



Redox reactions in food fermentations

Hansen, Egon Bech

Published in:
Current Opinion in Food Science

Link to article, DOI:
[10.1016/j.cofs.2018.03.004](https://doi.org/10.1016/j.cofs.2018.03.004)

Publication date:
2018

Document Version
Peer reviewed version

[Link back to DTU Orbit](#)

Citation (APA):
Hansen, E. B. (2018). Redox reactions in food fermentations. *Current Opinion in Food Science*, 19, 98-103.
<https://doi.org/10.1016/j.cofs.2018.03.004>

General rights

Copyright and moral rights for the publications made accessible in the public portal are retained by the authors and/or other copyright owners and it is a condition of accessing publications that users recognise and abide by the legal requirements associated with these rights.

- Users may download and print one copy of any publication from the public portal for the purpose of private study or research.
- You may not further distribute the material or use it for any profit-making activity or commercial gain
- You may freely distribute the URL identifying the publication in the public portal

If you believe that this document breaches copyright please contact us providing details, and we will remove access to the work immediately and investigate your claim.

Accepted Manuscript

Title: Redox reactions in food fermentations

Author: Egon Bech Hansen

PII: S2214-7993(17)30103-0

DOI: <https://doi.org/doi:10.1016/j.cofs.2018.03.004>

Reference: COFS 345



To appear in:

Received date: 15-11-2017

Revised date: 27-2-2018

Accepted date: 3-3-2018

Please cite this article as: Hansen, E.B., Redox reactions in food fermentations, *COFS* (2018), <https://doi.org/10.1016/j.cofs.2018.03.004>

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

1 **Redox reactions in food fermentations**

2

3

4 Egon Bech Hansen

5

6

7

8 National Food Institute

9 Technical University of Denmark

10 Kemitorvet bldg.. 202

11 DK2800 Kgs. Lyngby

12 Denmark

13 Phone: +45 3588 6203

14 e-mail: egbh@food.dtu.dk

15

Accepted Manuscript

15

16 **Abstract**

17 Food fermentations are typically performed without actively supplying air. Except for possible surface
18 microorganisms, oxygen will only be transiently available and the redox reactions during the fermentation
19 need to be in balance. Production of ATP from fermentation of carbohydrates typically involves oxidative steps
20 in the early part of the pathways whereas a multitude of different reactions are used as compensating
21 reductions. Much of the diversity seen between food fermentations arise from the different routes and the
22 different electron acceptors used by microorganisms to counterbalance the initial oxidative steps.

23 This review gives a short overview of the routes employed by microorganisms in food fermentations to find
24 ultimate electron acceptors allowing them to balance their fermentative metabolism.

25 The diversity of acceptors used leads to diversity of metabolic end products and this contributes to the
26 diversity in flavor, color, texture, and shelf life. The review concludes that these reactions are still only
27 incompletely understood and that they represent an interesting area for fundamental research and also
28 represent a fertile field for product development through a more conscious use of the redox properties of
29 strains used to compose food cultures.

30

30

31 **Introduction**

32 Fermented foods have during centuries been produced without any knowledge of microbiology and even today
33 our knowledge of the beneficial microorganisms is still quite limited. More than 200 species of microorganisms
34 have a documented history of use in food fermentations [1] and a handful of those are produced and made
35 available as commercial starter cultures [2]. Fermented food spans a large range of products with the major
36 categories being: alcoholic beverages (beer and wine); fermented doughs; vinegar; fermented dairy products
37 (cheese, yoghurt, and fermented milks); fermented soy (miso, tempeh, natto, and soy sauce); fermented fish;
38 fermented meat; fermented coffee; and fermented cocoa [1]. *Sensu stricto* fermentation was defined as life
39 without air as opposed to respiration. However, the above list of microorganisms in fermented foods is based
40 on a less strict definition and include some aerobic microbial processes like production of vinegar and surface
41 ripening of cheese and sausages. Nevertheless, the majority of food fermentations are performed with no
42 supply of air.

43 It might seem surprising that imposing a limitation on the microbial metabolism by withholding air should lead
44 to an increased diversity of flavors and textures produced by the cultures. In comparison to a respiratory
45 metabolism which mainly produce CO₂ and water as end products, the anaerobic metabolisms give a wide
46 range of end products as ethanol, acetoin, diacetyl, acetaldehyde, lactic acid, acetic acid, and other acids in
47 addition to water and CO₂ [3].

48 Redox reactions are chemical reactions involving the transfer of electrons between molecules where the donor
49 molecule is said to be oxidized and the recipient reduced. Although the two reactions must be simultaneous,
50 they can, in a galvanic cell, be separated to occur at different electrodes. For each reaction, the standard
51 potential E^0 (measured in volts, V) defines the condition where electrons are gained or lost at equal rates. The
52 oxidation/reduction potential (ORP, E_h) of an aqueous system can be measured using a redox electrode [4]. The

53 value of E_n relative to E^0 will determine the tendency for a molecule to receive or donate electrons (to oxidize
54 or to be oxidized).

55 Lactic acid bacteria (LAB) play a prominent role in food fermentations with respect to volume, diversity of raw
56 materials, and diversity of species. Traditionally the primary performance parameter for starter cultures for the
57 food industry has been the acidification activity. The second parameter has been robustness towards phage
58 infections, which is another manifestation of reliability of acidification [2]. Texture and taste have been of
59 lower priority and for this reason less attention has been given to E_n compared to pH.

60 This review focus on redox reactions in LAB and the conclusion will be that innovation in food fermentations
61 can be dramatically stimulated with increased knowledge on redox reactions when composing the starter
62 cultures for food fermentation.

63

64 **Fermentation**

65 The fermentative metabolism is most easily understood by separating energy production from maintenance of
66 redox homeostasis. Off course in reality, this is not possible.

67 ATP is typically generated from metabolizing carbohydrates into pyruvate by oxidation. Different pathways to
68 pyruvate can be used depending on the organism and the sugar. Glycolysis by the Embden–Meyerhof–Parnas
69 pathway is a main route but the so-called hetero-fermentative pathways involving phosphoketolase enzymes in
70 key metabolic steps are also quite common [3]. The net gain of ATP differs between the pathways and the ATP
71 yield depends on the uptake mechanism and the length of the carbohydrate. Gänsle has recently reviewed the
72 main carbohydrate metabolisms of LAB [5]. The oxidation of carbohydrates to pyruvate consumes NAD in
73 addition to the production of ATP, and this redox-debt must be paid back. An additional reward in the form of
74 gaining extra ATP by shifting from ethanol to acetate production is available for the hetero-fermentative LAB if
75 they can mobilize extra NAD generating capacity [6–8].

76 Regeneration of NAD is accomplished by the concerted action of all cellular oxidoreductases. However, only
77 the ones having an available electron acceptor will contribute under any given condition. We have probably
78 only identified the most obvious electron acceptors as we tend to reduce complexity when we study microbial
79 metabolism. The discovery of relevant electron acceptors utilized during food fermentations will require the
80 researcher to use the relevant food as medium in the research.

81 Pyruvate is the primary electron acceptor for LAB. This is in fact what unifies the group of lactic acid bacteria,
82 they produce lactic acid as the major end product. Pyruvate is reduced to lactate by the enzyme lactate
83 dehydrogenase (LDH) with concomitant oxidation of NADH to NAD [3]. Homo-fermentative LAB rely mainly on
84 pyruvate and LDH for NAD regeneration [9]. NAD regeneration solely by LDH leaves very little flexibility in the
85 metabolic network and LAB will therefore benefit by having alternative routes to NAD regeneration and even
86 homo-fermentative LAB will usually possess alternative routes to regenerate NAD [10].

87 Oxygen, O₂, offers, if present, such an alternative to regenerate NAD by oxidizing NADH via the NADH oxidases,
88 NoxE or NoxAB [11–13]. Some LAB even have a rudimentary electron transport chain including cytochromes
89 [11,14–16]. Traditionally most LAB have been considered to be anaerobic and much research has been devoted
90 to study the relationship between LAB and oxygen from the angle of oxidative stress [12,17–19]. It was quite a
91 surprise when Duwatt et al in 2001 showed that *Lactococcus* is able to respire if hemin is supplied in the
92 medium [15]. It now seems clear that the LAB ancestors were aerobes and that the ability to respire has since
93 been lost to various degrees as a consequence of genome reduction in the course of specialization to the
94 nutrient rich ecological niches where LAB are commonly found [20–22]. In the light of this ancestry it is not
95 surprising that LAB are able to use oxygen when available and that they possess the functions allowing them to
96 deal with aerobic stress. It might therefore be fruitful to look at the redox reactions from the angle of
97 regeneration of NAD rather than aerobic stress management. In a nutrient rich environment, speed might be
98 more important than economy and oxygen might anyhow be the first “nutrient” to be depleted. By losing the

99 ability to use the entire chain of oxidative phosphorylation, LAB will use oxygen less efficiently and consume
100 more oxygen and thereby deplete oxygen faster. Rapid consumption of oxygen by LAB might confer an
101 advantage over aerobic bacteria, with which they are commonly in competition in food matrices. By
102 reorienting the metabolic pathways towards the use of additional electron acceptors, LAB might have become
103 better adapted to efficiently remove air and to live well without it. The diversity of the routes developed by
104 LAB to use alternative electron acceptors are illustrated in Figure 1 and several of the electron acceptors used
105 by some LAB are listed in Table 1 and described further in the following sections. However, an increased focus
106 on the positive aspects of oxygen in NAD regeneration should not lead to a neglect of the negative aspects of
107 reactive oxygen species.

108

109 **Alternative electron acceptors**

110 One way to gain more flexibility is to acquire pyruvate with no “NAD-debt” and this is the main benefit of
111 utilizing citrate. Several LAB are metabolizing citrate without generating ATP from the citrate to pyruvate
112 pathway [23]. However, as no NAD has been consumed, the pyruvate from citrate can be used with greater
113 flexibility than pyruvate from sugar metabolism. It can be reduced to lactate with concomitant NAD
114 production, or the pyruvate can be directed towards other products than lactate including ATP-generating
115 routes [23].

116 Sugars can be used as electron acceptors by several fermentative LAB leading to the production of sugar
117 alcohols as mannitol and erythritol [24–26]. Similarly, fumarate and malate can serve as electron acceptors and
118 be reduced to succinate [27].

119 Phenolics, which are frequently found in fruits, are generally antimicrobial but some LAB use phenolics as
120 electron acceptors; their growth are stimulated by phenolics and the ratio of fermentation end products is
121 altered [28,29]. Similarly, LAB able to use other molecules for NAD regeneration can probably be isolated from

122 nature or constructed by engineering to make LAB become a general tool for reductions in bio-refinery
123 processes [30].

124

125 **Inside or outside**

126 The location of the electron acceptor molecule would seem of minor importance as long as NAD is
127 regenerated. However, it is energetically favorable to keep the negatively charged electrons inside the cell and
128 the positively charged protons outside of the cell membrane [31]. If the electron acceptor is uncharged and
129 able to diffuse through the membrane, reducing on the inside is likely to be more favorable. Oxygen, O_2 , can be
130 reduced on the outside of the cell to O_2^- by direct reduction via menaquinones or reduced on the inside by
131 NoxE or NoxAB and cytochrome bc [32,33]. The cytochrome reaction is the most efficient as charge is
132 separated by releasing protons on the outside while reducing O_2 on the inside [20]. Fructose is another
133 example of a molecule, which can be imported for the purpose of being reduced to mannitol and then again
134 exported [27].

135 The bacterial membrane serves as the barrier over which a pH and charge difference builds an electrochemical
136 gradient able to drive transport and ATP production. In addition to carrying the energy potential, the
137 membrane also serves as a reservoir and buffer for redox-equivalents [34].

138

139 **Redox reactions on the membrane and cell wall**

140 Menaquinones and menaquinols serve as carriers of reducing equivalents between oxidoreductases located in
141 the membrane [34]. They constitute an important component in the electron transport chain in oxidative
142 phosphorylation; other components of the respiratory process are cytochrome-bd and NoxAB [16,22,35–38].
143 Due to the link to the respiratory pathway, research has focused on understanding the role of menaquinones in
144 oxygen metabolism and relief of oxidative stress. The role of menaquinones in the anaerobic metabolism has

145 been somewhat out of focus although it has been recognized that lactococci produce menaquinones in
146 anaerobic growth [39] and that the production is twofold higher during anaerobic conditions compared to
147 aerobic growth [16]. The experiments of Tachon, Brandsma, and Yvon [11] demonstrated that during
148 fermentation in milk rapid removal of oxygen by *Lactococcus lactis* is mainly accomplished by the NoxE enzyme
149 whereas menaquinones and NoxAB are responsible for maintaining the low redox potential during the
150 stationary (anaerobic) phase. The same authors also demonstrated that in *Lactococcus* menaquinones are
151 participating in redox reactions on either face of the bacterial membrane and that NoxAB can use other
152 electron acceptors than menaquinones [11]. This seems to indicate that lactococci mainly utilize menaquinones
153 and NoxAB when air is absent or scarce.

154 Menaquinones might be a vehicle to use extracellular electron acceptors as direct reduction of tetrazolium
155 salts and metal ions have been demonstrated [11,32]. Using an extracellular electron acceptor would appear to
156 be less favorable than using an intracellular one as export of an electron will reduce the electrochemical
157 gradient over the membrane. One would therefore expect that this option should be reserved for molecules
158 which cannot be imported or which are unfavorable to import. It is unclear if this route contributes to NAD
159 regeneration under fermentation of milk. If it does, the terminal electron acceptors in milk remain to be
160 determined. Lab species commonly used in dairy fermentations differ widely in their ability to lower the redox
161 potential during fermentation [40]. Also strains within the same species show large difference [11,41].

162 The thiol group of cysteine containing peptides and proteins can, similar to the menaquinones, serve as carriers
163 of electrons between different oxidoreductases. In the oxidized form, two cysteines are bridged covalently via
164 the sulfur atoms; whereas the sulfur atoms will be free thiol groups in the reduced form. The two cysteines can
165 be in the same polypeptide chain as in thioredoxin, or located on different molecules, or, as in glutathione,
166 between two identical molecules. Thioredoxin and glutathione participate in a variety of redox reactions
167 involving the formation and breakage of disulfide bridges [42]. The sulfur redox reactions are coupled to the

168 NAD/NADH catalyzed redox reactions through the enzymes thioredoxin reductase and glutathione reductase.
169 Neither thioredoxin nor glutathione are essential for *Lactococcus lactis* [43]. The maintenance of components
170 of the thioredoxin and glutathione systems without being essential could point towards a function in the
171 transport of electrons allowing efficient regeneration of NAD, i.e. to transport electrons towards an electron
172 acceptor. Michelon et al showed that the very low redox potential reached by *Lactococcus lactis* in MRS
173 medium is due to exofacial thiol groups of membrane proteins maintained in the reduced state [44]. The
174 authors found that only the exofacial thiol groups contributed to the low redox potential and that reduction of
175 media components did not contribute [44]. It is difficult to understand why such a system would be maintained
176 in the course of evolution if the only outcome would be a dead end for the electrons in the form of reduced
177 thiol groups at the surface of the cell. It would make more sense if the reduced thiol represented a channel
178 through which the electrons can flow towards an ultimate acceptor. Obviously, the MRS medium used did not
179 contain such an acceptor. If the dairy associated *Lactococcus lactis* strains have evolved to perform optimally in
180 milk, one would assume that milk would contain a final acceptor for electrons transferred via the exofacial thiol
181 groups. Milk proteins would seem to be the most likely candidates. Titration of free thiol groups during milk
182 fermentation could possibly reveal if milk contrary to MRS can serve as electron acceptor. To date, this analysis
183 has not been conducted yet. Interestingly the analysis for free thiol groups have been done during sourdough
184 fermentation and gave a clear difference. Sourdoughs fermented by Lactobacilli show a difference of 3-5 mM
185 of free thiol groups compared to chemically acidified doughs [45]. Interesting strategies to identify the
186 exofacial thiol groups in *Lactococcus lactis* have recently been described by C. Roussel [46].

187

188 **Perspectives for innovation on fermented food products through redox engineering**

189 A shift in focus from acidification activity towards diversity of food products could be released through a better
190 understanding of redox reactions. It is not surprising that the industrial implementation of aerobic respiring

191 LAB was used to increase the yield of the acidification activity without changing the actual food fermentation
192 [37]. It is, however, surprising that this shift in paradigm has not yet led to a creative use of air and other
193 electron acceptors in food fermentations.

194 In the applied field, it seems obvious to combine strains with different reducing potential to compose cultures
195 with new and improved properties regarding shelf life and flavor. An approach so far mainly used for sour
196 dough cultures [47] but likely to be productive for all food cultures including cultures for dairy. A wider use of
197 E_h measurement as a control parameter in food fermentations might also be useful as texture and taste have
198 been demonstrated to vary depending on the reduction potential [48,49].

199 Fundamental research on the transport of electrons over the membrane and on expanding the range of
200 identified electron acceptors in food products used for fermentation would seem worthwhile. It is surprising
201 that we do not know which milk molecules are used by *Lactococcus lactis* to reach the typical low redox
202 potential. It would be interesting to know if the disulfides of milk proteins serve as electron acceptors and to
203 investigate if there is a link to the proteolytic systems of dairy adapted LAB.

204

204

205

206 Table 1.

207 Molecules used as alternative electron acceptors by lactic acid bacteria

208

electron acceptor	reduced molecule	organism	reference
fructose	mannitol	<i>Lactobacillus sanfranciscensis</i> , <i>Lactobacillus pontis</i> , <i>Lactobacillus fermentum</i> , <i>Lactobacillus florum</i> , <i>Leuconostoc citreum</i> , <i>Leuconostoc pseudomesenteroides</i> , <i>Oenococcus oeni</i> , <i>Weissella paramesenteroides</i>	[25,27,50]
citrate	lactate	<i>Leuconostocs</i> , <i>Lactobacilli</i> , <i>Weissella</i> , <i>Lactococcus lactis subsp.</i> <i>diacetylactis</i>	[23,27,51]
fumarate	succinate	<i>Lactobacillus pontis</i> , <i>Lactobacillus reuteri</i> , <i>Lactobacillus amylovorus</i> , <i>Lactobacillus fermentum</i>	[27]
malate	succinate	<i>Lactobacillus pontis</i> , <i>Lactobacillus reuteri</i> , <i>Lactobacillus amylovorus</i> , <i>Lactobacillus fermentum</i>	[27]
glucose, fructose	erythritol	<i>Lactobacillus sanfranciscensis</i> , <i>Oenococcus kitaharae</i> , <i>Oenococcus oeni</i>	[26,27,50]
α -ketoglutarate	2-hydroxyglutarate	<i>Lactobacillus sanfranciscensis</i> , <i>Lactobacillus reuteri</i>	[52]
disulphides	thiols	<i>Lactobacillus sanfranciscensis</i> ,	[45]

phenolics: caffeic acid p-coumaric acid ferulic acid	dihydrocaffeic acid phloretic acid dihydroferulic acid ethylcatechol ethylphenol ethylguaiacol	<i>Lactobacillus plantarum</i> <i>Weissella cibaria</i> <i>Weissella confuse</i> <i>Lactobacillus brevis</i> <i>Lactobacillus curvatus</i> <i>Lactobacillus rossiae</i>	[28]
---	---	--	------

209

Accepted Manuscript

209

210 **Bibliography**

- 211 1. Bourdichon F, Casaregola S, Farrokh C, Frisvad JC, Gerds ML, Hammes WP, Harnett J, Huys G, Laulund S,
 212 Ouwehand A, et al.: **Food fermentations: Microorganisms with technological beneficial use.** *Int J Food*
 213 *Microbiol* 2012, **154**:87–97.
- 214 2. Hansen EB: **Starter Cultures: Uses in the Food Industry.** In Edited by Batt CA, Tortorello ML. Elsevier Ltd,
 215 Academic Press,; 2014:529–534.
- 216 3. Caplice E, Fitzgerald GF: **Food fermentations: Role of microorganisms in food production and**
 217 **preservation.** *Int J Food Microbiol* 1999, **50**:131–149.
- 218 4. Abraham S, Cachon R, Jeanson S, Ebel B, Michelon D, Aubert C, Rojas C, Feron G, Beuvier E, Gervais P, et
 219 al.: **A procedure for reproducible measurement of redox potential (E h) in dairy processes.** *Dairy Sci*
 220 *Technol* 2013, **93**:675–690.
- 221 5. Gänzle MG: **Lactic metabolism revisited: Metabolism of lactic acid bacteria in food fermentations and**
 222 **food spoilage.** *Curr Opin Food Sci* 2015, **2**:106–117.
- 223 6. Vedamuthu ER: **The Dairy Leuconostoc: Use in Dairy Products.** *J Dairy Sci* 1994, **77**:2725–2737.
- 224 7. Condon S: **Responses of lactic acid bacteria to oxygen.** *FEMS Microbiol Lett* 1987, **46**:269–280.
- 225 8. Ravyts F, Vuyst L De, Leroy F: **Bacterial diversity and functionalities in food fermentations.** *Eng Life Sci*
 226 2012, **12**:356–367.
- 227 9. Henriksen CM, Nilsson D: **Redirection of pyruvate catabolism in Lactococcus lactis by selection of**
 228 **mutants with additional growth requirements.** *Appl Microbiol Biotechnol* 2001, **56**:767–775.
- 229 10. de Felipe FL, Kleerebezem M, Vos WM de, Hugenholtz J: **Cofactor Engineering: a Novel Approach to**
 230 **Metabolic Engineering in Lactococcus lactis by Controlled Expression of NADH Oxidase.** *J Bacteriol*
 231 1998, **180**:3804–3808.
- 232 11. Tachon S, Brandsma JB, Yvon M: **NoxE NADH Oxidase and the Electron Transport Chain Are**
 233 **Responsible for the Ability of Lactococcus lactis To Decrease the Redox Potential of Milk.** *Appl Environ*
 234 *Microbiol* 2010, **76**:1311–1319.
- 235 12. Higuchi M, Yamamoto Y, Poole LB, Shimada M, Sato Y, Takahashi N, Kamio Y: **Functions of Two Types of**
 236 **NADH Oxidases in Energy Metabolism and Oxidative Stress of Streptococcus mutans.** *J Bacteriol* 1999,
 237 **181**:5940–5947.
- 238 13. de Felipe FL, Starrenburg MJC, Hugenholtz J: **The role of NADH-oxidation in acetoin and diacetyl**
 239 **production from glucose in Lactococcus lactis subsp. lactis MG1363.** *FEMS Microbiol Lett* 1997, **156**:15–
 240 19.
- 241 14. Bolotin A, Wincker P, Mauger S, Jaillon O, Malarme K, Weissenbach J, Ehrlich SD, Sorokin A: **The**
 242 **complete genome sequence of the lactic acid bacterium Lactococcus lactis ssp. lactis IL1403.** *Genome*
 243 *Res* 2001, **11**:731–53.
- 244 15. Duwat P, Sourice S, Cesselin B, Lamberet G, Vido K, Gaudu P, Loir Y Le, Violet F, Loubière P, Gruss A:
 245 **Respiration Capacity of the Fermenting Bacterium Lactococcus lactis and Its Positive Effects on Growth**
 246 **and Survival.** *J Bacteriol* 2001, **183**:4509–4516.
- 247 16. Brooijmans R, Smit B, Santos F, van Riel J, de Vos WM, Hugenholtz J: **Heme and menaquinone induced**
 248 **electron transport in lactic acid bacteria.** *Microb Cell Fact* 2009, **8**:28.
- 249 17. Rezaiki L, Cesselin B, Yamamoto Y, Vido K, Van West E, Gaudu P, Gruss A: **Respiration metabolism**
 250 **reduces oxidative and acid stress to improve long-term survival of Lactococcus lactis.** *Mol Microbiol*
 251 2004, **53**:1331–1342.
- 252 18. Efler P, Kilstrup M, Johnsen S, Svensson B, Hägglund P: **Two Lactococcus lactis thioredoxin paralogues**
 253 **play different roles in responses to arsenate and oxidative stress.** *Microbiology* 2015, **161**:528–538.

- 254 19. Chen J, Shen J, Solem C, Jensen PR: **Oxidative stress at high temperatures in *Lactococcus lactis* due to**
255 **an insufficient supply of Riboflavin.** *Appl Environ Microbiol* 2013, **79**:6140–7.
- 256 20. Brooijmans RJW, Poolman B, Schuurman-Wolters GK, Vos WM de, Hugenholtz J: **Generation of a**
257 **Membrane Potential by *Lactococcus lactis* through Aerobic Electron Transport.** *J Bacteriol* 2007,
258 **189**:5203–5209.
- 259 21. Makarova K, Slesarev A, Wolf Y, Sorokin A, Mirkin B, Koonin E, Pavlov A, Pavlova N, Karamychev V,
260 Polouchine N, et al.: **Comparative Genomics of the Lactic Acid Bacteria.** *Proc Natl Acad Sci* 2006,
261 **103**:15611–15616.
- 262 22. Pedersen MB, Gaudu P, Lechardeur D, Petit M-A, Gruss A: **Aerobic Respiration Metabolism in Lactic**
263 **Acid Bacteria and Uses in Biotechnology.** *Annu Rev Food Sci Technol* 2012, **3**:37–58.
- 264 23. Hugenholtz J: **Citrate metabolism in lactic acid bacteria.** *FEMS Microbiol Rev* 1993, **12**:165–178.
- 265 24. Ortiz ME, Bleckwedel J, Raya RR, Mozzi F: **Biotechnological and in situ food production of polyols by**
266 **lactic acid bacteria.** *Appl Microbiol Biotechnol* 2013, **97**:4713–4726.
- 267 25. Wisselink H., Weusthuis R., Eggink G, Hugenholtz J, Grobben G.: **Mannitol production by lactic acid**
268 **bacteria: a review.** *Int Dairy J* 2002, **12**:151–161.
- 269 26. Veiga-da-Cunha M, Santos H, Van Schaftingen E: **Pathway and regulation of erythritol formation in**
270 ***Leuconostoc oenos*.** *J Bacteriol* 1993, **175**:3941–8.
- 271 27. Stolz P, Vogel RF, Hammes WP: **Utilization of electron acceptors by lactobacilli isolated from**
272 **sourdough - II. *Lactobacillus pontis*, *L. reuteri*, *L. amylovorus*, and *L. fermentum*.** *Z Lebensm Unters*
273 *Forsch* 1995, **201**:402–410.
- 274 28. Filannino P, Gobetti M, De Angelis M, Di Cagno R: **Hydroxycinnamic acids used as external acceptors**
275 **of electrons: An energetic advantage for strictly heterofermentative lactic acid bacteria.** *Appl Environ*
276 *Microbiol* 2014, **80**:7574–7582.
- 277 29. Rodríguez H, Curiel JA, Landete JM, de las Rivas B, de Felipe FL, Gómez-Cordovés C, Mancheño JM,
278 Muñoz R: **Food phenolics and lactic acid bacteria.** *Int J Food Microbiol* 2009, **132**:79–90.
- 279 30. Sauer M, Russmayer H, Grabherr R, Peterbauer CK, Marx H: **The Efficient Clade: Lactic Acid Bacteria for**
280 **Industrial Chemical Production.** *Trends Biotechnol* 2017, **35**:756–769.
- 281 31. Uden G, Bongaerts J: **Alternative respiratory pathways of *Escherichia coli*: Energetics and**
282 **transcriptional regulation in response to electron acceptors.** *Biochim Biophys Acta - Bioenerg* 1997,
283 **1320**:217–234.
- 284 32. Tachon S, Michelon D, Chambellon E, Cantonnet M, Mezange C, Henno L, Cachon R, Yvon M:
285 **Experimental conditions affect the site of tetrazolium violet reduction in the electron transport chain**
286 **of *Lactococcus lactis*.** *Microbiology* 2009, **155**:2941–2948.
- 287 33. Michelon D, Tachon S, Ebel B, De Coninck J, Feron G, Gervais P, Yvon M, Cachon R: **Screening of lactic**
288 **acid bacteria for reducing power using a tetrazolium salt reduction method on milk agar.** *J Biosci*
289 *Bioeng* 2013, **115**:229–32.
- 290 34. LeBlanc JG, Milani C, de Giori GS, Sesma F, van Sinderen D, Ventura M: **Bacteria as vitamin suppliers to**
291 **their host: a gut microbiota perspective.** *Curr Opin Biotechnol* 2013, **24**:160–168.
- 292 35. Lechardeur D, Cesselin B, Fernandez A, Lamberet G, Garrigues C, Pedersen M, Gaudu P, Gruss A: **Using**
293 **heme as an energy boost for lactic acid bacteria.** *Curr Opin Biotechnol* 2011, **22**:143–149.
- 294 36. Vido K, Bars D le, Mistou M-Y, Anglade P, Gruss A, Gaudu P: **Proteome Analyses of Heme-Dependent**
295 **Respiration in *Lactococcus lactis*: Involvement of the Proteolytic System.** *J Bacteriol* 2004, **186**:1648–
296 1657.
- 297 37. Pedersen MB, Garrigues C, Tuphile K, Brun C, Vido K, Bennedsen M, Møllgaard H, Gaudu P, Gruss A:
298 **Impact of Aeration and Heme-Activated Respiration on *Lactococcus lactis* Gene Expression:**
299 **Identification of a Heme-Responsive Operon.** *J Bacteriol* 2008, **190**:4903–4911.

- 300 38. Guidone A, Ianniello RG, Ricciardi A, Zotta T, Parente E: **Aerobic metabolism and oxidative stress**
 301 **tolerance in the Lactobacillus plantarum group.** *World J Microbiol Biotechnol* 2013, **29**:1713–1722.
- 302 39. Morishita T: **Production of menaquinones by lactic acid bacteria.** *J Dairy Sci* 1999, **82**:1897–1903.
- 303 40. Brasca M, Morandi S, Lodi R, Tamburini A: **Redox potential to discriminate among species of lactic acid**
 304 **bacteria.** *J Appl Microbiol* 2007, **103**:1516–1524.
- 305 41. Larsen N, Moslehi-Jenabian S, Werner BB, Jensen ML, Garrigues C, Vogensen FK, Jespersen L:
 306 **Transcriptome analysis of Lactococcus lactis subsp. lactis during milk acidification as affected by**
 307 **dissolved oxygen and the redox potential.** *Int J Food Microbiol* 2016, **226**:5–12.
- 308 42. Sevier CS, Kaiser CA: **Formation and transfer of disulphide bonds in living cells.** *Nat Rev Mol Cell Biol*
 309 2002, **3**:836–847.
- 310 43. Li Y, Hugenholtz J, Abee T, Molenaar D: **Glutathione protects Lactococcus lactis against oxidative**
 311 **stress.** *Appl Environ Microbiol* 2003, **69**:5739–45.
- 312 44. Michelon D, Abraham S, Ebel B, De Coninck J, Husson F, Feron G, Gervais P, Cachon R: **Contribution of**
 313 **exofacial thiol groups in the reducing activity of Lactococcus lactis.** *FEBS J* 2010, **277**:2282–2290.
- 314 45. Jänsch A, Korakli M, Vogel RF, Gänzle MG: **Glutathione reductase from Lactobacillus sanfranciscensis**
 315 **DSM20451 T: Contribution to oxygen tolerance and thiol exchange reactions in wheat sourdoughs.**
 316 *Appl Environ Microbiol* 2007, **73**:4469–4476.
- 317 46. Roussel C: **Compréhension des mécanismes physiologiques et génétiques impliqués dans l'activité**
 318 **réductrice de Lactococcus lactis.** These PhD, Université de Bourgogne, Dijon, France; 2015.
- 319 47. Capuani A, Werner S, Behr J, Vogel RF: **Effect of controlled extracellular oxidation-reduction potential**
 320 **on microbial metabolism and proteolysis in buckwheat sourdough.** *Eur Food Res Technol* 2014,
 321 **238**:425–434.
- 322 48. Kieronczyk A, Cachon R, Feron G, Yvon M: **Addition of oxidizing or reducing agents to the reaction**
 323 **medium influences amino acid conversion to aroma compounds by Lactococcus lactis.** *J Appl Microbiol*
 324 2006, **101**:1114–22.
- 325 49. Abraham S, Cachon R, Colas B, Feron G, De Coninck J: **Eh and pH gradients in Camembert cheese during**
 326 **ripening: Measurements using microelectrodes and correlations with texture.** *Int Dairy J* 2007,
 327 **17**:954–960.
- 328 50. Tyler CA, Kopit L, Doyle C, Yu AO, Hugenholtz J, Marco ML: **Polyol production during**
 329 **heterofermentative growth of the plant isolate Lactobacillus florum 2F.** *J Appl Microbiol* 2016,
 330 **120**:1336–1345.
- 331 51. Drider D, Bekal S, Prevost H: **Genetic organization and expression of citrate permease in lactic acid**
 332 **bacteria.** *Genet Mol Res* 2004, **3**:273–281.
- 333 52. Zhang C, Gänzle MG: **Metabolic pathway of α -ketoglutarate in Lactobacillus sanfranciscensis and**
 334 **Lactobacillus reuteri during sourdough fermentation.** *J Appl Microbiol* 2010, **109**:1301–1310.

335
 336

337 Annotation to references

- 338 **7.** This review by Seamus Condon from 1987 gives an excellent overview of the effect of oxygen on the
 339 metabolism of lactic acid bacteria. Although the ability to respire was not known at the time, the
 340 review describes the beneficial effect of oxygen on some LAB, as well as the oxidative stress.
- 341 **11.** This paper by Tachon et al. from 2010 describes a thorough genetic and physiological analysis of
 342 enzymes and cofactors responsible for oxygen removal and lowering of the redox potential of
 343 *Lactococcus lactis* in milk. This paper will become a key paper in the field of LAB metabolism.

- 344 **15.** Duwat et al. demonstrates in this paper from 2001 with excellent clarity that *Lactococcus lactis* is able
345 to respire. This paper opens a new scientific field on aerobic respiration in LAB.
- 346 **31.** With simple means Tachon et al. showed that redox reactions take place at both sides of the bacterial
347 membrane.
- 348 **40.** Brasca, Morandi, and Tamburini describes the typical evolution of the redox potential during
349 fermentation in milk by 88 strains from 10 different species. It is surprising that such reference data set
350 is established as late as 2007. It is also remarkable how different commonly used acidifying LAB behave
351 regarding the final redox potential reached.
- 352 **44.** Michelon et al. 2010 demonstrates clearly that the low redox potential reached by *Lactococcus lactis* is
353 due to exofacial thiol groups. However, the proteins carrying the thiol groups and the biological
354 function of the thiol groups are not identified.
355
356
357
358
359
360
361
362
363
364
365
366
367
368
369
370
371
372
373
374
375
376
377
378
379
380
381
382
383
384
385

386
387

Figure legends

Figure 1

388 Reactions contributing to maintaining redox homeostasis in lactic acid bacteria during fermentative growth
389 [11,32,44,46].

Lactate dehydrogenase (LDH) is the primary electron acceptor for homo-fermentative lactic acid bacteria and a major electron acceptor for all lactic acid bacteria. Alternative electron acceptors (A) can be reduced intracellularly or extracellularly. The electrons are directed towards the acceptors through various dehydrogenases (DH) possibly via menaquinones or disulphides.

A: electron acceptor; RA: reduced form of A (examples of As and corresponding RAs are given in Table 1); DH: dehydrogenase; GlpD: glycerol-3-phosphate dehydrogenase; G3P: glycerol-3-phosphate; DHAP: dihydroxyacetonephosphate; MK: menaquinone; MKH₂: menaquinol; NAD: nicotinamide adenine dinucleotide; NADH: reduced form of nicotinamide adenine dinucleotide; NoxAB: NADH dehydrogenase AB; NoxE: NADH oxidase E; CytBC: cytochrome bc; LDH: lactate dehydrogenase; TR: thioredoxin reductase; TS₂: thioredoxin oxidized form; T(SH)₂: thioredoxin reduced form; GR: glutathione reductase; GSH: glutathione; (GS)₂: oxidized glutathione; ESP: exo facial thiol containing protein.

390

391

392

393

394

395

396

Highlights

397

- pathways of redox reactions distinguish cultures for food fermentations

398

- range of electron acceptors in food matrices differ between food and types of cultures

399

- redox engineering of cultures for food fermentations is underexploited

400

-

401

