



UNIVERSITY OF LEEDS

This is a repository copy of *Biotransformation of organic acids to their corresponding alcohols by Thermoanaerobacter pseudoethanolicus*.

White Rose Research Online URL for this paper:
<http://eprints.whiterose.ac.uk/145430/>

Version: Accepted Version

Article:

Scully, SM, Brown, A, Ross, AB et al. (1 more author) (2019) Biotransformation of organic acids to their corresponding alcohols by *Thermoanaerobacter pseudoethanolicus*. *Anaerobe*, 57. pp. 28-31. ISSN 1075-9964

<https://doi.org/10.1016/j.anaerobe.2019.03.004>

(c) 2019, Elsevier Ltd. This manuscript version is made available under the Creative Commons CC-BY-NC-ND 4.0 license <https://creativecommons.org/licenses/by-nc-nd/4.0/>.

Reuse

This article is distributed under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs (CC BY-NC-ND) licence. This licence only allows you to download this work and share it with others as long as you credit the authors, but you can't change the article in any way or use it commercially. More information and the full terms of the licence here: <https://creativecommons.org/licenses/>

Takedown

If you consider content in White Rose Research Online to be in breach of UK law, please notify us by emailing eprints@whiterose.ac.uk including the URL of the record and the reason for the withdrawal request.



eprints@whiterose.ac.uk
<https://eprints.whiterose.ac.uk/>

1 **Short running title: SCFA to Alcohols by *Thermoanaerobacter***

2 **Biotransformation of organic acids to their corresponding**
3 **alcohols by *Thermoanaerobacter pseudoethanolicus***

4

5 **Sean M. Scully¹**
6 **Aaron Brown²**
7 **Andrew B. Ross²**
8 **Johann Orlygsson^{1,#}**

9

10

11 ¹*Faculty of Natural Resource Science, University of Akureyri, Borgir v. Nordurslod, 600 Akureyri, Iceland.*

12 ²*School of Chemical and Process Engineering, University of Leeds, Leeds, LS2 9JT, United Kingdom.*

13

14 *E-mail: jorlygs@unak.is (corresponding author)*

15

16

17 **Keywords:** butanol, propanol, hexanol, bioreduction, thermophiles, biocatalysis

18

19

20 **ABSTRACT**

21 Higher order alcohols, such as 1-butanol and 1-hexanol, have a large number of applications but
22 are currently prepared from non-renewable feedstocks. Here, the ability of *Thermoanaerobacter*
23 *pseudoethanolicus* to reduce short-chain fatty acids to their corresponding alcohols using
24 reducing potential generated by glucose catabolism with yields between 21.0 and 61.0%. ¹³C-
25 labelled acetate, 1-propionate and 1-butyrate demonstrates that exogenously added fatty acids are
26 indeed reduced to their corresponding alcohols. This mode of producing primary alcohols from
27 fatty acids using a thermophilic anaerobe opens the door for the production of such alcohols
28 from low-value materials using an inexpensive source of reducing potential.

29

30

31 Organisms within the genera of *Thermoanaerobacter* and *Caldanaerobacter* have broad
32 metabolic capabilities and are noted for their biotechnological potential such as their highly
33 thermotolerant nature and ability to produce biofuels such as ethanol from a broad range of
34 monosaccharides and branched-chain alcohols from branched-chain amino acids [1–5]. Also,
35 *Thermoanaerobacter* species have been exploited for their thermostable enzymes such as
36 xylanases, pectases, and amylases, and the utility of their enzymes in organic synthesis [6–9].
37 The alcohol dehydrogenases (ADHs) of *Thermoanaerobacter* species have been of particular
38 interest; *Thermoanaerobacter pseudoethanolicus* 39E possess several ADHs, each with varying
39 substrate specificity and cofactor preference [6]. One of the functions of these multiple ADHs
40 seems to be to transfer an electron from NAD(P)H to NAD(P)⁺ which can be used for the
41 reduction of acetaldehyde to ethanol. The use of *Thermoanaerobacter* (formerly
42 *Thermoanaerobium*) *brockii* ADHs have previously been commercially available and applied to

43 the enantioselective biological reductions of ketones to their corresponding secondary alcohols
44 [8,10,11]. However, the use of thermophilic *Clostridia* has not previously been investigated for
45 their ability to produce higher-order alcohols (C3-C7) from short-chain fatty acids (SCFAs)
46 although the addition of acetate to the fermentation medium is known to stimulate ethanol
47 production [12]. Beyond the production of 1-butanol via the acetone-butanol-ethanol (ABE)
48 process [13,14] and some limited work on the production 1-hexanol from the fermentation of
49 syngas [15–17], there has been little focus on the production of C3 and larger primary alcohols.
50 Historically, such alcohols are prepared using the oxo process or by the reduction of carboxylic
51 acids using strong reducing agents, both of which have substantial drawbacks. There have been
52 recent reports of *Clostridium saccharoperbutylacetonicum* N1-4 to convert 1-butyric acid to 1-
53 butanol in the presence of glucose in the context of ABE fermentations [18,19] as well as by
54 acetogens such as “*Clostridium ragsdaleii*” [20] and engineered *Clostridium autoethanogenum*
55 [21] and in mixed culture with propionate-producing *Clostridium propionicum* and
56 *Alkalibaculum bacchi* [22].

57
58 Recently, *Thermoanaerobacter* strains have been shown to produce branched-chain fatty acids
59 and alcohols during the fermentation of branched-chain amino acids [23–25] although no wild
60 type strains have been reported to produce 1-butanol or higher alcohols as a product of glucose
61 fermentation while the mechanism behind branched-chain alcohols formation from amino acids
62 remains unclear.

63
64 The present study investigates the ability of *Thermoanaerobacter pseudoethanolicus* (DSM
65 2355) to reduce C1-C8 short-chain fatty acids (SCFAs) to their corresponding alcohols using

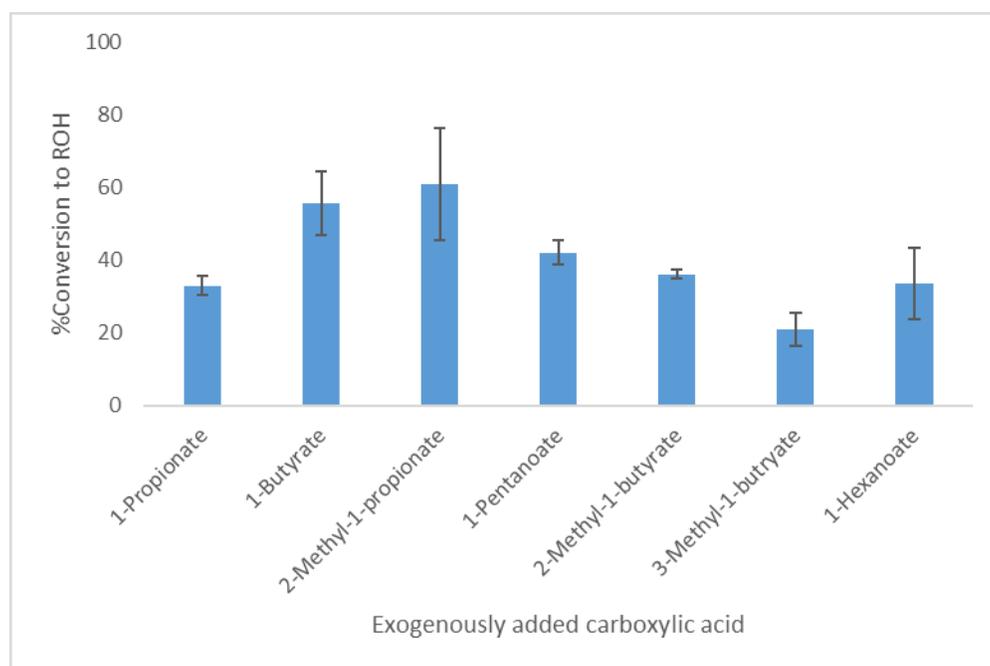
66 glucose as a carbon source and as a source of reducing potential. *T. pseudoethanolicus* was
67 cultivated anaerobically with nitrogen in the gas phase in Hungate tubes (16x150 mm)
68 containing 8.3 mL of Basal Mineral (BM) medium [26] containing 20 mM of glucose and 20
69 mM of the SCFA. Cultivations were performed at 65°C and pH 7 in triplicate without agitation
70 for 5 days. All materials were purchased from Sigma Aldrich except for ¹³C-labelled compounds
71 (Cambridge Isotope Laboratories, MA, USA).

72
73 Hydrogen was quantified by gas chromatograph equipped with a thermoconductivity detector
74 while short-chain fatty acids and alcohols were quantified by gas chromatograph with a flame
75 ionization detector as described previously [26]. The products were separated on a
76 nitroterephthalic-acid-modified polyethylene glycol (DB-FFAP, J&W Scientific) 30 m capillary
77 column, 0.32 i.d., 0.25 µm film thickness, using a temperature program of 40°C, hold time 1 min,
78 ramped to 240°C (10°C/min), hold time 10 min and a constant column head space pressure of 9.9
79 kPa. Optical densities were measured after 5 days of cultivation using a Shimadzu UV-1800 at
80 600 nm (*l*=1 cm). The mass spectrometer ion source was set to 260°C and the interface 250°C,
81 scanning took place once every 0.30 seconds in the range of 40 to 60 m/z. Peaks were identified
82 using the NIST mass spectral database, versions 147 and 27, with an identity threshold cut-off of
83 80. Samples were filtered prior to injection using a 0.45 µm syringe filter. ¹³C nuclear magnetic
84 resonance (NMR) spectroscopy of fermentations containing ¹³C1-labelled SCFAs were
85 performed on a Bruker AV400 NMR spectrometer at 298K after spiking with D₂O to obtain a
86 signal lock (0.3 mL addition to 1 mL of aqueous sample).

87

88 The ability of *T. pseudoethanolicus* to reduce short-chain carboxylic acids was evaluated by
89 cultivation on single SCFAs and 20 mM of glucose and detection of the corresponding alcohol
90 by gas chromatograph with flame ionizing detector. The ability of *T. pseudoethanolicus* to
91 reduce fatty acids in the presence of glucose was evaluated in batch culture after 5 days (Figure
92 1); the addition of heptanoate, and octanoate was also attempted although alcohol end products
93 were not detected above background (Supplemental Table 1). The data shows that the addition of
94 external fatty acids increase the final optical density in the cultures (above controls) and carbon
95 recoveries were between 70.8 and 104.4% (Supplemental Table 1).

96



97

98

99 **Figure 1-** Conversion of exogenously added SCFAs to fermentations containing 20 mM of
100 glucose by *T. pseudoethanolicus*. Values represent the average of triplicates with error bars
101 presented as standard deviation.

102

103 Apart from alcohol formation from the conversion of C2-C6 fatty acids, a peak attributable to
104 methanol was detected when formate was added exogenously although it could not be quantified
105 due to peak shouldering with ethanol. As ethanol is a normal end product of glucose
106 fermentation from *T. pseudoethanolicus*, ethanol formation from acetate was unclear. SCFAs
107 (C3 to C6) were converted to their corresponding alcohols with yields between 21.0 and 61.0%
108 (Figure 1). For instance, the addition of 1-propionate gave a 1-propanol titer of 6.6 mM (33 %)
109 and a 2-methyl-1-propionic acid addition yielded 12.2 mM of 2-methyl-1-propanol (61%).
110 Alcohol formation from 1-hexanoic acid was observed by GC-FID with the appearance of a peak
111 attributable to 1-hexanol which was confirmed by GC-MS (Supplemental Figure 1). As end
112 products were only analyzed after 5 days, the rates of carboxylic acid conversion to alcohols and
113 the impact that the addition of these exogenously added acids may have on growth is not
114 available. However, the decreased concentration of ethanol in the presence of carboxylic acids
115 suggest that electron flow from glucose is redirected to the corresponding alcohols.

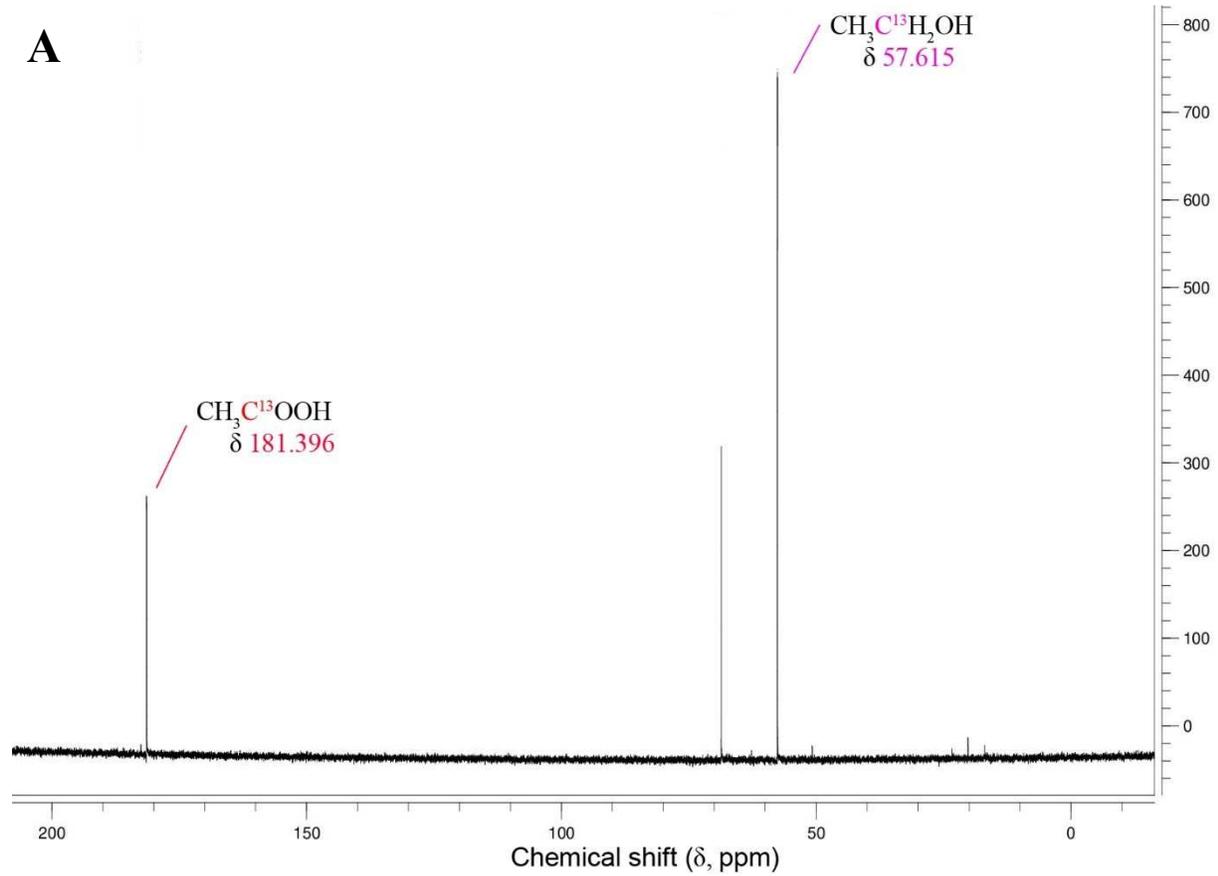
116

117 To confirm that the organic acid was indeed reduced to the corresponding alcohol, ¹³C1-acetate,
118 propionate, and butyrate were added at a concentration of 20 mM in addition to 20 mM of
119 glucose. Figure 2A-C shows the NMR spectra of fermentations containing exogenously added
120 ¹³C1 acetate (181.4 ppm), ¹³C1 propionate (181.3 ppm) and ¹³C1 butyrate (184.0 ppm),
121 respectively. After fermentation with glucose, new peaks of alcohol formation were observed
122 (ethanol at 57.6 ppm, propanol at 63.8 ppm and butanol at 63.7 ppm). This supports that the
123 exogenously added SCFAs are indeed being reduced to ethanol rather than appearing direct as
124 end products of glucose fermentation.

125

126 **Figure 2** – ^{13}C NMR spectra of glucose fermentations (20 mM) by *T. pseudoethanolicus* with
127 exogenously added ^{13}C 1-acetate (A), ^{13}C 1-propionate (B) and ^{13}C 1-butyrate (C) at 20 mM final
128 concentration.

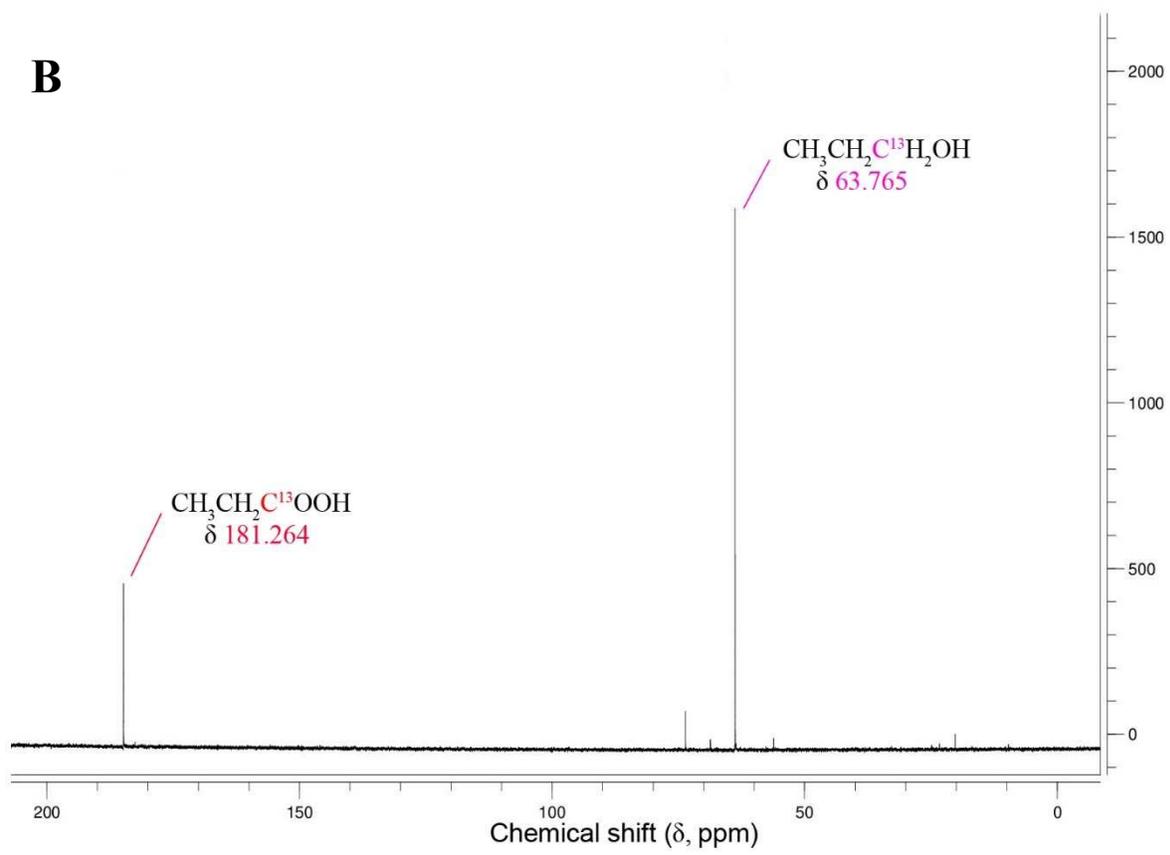
129



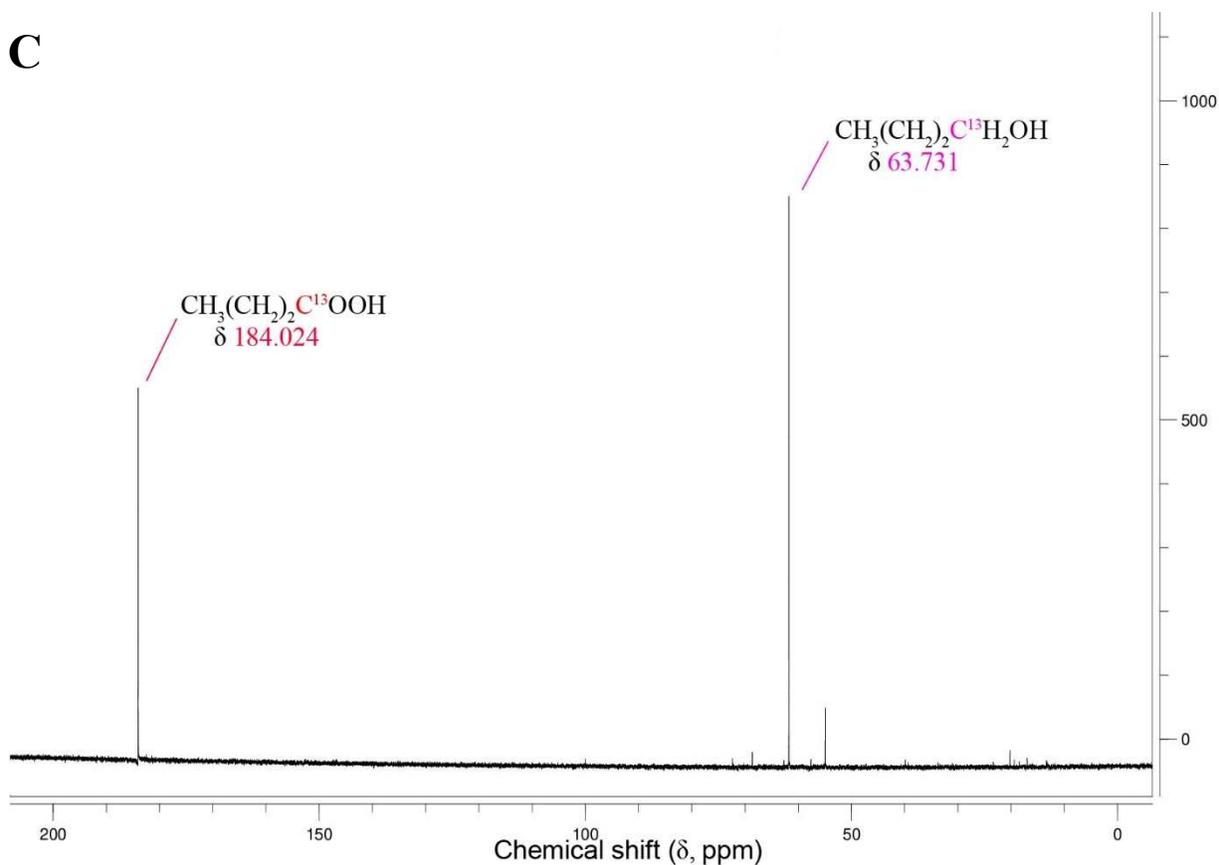
130

131

B



C



133

134

135 The ability to reduce carboxylic acids might suggest an alternative mechanism for the formation
136 of fusel alcohols (3-methyl-1-butanol, 2-methyl-1-butanol, and 2-methyl-1-propanol) during the
137 fermentation of branched chain amino acids [23–25]. The presence of such alcohols as end
138 products can alter fermentation kinetics as well as cell viability due to disruption of the cellular
139 membrane, particularly those with a more non-polar character. Thus, higher order alcohols are
140 often more inhibitory as compared to ethanol [27]. While *Thermoanaerobacter*
141 *pseudoethanolicus* is known to tolerate up to 2.5% ethanol although growth rates are diminished
142 with 1-butanol and 2-methyl-1-butanol having a noticeable impact on growth rates at less than
143 1.0% v/v [28].

144

145 The capability of *Thermoanaerobacter* species to reduce carboxylic acids in the presence of
146 glucose could present a novel route to the production of higher-order alcohols from inexpensive
147 feedstocks using the reducing power of glucose or other inexpensive materials.
148 *Thermoanaerobacterium* strains have been noted to rapidly reduce ¹³C-labeled exogenously
149 added acetate to ethanol via an acetyl-CoA intermediate [29]. Carboxylic acid reassimilation
150 may provide insight as to why some *Thermoanaerobacterium* and *Thermoanaerobacter* strains
151 are such efficient ethanol producers during glucose fermentation. To the author's knowledge, this
152 is the first demonstration of the ability of a whole cell system of *Thermoanaerobacter* to convert
153 carboxylic acids to their corresponding alcohols using inexpensive carbohydrates as a source of
154 reducing potential and could present a novel route to the production of higher primary alcohols,
155 including branched-chain alcohols, from renewable substrates.

156

157 REFERENCES

- 158 [1] T. Chades, S.M. Scully, E.M. Ingvadottir, J. Orlygsson, Fermentation of Mannitol
159 Extracts From Brown Macro Algae by Thermophilic Clostridia, *Front. Microbiol.* 9
160 (2018) 1–13. doi:10.3389/fmicb.2018.01931.
- 161 [2] S.M. Scully, J. Orlygsson, Branched-chain alcohol formation from branched-chain amino
162 acids by *Thermoanaerobacter brockii* and *Thermoanaerobacter yonseiensis*, *Anaerobe.* 30
163 (2014) 82–84. doi:10.1016/j.anaerobe.2014.09.003.
- 164 [3] R.U. Onyenwoke, V. V. Kevbrin, A.M. Lysenko, J. Wiegel, *Thermoanaerobacter*
165 *pseudethanolicus* sp. nov., a thermophilic heterotrophic anaerobe from Yellowstone
166 National Park, *Int. J. Syst. Evol. Microbiol.* 57 (2007) 2191–2193.
167 doi:10.1099/ijs.0.65051-0.

- 168 [4] S. Scully, J. Orlygsson, Recent Advances in Second Generation Ethanol Production by
169 Thermophilic Bacteria, *Energies*. 8 (2015) 1–30. doi:10.3390/en8010001.
- 170 [5] M.P. Taylor, K.L. Eley, S. Martin, M.I. Tuffin, S.G. Burton, D.A. Cowan, Thermophilic
171 ethanogenesis: future prospects for second-generation bioethanol production., *Trends*
172 *Biotechnol.* 27 (2009) 398–405. doi:10.1016/j.tibtech.2009.03.006.
- 173 [6] D. Burdette, J.G. Zeikus, Purification of acetaldehyde dehydrogenase and alcohol
174 dehydrogenases from *Thermoanaerobacter ethanolicus* 39E and characterization of the
175 secondary-alcohol dehydrogenase (20 Adh) as a bifunctional alcohol dehydrogenase-
176 acetyl-CoA reductive thioes, *Biochem. J.* 302 (1994) 163–170.
- 177 [7] M. V Sizova, J. a Izquierdo, N.S. Panikov, L.R. Lynd, Cellulose- and xylan-degrading
178 thermophilic anaerobic bacteria from biocompost., *Appl. Environ. Microbiol.* 77 (2011)
179 2282–91. doi:10.1128/AEM.01219-10.
- 180 [8] C.-H. Wong, G.M. Whitesides, *Enzymes in Synthetic Organic Chemistry*, Pergamon,
181 Oxford, 1994.
- 182 [9] M.K. Bhat, Cellulases and related enzymes in biotechnology., *Biotechnol. Adv.* 18 (2000)
183 355–83. <http://www.ncbi.nlm.nih.gov/pubmed/14538100>.
- 184 [10] K. Faber, *Biotransformations in Organic Chemistry: A Textbook*, 5th ed., Springer, New
185 York, 2011.
- 186 [11] G.M. Whitesides, C.-H. Wong, Enzyme in der organischen Synthese, *Angew. Chemie.* 97
187 (1985) 617–720.
- 188 [12] Q. He, P.M. Lokken, S. Chen, J. Zhou, Characterization of the impact of acetate and
189 lactate on ethanolic fermentation by *Thermoanaerobacter ethanolicus*, *Bioresour.*
190 *Technol.* 100 (2009) 5955–65. doi:10.1016/j.biortech.2009.06.084.

- 191 [13] D.T. Jones, D.R. Woods, Acetone-Butanol Fermentation, *Microbiol. Rev.* 50 (1986) 484–
192 524. doi:3540574.
- 193 [14] O. Tirado-Acevedo, M.S. Chinn, A.M. Grunden, Production of biofuels from synthesis
194 gas using microbial catalysts., in: *Adv. Appl. Microbiol.*, 1st ed., Elsevier Inc., 2010: pp.
195 57–92. doi:10.1016/S0065-2164(10)70002-2.
- 196 [15] H. Richter, B. Molitor, M. Diender, D.Z. Sousa, A Narrow pH Range Supports Butanol,
197 Hexanol, and Octanol Production from Syngas in a Continuous Co-culture of *Clostridium*
198 *ljungdahlii* and *Clostridium kluveri* with In-Line Product Extraction, *Front. Microbiol.* 7
199 (2016) 1–13. doi:10.3389/fmicb.2016.01773.
- 200 [16] Á. Fernández-Naveira, M.C. Veiga, C. Kennes, H-B-E (hexanol-butanol-ethanol)
201 fermentation for the production of higher alcohols from syngas/waste gas, *J. Chem.*
202 *Technol. Biotechnol.* 92 (2017) 712–731. doi:10.1002/jctb.5194.
- 203 [17] Y. Dekishima, E.I. Lan, C.R. Shen, K.M. Cho, J.C. Liao, Extending carbon chain length
204 of 1-butanol pathway for 1-hexanol synthesis from glucose by engineered *Escherichia*
205 *coli*, *J. Am. Chem. Soc.* 133 (2011) 11399–11401. doi:10.1021/ja203814d.
- 206 [18] S. Baba, Y. Tashiro, H. Shinto, K. Sonomoto, Development of high-speed and highly
207 efficient butanol production systems from butyric acid with high density of living cells of
208 *Clostridium saccharoperbutylacetonicum*, *J. Biotechnol.* 157 (2012) 605–612.
209 doi:10.1016/j.jbiotec.2011.06.004.
- 210 [19] N.K.N. Al-Shorgani, E. Ali, M.S. Kalil, W.M.W. Yusoff, Bioconversion of Butyric Acid
211 to Butanol by *Clostridium saccharoperbutylacetonicum* N1-4 (ATCC 13564) in a Limited
212 Nutrient Medium, *Bioenergy Res.* 4 (2012) 287–293. doi:10.1007/s12155-011-9126-6.
- 213 [20] C.E. Isom, M.A. Nanny, R.S. Tanner, Improved conversion efficiencies for *n*-fatty acid

214 reduction to primary alcohols by the solventogenic acetogen “*Clostridium ragsdalei*,” J.
215 Ind. Microbiol. Biotechnol. 42 (2015) 29–38. doi:10.1007/s10295-014-1543-z.

216 [21] F. Liew, A.M. Henstra, M. Köpke, K. Winzer, S.D. Simpson, N.P. Minton, Metabolic
217 engineering of *Clostridium autoethanogenum* for selective alcohol production, Metab.
218 Eng. 40 (2017) 104–114. doi:10.1016/j.ymben.2017.01.007.

219 [22] K. Liu, H.K. Atiyeh, B.S. Stevenson, R.S. Tanner, M.R. Wilkins, R.L. Huhnke, Mixed
220 culture syngas fermentation and conversion of carboxylic acids into alcohols, Bioresour.
221 Technol. 152 (2014) 337–346. doi:10.1016/j.biortech.2013.11.015.

222 [23] S.M. Scully, J. Orlygsson, Branched-chain alcohol formation from branched-chain amino
223 acids by *Thermoanaerobacter brockii* and *Thermoanaerobacter yonseiensis*, Anaerobe. 30
224 (2014) 82–84. doi:10.1016/j.anaerobe.2014.09.003.

225 [24] S.M. Scully, J. Orlygsson, Amino Acid Metabolism of *Thermoanaerobacter* Strain
226 AK90 : The Role of Electron-Scavenging Systems in End Product Formation, J. Amino
227 Acids. 2015 (2015). doi:10.1155/2015/410492.

228 [25] S.M. Scully, P. Iloranta, P. Myllymaki, J. Orlygsson, Branched-chain alcohol formation
229 by thermophilic bacteria within the genera of *Thermoanaerobacter* and
230 *Caldanaerobacter*., Extremophiles. 19 (2015) 809–18. doi:10.1007/s00792-015-0756-z.

231 [26] J. Orlygsson, S.R.B. Baldursson, Phylogenetic and physiological studies of four hydrogen-
232 producing thermoanareobes, Icelandic Agric. Sci. 20 (2007) 93–105.

233 [27] R.K. Bajapi, E.L. Iannotti, Product Inhibition, in: L.E. Erickson, D.Y.-C. Fung (Eds.),
234 Handb. Anaerob. Ferment., Marcel Dekker, Inc., New York, 1988: pp. 207–241.

235 [28] S. Huffer, M.E. Clark, J.C. Ning, H.W. Blanch, D.S. Clark, Role of Alcohols in Growth,
236 Lipid Composition , and Membrane Fluidity of Yeasts, Bacteria, and Archaea, Appl.

237 Environ. Microbiol. 77 (2011) 6400–6408. doi:10.1128/AEM.00694-11.

238 [29] A.J. Shaw, B.B. Miller, S.R. Rogers, W.R. Kenealy, A. Meola, A. Bhandiwad, W.R.

239 Sillers, I. Shikhare, D. a Hogsett, C.D. Herring, Anaerobic detoxification of acetic acid in

240 a thermophilic ethanologen, Biotechnol. Biofuels. 8 (2015) 1–12. doi:10.1186/s13068-

241 015-0257-4.

242