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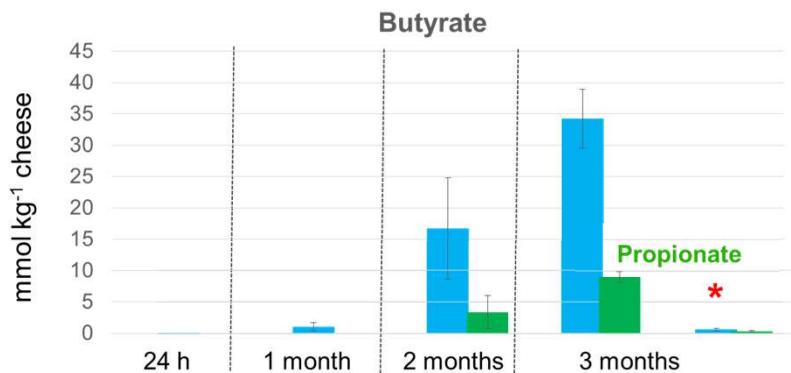
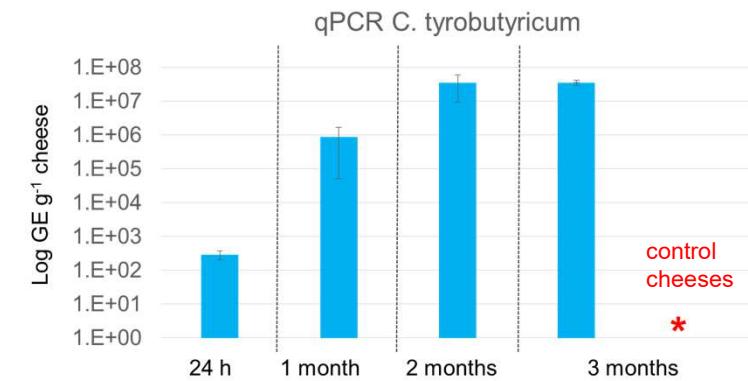
Deciphering the metabolism of *Clostridium tyrobutyricum* in cheese by transcriptomics

Emmanuelle Arias-Roth

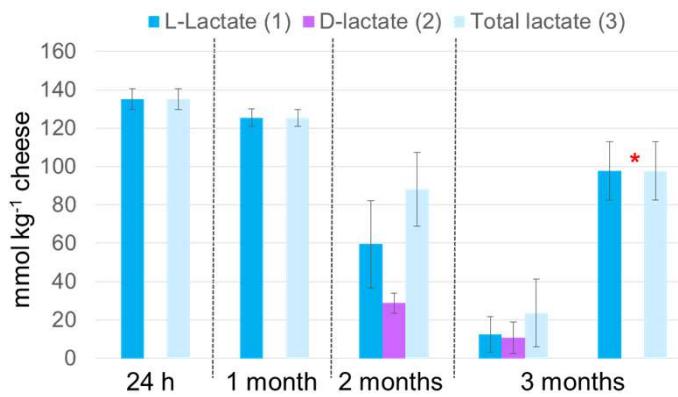
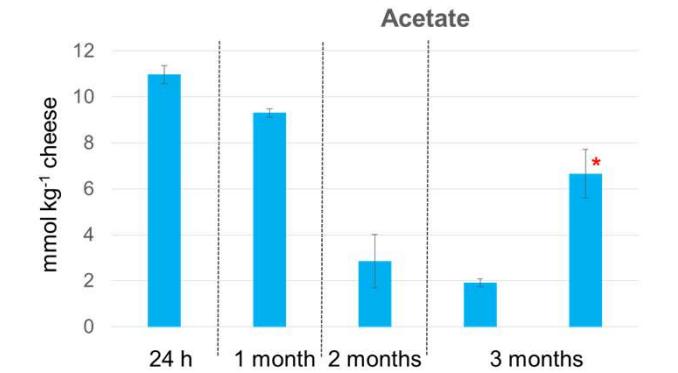
M. Storari, H. Berthoud, D. Wüthrich, S. Irmler



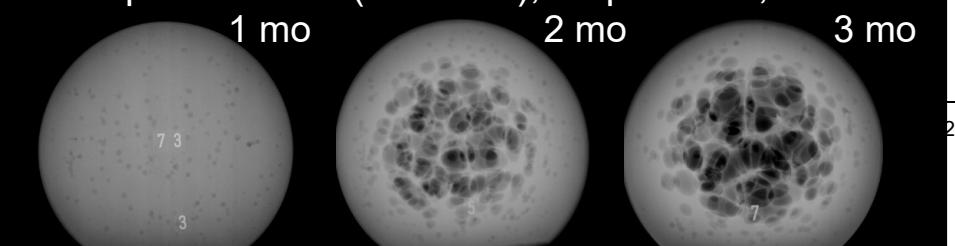
Growth of *C. tyrobutyricum* in cheese



+ other metabolites (Gómez-Torres, 2015)



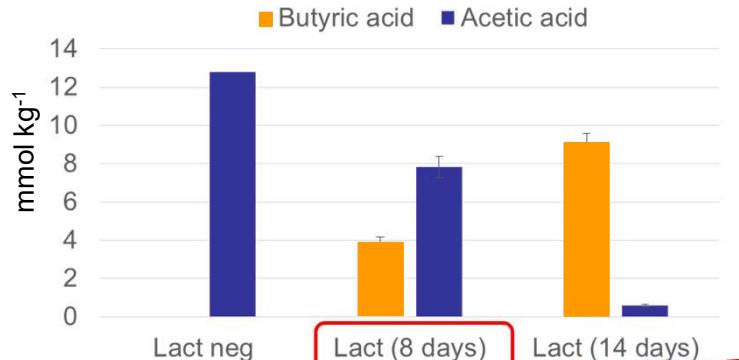
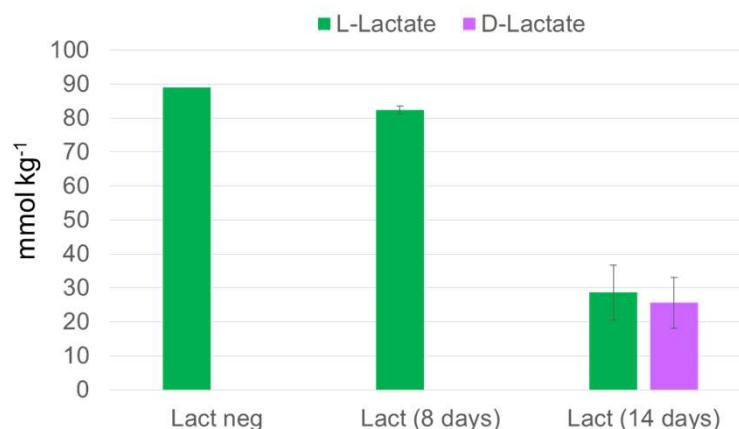
mesophilic starter (L-lactate), 3 spores/ml, 5 strains





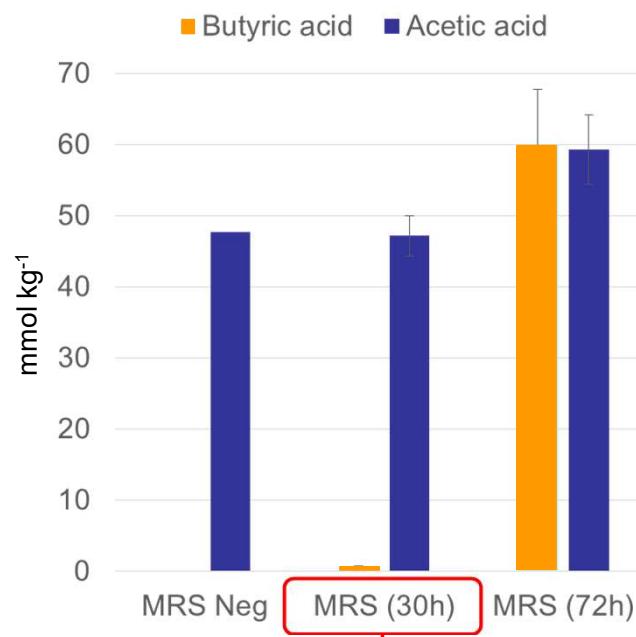
Growth of *C. tyrobutyricum* FAM22553 *in vitro*

Defined medium mimicking cheese
(L-lactate, acetate)



MRS
(glucose, acetate, yeast extract,...)

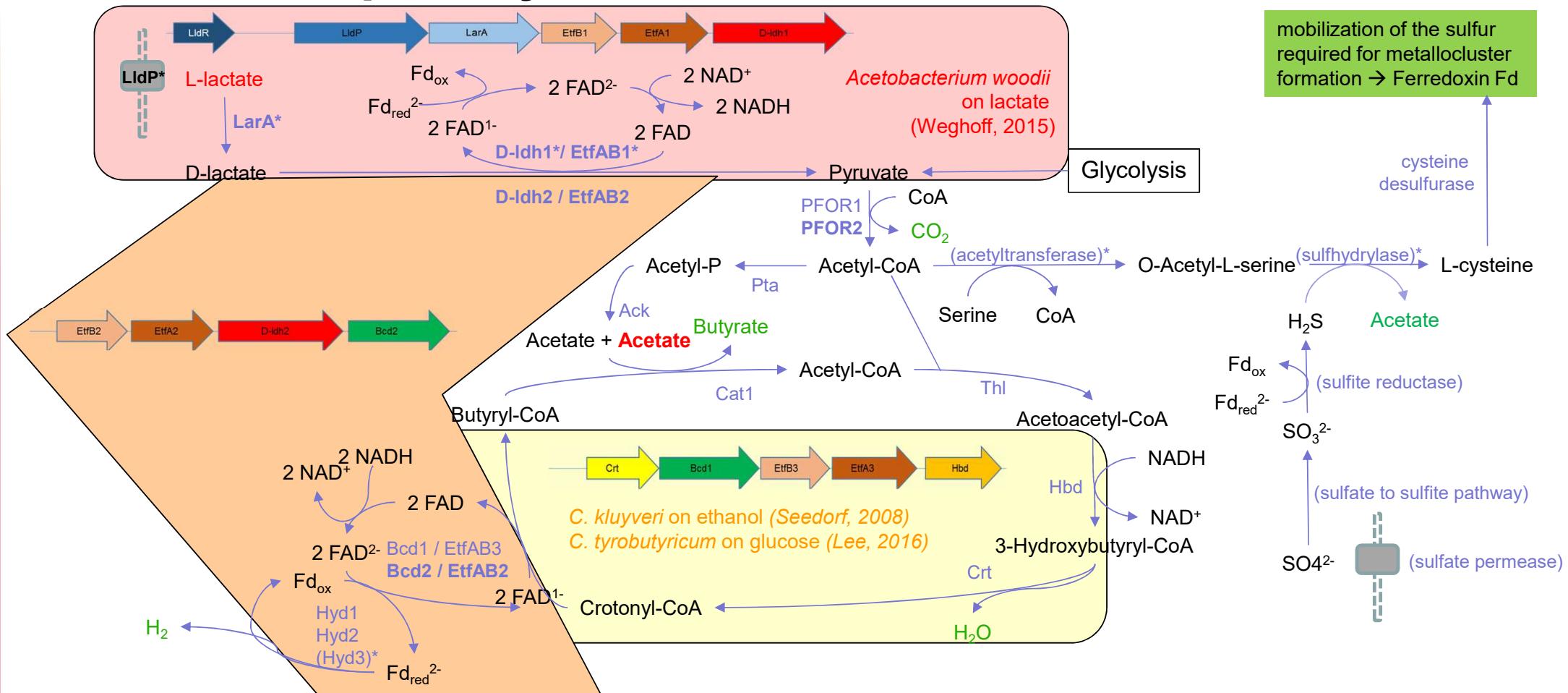
- no lactate detected, 0.4 mol/L propionate



RNA-Seq



RNA-seq – butyrate metabolism - FAM22553





Conclusion

Metabolism of *C. tyrobutyricum* in cheese = what did we learn from the transcriptomic study?

- Enzymes involved in the consumption of lactate: LarA, D-Ldh
- serine and methionine / H₂S seem to be key compounds for growth of *C. tyrobutyricum* in cheese
- Acetate producing starter and non-starter LAB promote late blowing (heterofermentative and citrate metabolizing LAB, propionic acid bacteria)



Thank you for your attention

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