

# Identifying chain elongation processes during the mixed-culture fermentation of proteins

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## HIGHLIGHTS:

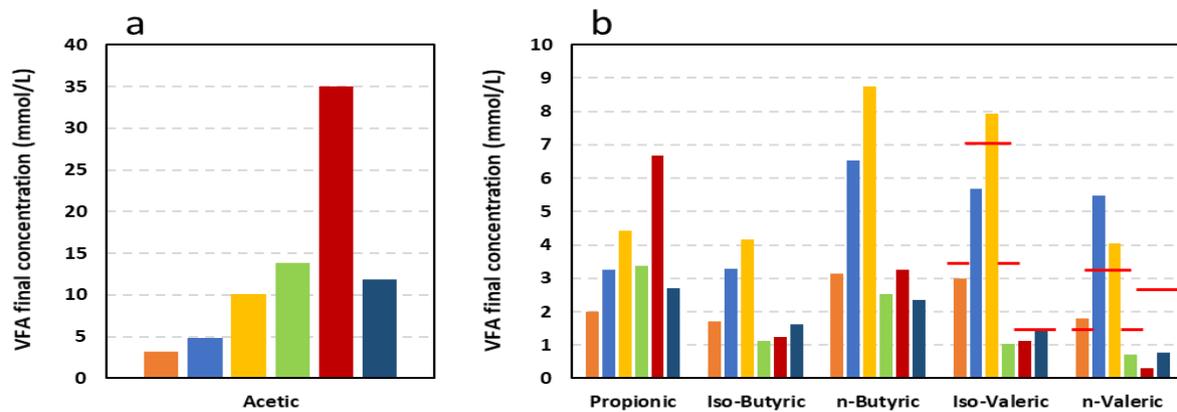
- Chain elongation was identified for the first time as a relevant process occurring in protein mixed-culture fermentation
- No external electron donor compound is required for the VFA elongation from proteins
- Its feasibility depends on both the chosen pH setpoint and protein composition

**BACKGROUND:** the chain elongation processes occurring during the fermentation of carbohydrate-rich residual streams have already been thoroughly studied, with multiple examples available in literature (Angenent et al., 2016). On the contrary, little is known on the role of proteins as substrates. The only previous studies focusing on amino acid-based chain elongation were performed with pure culture of a bacterium isolated from bovine rumen, *Eubacterium pyruvativorans* (Wallace et al., 2003), which consumes short chain VFAs to elongate butyric to caproic acid, using amino acids (e.g. alanine and leucine) as electron donor compounds (Wallace et al., 2004). Conversely, the feasibility of chain elongation process in mixed-culture microbiomes fermenting proteins has not been described before.

Hence, the aim of the present study was to verify the potential of proteins as a single substrate for the acidification and condensation of longer chain carboxylates while evaluating the related mechanisms and the required operational conditions.

**METHODOLOGY:** integrating the results of a previous study with two different proteins, casein and gelatin (Bevilacqua et al., 2020), two fermentation batch tests were performed at pH 5 using casein as the sole carbon source. The chosen substrate-to-inoculum ratio (SIR) was of 20 g COD protein/g VSS, with macronutrients being added accordingly, while the results from the previous study were obtained with a SIR equal to 10. Acetic acid was supplemented to one of the batch tests (approx. 10 mmol/L) to better understand the role of short chain carboxylates as electron acceptor compounds in protein-based chain elongation processes.

42 **RESULTS AND DISCUSSION:** Both tests lasted 384 hours, leading to the  
 43 final products spectra (mmol/L basis) observable in figure 1. In both  
 44 cases, the main products were n-butyric and iso-valeric acid (Fig. 1b),  
 45 while n-valeric acid production depended on whether acetic acid was  
 46 supplemented at the beginning of the batch operation.



47 **Figure 1.** VFA final concentration obtained in batch fermentation tests  
 48 performed with casein and gelatin at different pH values (a: acetic acid; b:  
 49 remaining VFAs). ■ Casein pH 5 SIR10; ■ Casein pH 5 SIR20; ■ Casein pH  
 50 5 SIR20 with acetic acid supplementation; ■ Casein pH 7 SIR10; ■ Gelatin  
 51 pH 7 SIR10; ■ Gelatin pH 5 SIR10. The red horizontal bars illustrate the  
 52 maximum theoretical production of iso and n-valeric acids based on the  
 53 chosen substrate.

54 Consumption of acetic and propionic acid in the first stages (24-72 hours)  
 55 of the casein batch tests was observed (concentration profile not shown  
 56 here), suggesting the existence of chain elongation processes.  
 57 Furthermore, no external electron donor supplementation was required, as  
 58 amino acids probably reacted with in-situ produced VFAs. The acetic acid  
 59 supplementation test helped identify the elongation process as the final  
 60 concentration of this acid was comparable to the initial one. In fact, the 5  
 61 mmol/L produced in the SIR 20 test plus the 10 mmol/L of acetic acid  
 62 supplemented should have led to a total 15 mmol/L at the end of this test.  
 63 As only 10 mmol/L were measured (Fig. 1a), it indicates that some 5  
 64 mmol/L of may have been used for chain elongation. The acetic acid  
 65 supplementation also increased the overall casein conversion and diverted  
 66 the condensation towards iso-valeric acid and possibly n-butyric acid. Still,  
 67 protein-based fermentation appears to be especially selective towards n-  
 68 valeric acid as its production was always greater than theoretically  
 69 possible by acidification only in both tests (Fig. 1b), given that this VFA  
 70 only originates from a well identified amino acid, proline (Bevilacqua et  
 71 al., 2020; Regueira et al., 2020). Comparing these results with the ones  
 72 obtained from the fermentation of different proteins (Bevilacqua et al.,  
 73 2020), it appears that protein composition and pH are fundamental in  
 74 determining the feasibility of the chain elongation. The process was only  
 75 identified during casein fermentation at low pH (5.0); gelatin did not  
 76 undergo chain elongation during its anaerobic conversion to VFAs (Fig.

77 1b). The relative abundance of amino acids acting as electron donors  
78 might explain the differences according to the substrate.

79 **CONCLUSION:** to the best of our knowledge, chain elongation was  
80 identified for the first time during mixed-culture fermentation of proteins.  
81 This kind of process is especially attractive as it does not require electron  
82 donor supplementation as in most cases described in literature. Also, its  
83 feasibility depends on the environmental conditions (i.e. low pH) and the  
84 protein composition. This work helps to shed some light on amino acid-  
85 based chain elongation in mixed microbiomes while constituting a starting  
86 point for further studies on the subject.

## 87 REFERENCES

- 88 1. Angenent, L.T., Richter, H., Buckel, W., Spirito, C.M., Steinbusch,  
89 K.J.J., Plugge, C.M., Strik, D.P.B.T.P., Grootsholten, T.I.M.,  
90 Buisman, C.J.N., Hamelers, H.V.M. Chain elongation with reactor  
91 microbiomes: open-culture biotechnology to produce biochemicals.  
92 *Environ. Sci. Technol.*, 2016, 50, 2796–2810.
- 93 2. Bevilacqua, R., Regueira, A., Mauricio-Iglesias, M., Lema, J.M.,  
94 Carballa, M. Steering the conversion of protein residues to short  
95 chain carboxylates by adjusting pH. Submitted
- 96 3. Regueira, A., Lema, J.M., Carballa, M., Mauricio-Iglesias, M.  
97 Metabolic modeling for predicting VFA production from protein-rich  
98 substrates by mixed-culture fermentation. *Biotechnology and*  
99 *Bioengineering*, 2020, 117, 73-84.
- 100 4. Wallace, R. J., Chaudhary, L. C., Miyagawa, E., McKain, N., Walker,  
101 N. D. Metabolic properties of *Eubacterium pyruvatorans*, a ruminal,  
102 "hyper-ammonia-producing," anaerobe with metabolic properties  
103 analogous to those of *Clostridium kluyveri*. *Microbiology*, 2004, 150  
104 (9) 2921– 2930.
- 105 5. Wallace, R. J., McKain, N., McEwan, N.R., Miyagawa, E., Chaudhary,  
106 L.C., King, T.P., Walker, N.D., Apajalahti, J.H.A., Newbold, C.J.  
107 *Eubacterium pyruvatorans* sp. nov., a novel non-saccharolytic  
108 anaerobe from the rumen that ferments pyruvate and amino acids,  
109 forms caproate and utilizes acetate and propionate. *Microbiology*,  
110 2003, 53 (4), 965-970.