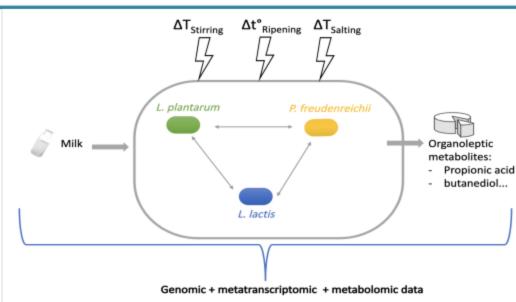


# MODELLING METABOLIC NETWORK DYNAMICS IN A CHEESE BACTERIAL COMMUNITY

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### Context

- How do industrial processes impact cheese's aroma and taste?
- Fermentation process enables bacteria to produce organoleptic compounds
- Bacteria uptake compounds in growth media and release final compounds in the shared environment
- **Variant itineraries** are tested and their effect is measured through multi-omics experiments [1]



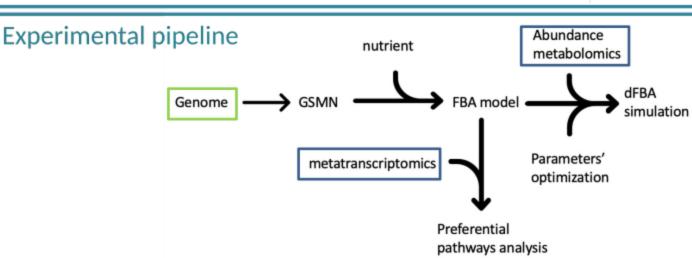


Figure 1: Overall pipeline used to produce results (is being automated).

Green square shows the minimal requirement to achieve FBA model and dFBA simulation

Blue squares show available data which improve the accuracy of the result

## Genome scale metabolic network (GSMN) of the cheese community

Bacteria	Genes	Reactions	Associated reactions	% reactions associated at least one gene
L. Plantarum	1433	1517	1016	67
L. Lactis	1272	1335	940	70
P. Freudenreichii	1473	1789	1177	66

Table 1: Representation of global characteristics of each GSMN [2,4]: number of genes and reactions identified, number of reactions associated to a gene and percentage of reactions linked at least one genes.

### Application and results

Small cheese bacterial community : *L. lactis* (Ll), *L.* plantarum (Lp) and *P.* freudenreichii (Pf)

Cooperation Ll/Pf and Lp/Pf --lactic acid

Cooperation Ll/Pf --> acetaldehyde

Competition Ll/Lp --> carbon source (lactose)

**Preferential** pathway identified in **accordance** with literature [5]

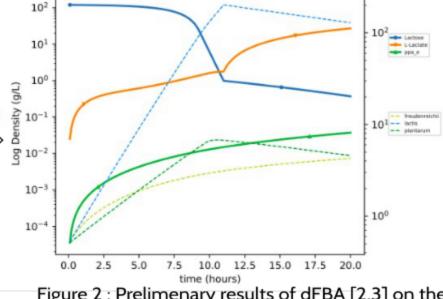


Figure 2 : Prelimenary results of dFBA [2,3] on the whole community:

- Production of organoleptic compound for Pf (propionic acid) from lactate
- Cooperation between lactic species and Pf

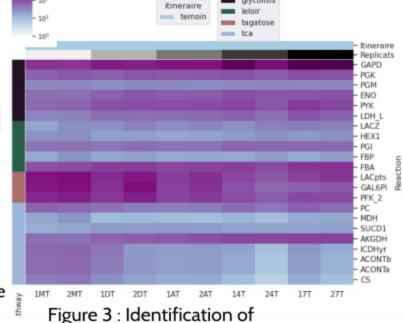


Figure 3 : Identification of preferential pathway for Ll --> uptake lactose first with tagatose then glycolysis pathway

### Discussion

- Identify **preferential pathways** thanks to metatranscriptomic data in the production of organoleptic compounds
- Decipher Interactions bacterium-bacterium within the bacterial community
- Salting variant has no effect in the production of organoleptic compounds

### References

- [1] Falentin et al, Agricultural and Food Chemistry, accepted
- [2] A. Varma et al, Appl. Environ. Microbiol., 1994
- [3] Mahadevan et al, biophys, 2002
- [4] Belcour et al, eLife, 2020
- [5] Himmi et al, Appl. Environ. Microbiol, 2000





