loïc PAULEVE (LaBRI)
Clémence FRIOUX (Inria)

A logic programming exploration of metabolic networks to explain microbial community models

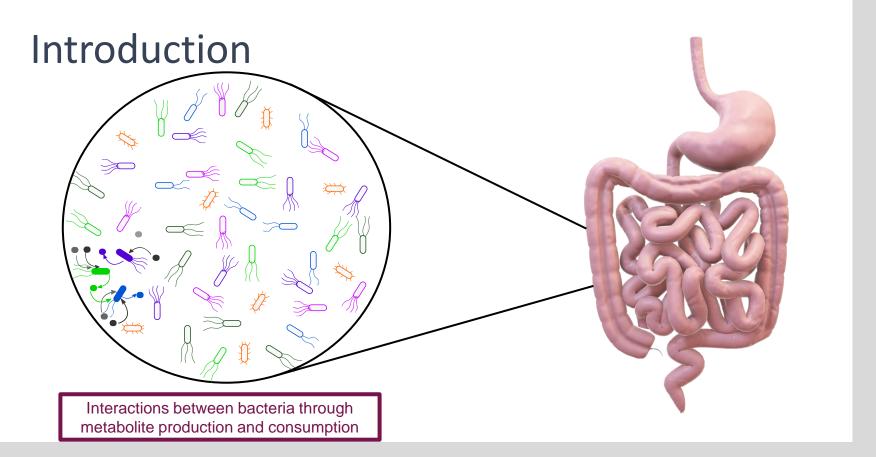
LaBRI





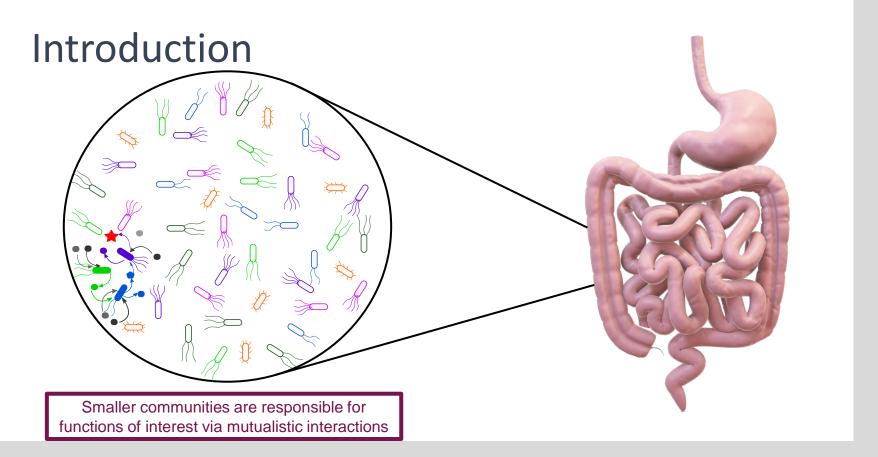










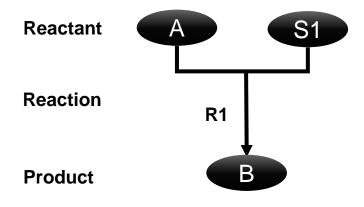




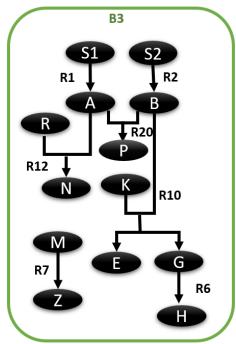


Genome Scale Metabolic Networks (GSMNs)

How to represent metabolism?



GSMNs: set of hundreds of reactions associated to a genome





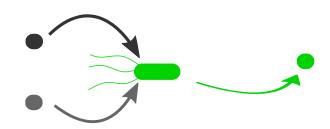
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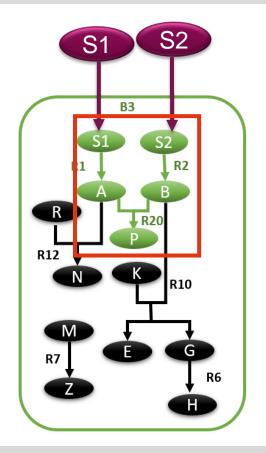
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Individual Scope

Boolean abstraction of metabolic producibility in an environment



Metabolites produced by individual species



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Logic programming to model metabolism

Use of Answer Set Programming (ASP)

Reasoning on knowledge and solving combinatorial problems

```
bacteria("B3").

species("M_A_c","A","c","B3").

reaction("R12"," B3").

reactant ("M_A_c","R12","B3").

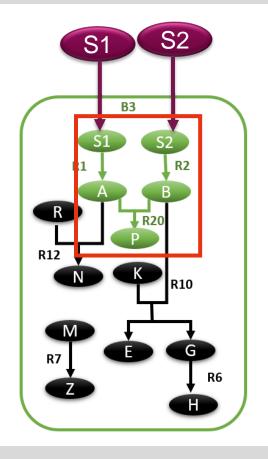
reactant ("M_R_c","R12","B3").

product("M_N_c","R12","B3").

seed("S1").

seed("S2").
```

RULES



Explainable models



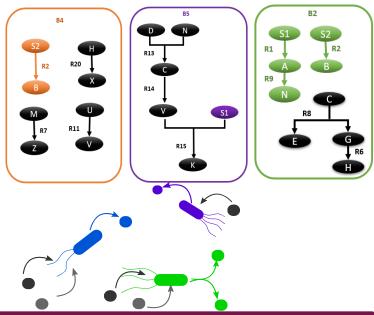
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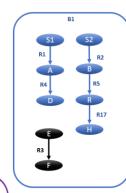
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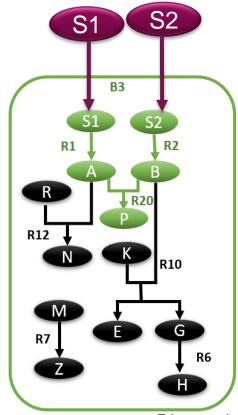
Community of bacteria

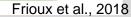
Hundreds of bacteria live in an environment











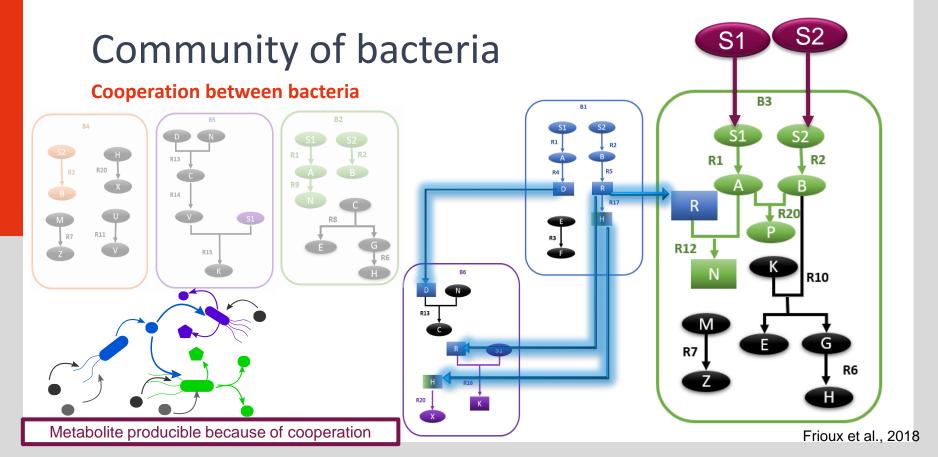


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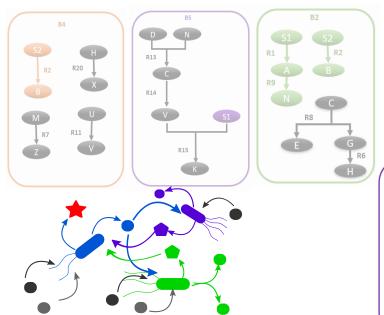
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S1

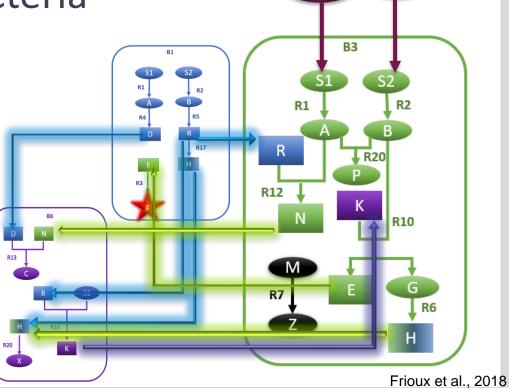
S2

Community of bacteria

Metabolic potential



Target metabolites become producible through metabolites exchanges





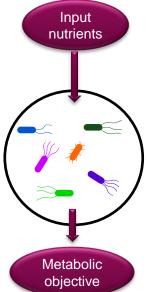
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Effect of Metabolic redundancy on community selection



Community identified with metabolic potential to produce a target Redundancy of metabolism in different species

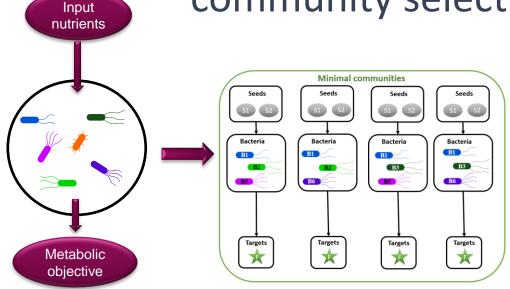
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Frioux et al., 2018 Belcour, Frioux et al. 2020





Effect of Metabolic redundancy on community selection



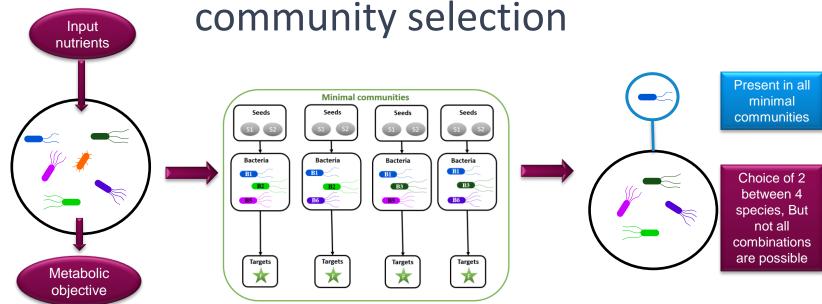
Community identified with metabolic potential to produce a target Redundancy of metabolism in different species

Frioux et al., 2018
Belcour, Frioux et al. 2020





Effect of Metabolic redundancy on community selection



Community identified with metabolic potential to produce a target Redundancy of metabolism in different species

Frioux et al., 2018 Belcour, Frioux et al. 2020

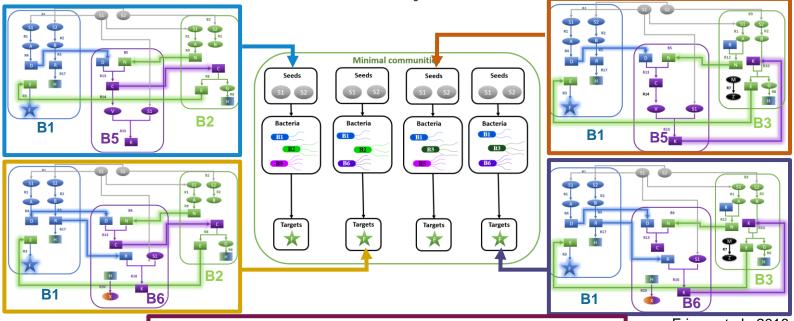
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17/11/2022 Chabname GHASSEMI Effect of Metabolic redundancy on community selection



Mechanistic model to predict smaller communities

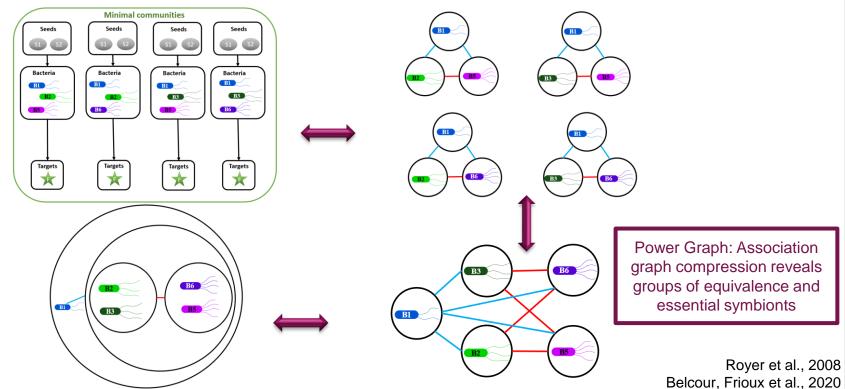
Frioux et al., 2018
Belcour, Frioux et al. 2020





Explainable models

How visualize all communities

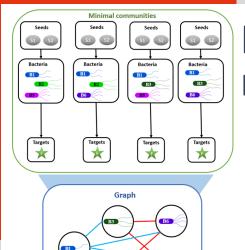




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Power graph

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Metage2Metabo: Application on real data

Lipids Lipids Actinobacteria Bacteroidetes Firmicutes Fusobacteria Proteobacteria

How visualize all communities?

Scaling up to 1,500 bacteria, with a set of 27 lipids as targets

- Community structure
 - > 185 bacteria involved
 - > 7 bacteria per community
 - > 58,520 communities

Belcour, Frioux et al., 2020

Functions not identified on equivalence group.

Why the species are equivalent?





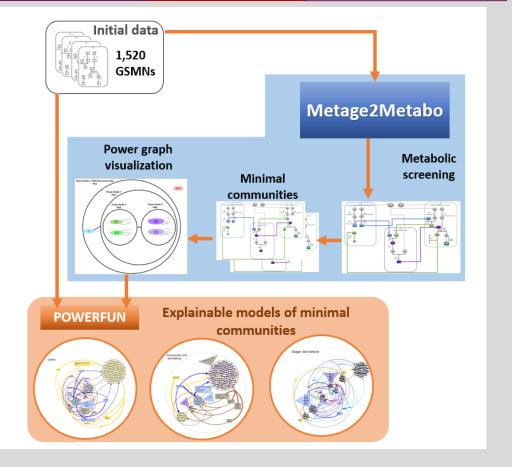
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Explainable models of minimal community assembly

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What functions are carried by a group of equivalence?

Which metabolites are redundant between species?



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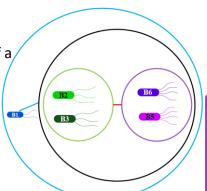
Equivalence group: metabolite produced by a reaction both existing only on a node

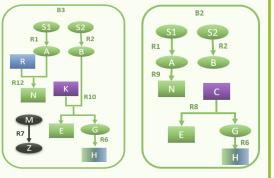
Which functions are carried by equivalence group?

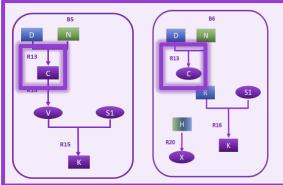
Rules to reveal shared functions of a group of bacteria

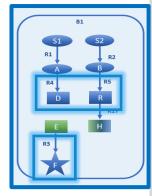
 Metareact: unique reaction producing unique metabolite

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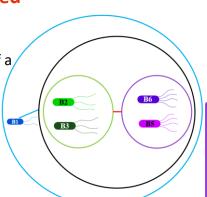
Equivalence group: node specific producible

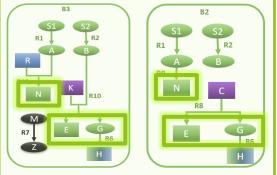
metabolites

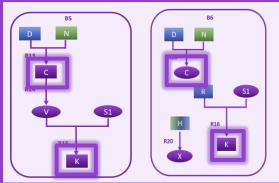
Which functions are carried by equivalence group?

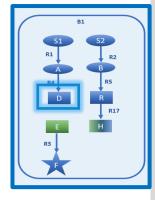
Rules to reveal shared functions of a group of bacteria

- Metareact: unique reaction producing unique metabolite
- Node specific metabolite













Equivalence group: node specific activated

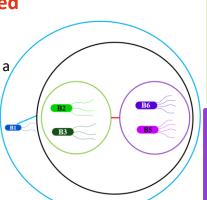
reactions

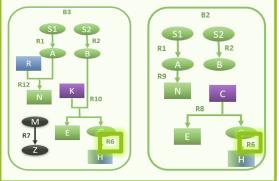
Which functions are carried by equivalence group?

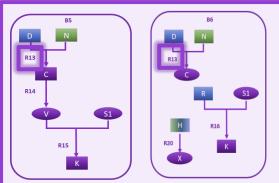
Rules to reveal shared functions of a group of bacteria

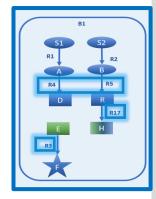
- Metareact: unique reaction producing unique metabolite
- Node specific metabolite
- Node specific reaction

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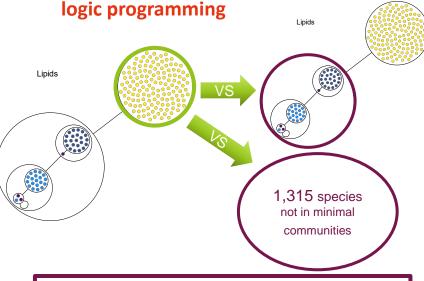


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PowerFun: Powernodes Functions

ASP based tool to explain assembly and redundancy of microbial communities with logic programming



From genome to shared functions of bacteria on minimal communities

ATOMS

```
powernode(P,T).
included(B,P).
```

RULES

```
1{selected pwrn(P) : powernode(P), included(B,P), bacteria(B)}1.
selected_bact(B,P) :- bacteria(B), included(B,P),
                      selected pwrn(P).
producible_by_select(M,B) :- select_activ_reaction(R,B),
                             product(M,R,B).
producible_by_other(M,B) :- other_activ_reaction(R,B),
                            product(M,R,B).
producible_only_select(M,B) :- producible_by_select(M,B),
                                not producible by other(M, ).
prod only all select(M) :-
      producible only select(M,B1) : selected bact(B1, );
      producible only select(M,B2) : selected bact(B2, );
      producible only select(M,B1), producible only select(M,B2),
      B1 != B2.
prod_only_all_select(M) :- nb_selected_bact(N), N=1;
                 producible only select(M,B).
```

https://gitlab.inria.fr/pleiade/powerfun



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Application on real data

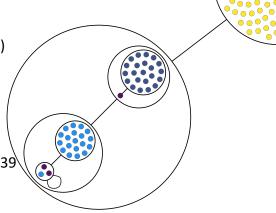
PowerFun computed for the lipids powergraph

Data and performance

• 30 min to compute (1CPU, 50GB)

85 node specific reactions, 24 activated

 209 node specific metabolites, 139\ carried by one powernode Lipids



Bacteroidetes
Firmicutes
Fusobacteria
Proteobacteria

Actinobacteria

1,500 bacteria, with a set of 27 lipids as targets

- Community structure
 - > 185 bacteria involved
 - > 7 bacteria per community
 - > 58,520 communities

Scaling up and analyses of node specific metabolite and reactions





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Node specific metabolites



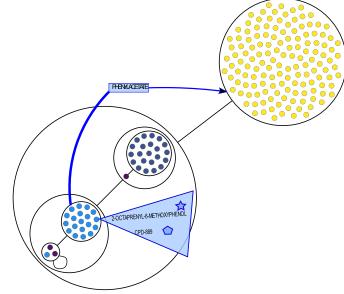
Bacteroidetes

Firmicutes

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Fusobacteria

Proteobacteria



Focus on metabolites produced by all species of a node that are transferred, target and/or from reaction existing only on a node



Metabolite produce only by powernode by a reaction existing only in powernode



Target produced only by the ponernode



Metabolite transferred to other powernode



Metabolite produced only by the ponernode



Metabolite produced only by the ponernode and transferred



Metabolite produced by one of the bacterium in the ponernode



Metabolite produced by one of the bacterium of the ponernode and transferred





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Application on real data: Power nodes

flow

Actinobacteria

Bacteroidetes

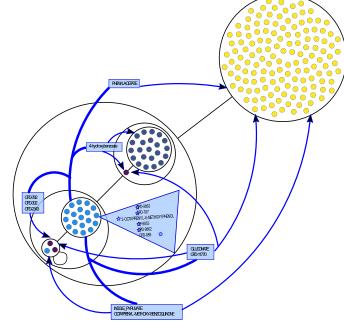
Firmicutes

Fusobacteria

Proteobacteria

Equivalence group produces alone a set of targets or transfers node specific metabolites

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Metabolite produce only by powernode by a reaction existing only in powernode



Target produced only by the ponernode



Metabolite transferred to other powernode



Metabolite produced only by the ponernode



Metabolite produced only by the ponernode and transferred



Metabolite produced by one of the bacterium in the ponernode



Metabolite produced by one of the bacterium of the ponernode and transferred



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Application on real data: Power nodes

flow

Actinobacteria

Bacteroidetes

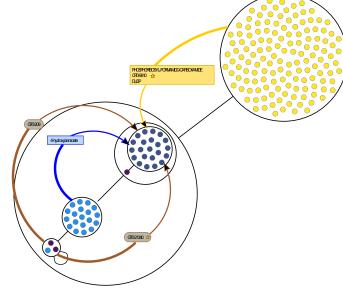
Firmicutes

Fusobacteria

Proteobacteria

Power node not explained by defined rules: no specific metabolites produced by the node

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Metabolite produce only by powernode by a reaction existing only in powernode



Target produced only by the ponernode



Metabolite transferred to other powernode



Metabolite produced only by the ponernode



Metabolite produced only by the ponernode and transferred



Metabolite produced by one of the bacterium in the ponernode



Metabolite produced by one of the bacterium of the ponernode and transferred



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Application on real data: Complete flow



Bacteroidetes

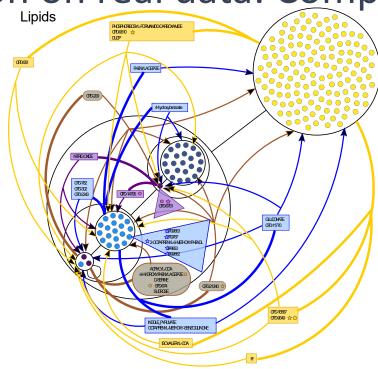
Firmicutes

Fusobacteria

Proteobacteria

Complex model of node specific metabolite production and transfers for a small community

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Metabolite produce only by powernode by a reaction existing only in powernode



Target produced only by the ponernode



Metabolite transferred to other powernode



Metabolite produced only by the ponernode



Metabolite produced only by the ponernode and transferred



Metabolite produced by one of the bacterium in the ponernode



Metabolite produced by one of the bacterium of the ponernode and transferred



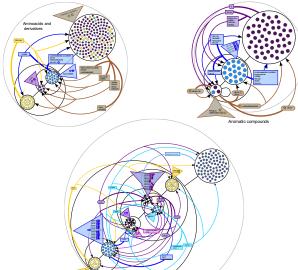


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Discussion

- PowerFun is able to explain equivalence group of species
- Some groups are not explained
- Explainable models remain complex



Sugar derivatives



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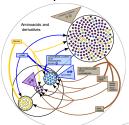


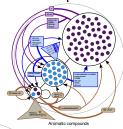
Discussion and perspectives

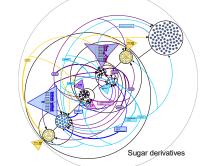
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Discussion

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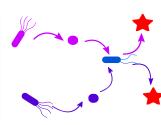


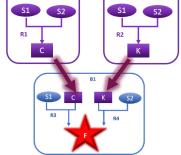
Perspectives

Define new rules and models: A non-explained group can have two different metabolite activating different path

Explainable models







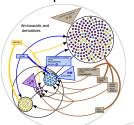




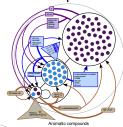
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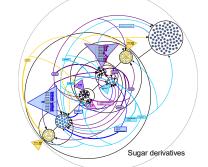
Discussion

- PowerFun is able to explain equivalence group of species
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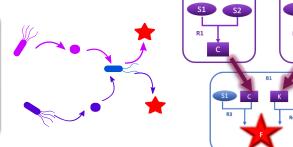




Perspectives

Define new rules and models: A non-explained group can have two different metabolite activating different path

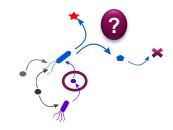
completion



Constrain definition of exchange metabolites



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Thank You!

Team Pleiade



