

1 Utilization of nitrate abolishes the “Custers effect” in *Dekkera bruxellensis* and
2 determines a different pattern of fermentation products.

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17

18 **Abstract**

19 Nitrate is one of the most abundant nitrogen sources in nature. Several yeast species have been
20 shown able to assimilate nitrate and nitrite, but the metabolic pathway has been studied in very few
21 of them. *D. bruxellensis* can use nitrate as sole nitrogen source and this metabolic characteristic can
22 render *D. bruxellensis* able to overcome *S. cerevisiae* populations in industrial bioethanol
23 fermentations. In order to better characterise how nitrate utilization affects carbon metabolism and
24 the yields of the fermentation products, we investigated this trait in defined media under well
25 controlled aerobic and anaerobic conditions. Our experiments showed that in *D. bruxellensis*
26 utilization of nitrate determines a different pattern of fermentation products. Acetic acid, instead of
27 ethanol, became in fact the main product of glucose metabolism under aerobic conditions. We have
28 also demonstrated that under anaerobic conditions nitrate assimilation abolishes the “Custers
29 effect”, in this way improving its fermentative metabolism. This can offer a new strategy, besides
30 aeration, to sustain growth and ethanol production for the employment of this yeast in industrial
31 processes.

32

33 **Keywords:** *Dekkera bruxellensis*, nitrate metabolism, Custers effect, ethanol production

34

35 **Introduction**

36 Nitrate is one of the most abundant nitrogen sources in nature. In the biosphere, nitrate assimilation
37 is the major pathway converting inorganic nitrogen to organic forms. It has been estimated that
38 more than 2×10^4 megatons of organic nitrogen per year are produced by nitrate assimilation in a
39 variety of organisms, including bacteria, fungi, algae and plants [15]. Several yeast species have
40 been shown able to assimilate nitrate and nitrite, but the metabolic pathway has been studied in
41 very few of them so far, especially in that ones which have been receiving biotechnological interest
42 such as *Hansenula polymorpha*, *Pichia anomala*, *Pachysolen tannophilus* and *Arxula*
43 *adenivorans* [16, 2, 14, 7]. The nitrate assimilation pathway in yeast is the same as the one
44 described for plants and filamentous fungi [29]. Two kinds of nitrate transporters have been
45 characterized, with high and low affinity [19]. After its uptake, nitrate is converted to ammonium by
46 two successive reductions catalyzed by nitrate reductase and nitrite reductase respectively (Fig. 1).
47 In *H. polymorpha* the genes for nitrate transporter, nitrate reductase and nitrite reductase are
48 clustered [3]. These genes are induced by nitrate and nitrite and repressed by ammonium as well as
49 by other factors involved in the utilization of secondary nitrogen sources [27, 29]. This is in
50 agreement with the preference exhibited by yeast to use inorganic compounds as nitrogen sources,
51 like ammonium, as well as various amino acids.

52 *Dekkera bruxellensis* is often associated with wine production and lambic beer fermentation and it
53 may contribute in a positive or negative way to the flavor development [8, 12, 21]. *D. bruxellensis*
54 has also been reported to contaminate distilleries producing fuel ethanol, specially in continuous
55 fermentation systems where it has been observed that this yeast can outcompete *S. cerevisiae* [18,
56 23]. Although *D. bruxellensis* and *S. cerevisiae* are considered as two phylogenetically very distant
57 relatives, they share several peculiar traits, such as the ability to produce ethanol under aerobic
58 conditions, high tolerance towards ethanol and acid, and the ability to grow without oxygen [28, 13,
59 17]. Apparently, these traits have evolved in parallel in both groups, but the molecular mechanisms
60 involved may be different [28]. *D. bruxellensis* can use nitrate as sole nitrogen source [10], whereas

61 *S. cerevisiae* cannot. Woolfit et al. [32] reported the presence of five genes encoding for nitrate
62 assimilation pathway in *D. bruxellensis*. Recently it has been shown that this metabolic
63 characteristic can render *D. bruxellensis* able to overcome *S. cerevisiae* populations in industrial
64 fermentations [11]. The presence of these skills together with a wide range of carbon sources
65 utilisation by this yeast species has been leading to the idea that *D. bruxellensis* could become a
66 new industrially relevant ethanol producing organism [4, 13, 6, 24]. In order to better characterise
67 how nitrate utilization affects carbon metabolism and the yields of fermentation products, we
68 investigated this trait under well controlled aerobic and anaerobic conditions and in well defined
69 media. Our experiments showed that utilization of nitrate determines in *D. bruxellensis* a different
70 pattern of fermentation products, in comparison to the one obtained by ammonium utilization. We
71 have also demonstrated that nitrate assimilation abolishes the “Custers effect” under anaerobic
72 conditions, improving its growth and fermentative metabolism.

73

74

75 **Materials and methods**

76

77 Yeast strains

78 The yeast strain used in this work is *Dekkera bruxellensis* CBS 2499. Stocks of the strain were
79 stored at -80 °C in 15% v/v glycerol and revitalized prior to each experiment in liquid mineral
80 medium (20 g l⁻¹ glucose; 1.7 g l⁻¹ YNB w/o amino acid and ammonium sulfate; 5 g l⁻¹ ammonium
81 sulfate).

82

83 Media and growth conditions

84 Aerobic batch cultivations were performed in shake-flasks and in a Biostat-Q system bioreactor (B-
85 Braun) with a working volume of 0.8 l. The temperature was set at 30°C, the stirring speed at 500
86 rpm and the pH, measured by Mettler Toledo pH electrode, was adjusted at 5.0 by automatic

87 addition of 2M KOH. The dissolved oxygen concentration (more than 30% of air saturation) was
88 measured by Mettler Toledo polarographic oxygen probe. The medium used was a defined synthetic
89 medium as reported by Merico et al. [20] with the only exceptions that nitrogen sources were:
90 ammonium sulfate, 5.0 g l⁻¹; sodium nitrate, 6.43 g l⁻¹; mixtures of ammonium sulfate, 5.0 g l⁻¹ and
91 sodium nitrate, 1 g l⁻¹, as specified. The media for anaerobic cultures were supplemented with
92 uracil, 50 mg l⁻¹; ergosterol, 10 mg l⁻¹; and Tween 80, 420 mg l⁻¹.

93 The anaerobic batch cultivations were performed in Biostat-Q system. The bioreactor was flushed
94 with nitrogen (<3 p.p.m. O₂) with a flow of 0.1 - 0.3 l l⁻¹ min⁻¹. The stirring was kept constant at 500
95 r.p.m. Norprene tubes (Cole-Palmer, General Control, Milan, Italy) were used to minimize the
96 diffusion of oxygen into the bioreactor. All the cultivations were repeated at least two times.

97

98 Anaerobic plate test

99 The plates for anaerobic test were performed on mineral medium (glucose, 20 g l⁻¹; Yeast Nitrogen
100 Base without amino acids and ammonium sulfate, 1.7 g l⁻¹; agarose, 20 g l⁻¹) supplemented with
101 uracil (50 mg l⁻¹), ergosterol (10 mg l⁻¹) and Tween 80 (420 mg l⁻¹) and nitrogen sources
102 (ammonium sulfate, 5.0 g l⁻¹; sodium nitrate 6.43 g l⁻¹; mixtures of ammonium sulfate, 5.0 g l⁻¹ and
103 sodium nitrate, 1 g l⁻¹). Cells grown on liquid YPD until the exponential phase were harvested and
104 suspended in distilled water. Approximately 500 cells were spotted on the respective plates and
105 were grown anaerobically for two weeks. The anaerobic environment was established using
106 Anaerocult A system (Merck, cat. no. 1138290001 and 116387) and the strength of the anaerobiosis
107 was checked with Anaerotest strips (Merck, cat. no.115112), ensuring an oxygen content below 1
108 p.p.m. O₂. The assembly was prepared and the growth was followed according to Merck
109 instructions. Each plate included the positive and negative controls, *S. cerevisiae* and *K. lactis*,
110 respectively.

111

112 Biomass and metabolites quantification

113 Samples were withdrawn from the bioreactor at appropriate intervals and used to monitor the cell
114 growth measuring the optical density at 600 nm with a spectrophotometer, after appropriate
115 dilution. For dry weight determination, washed culture samples were filtered on a 0.45 μm glass
116 microfiber GF/A filter (Whatman) and dried 24 h at 80°C. The concentration of extracellular
117 metabolites, such as glucose, ethanol, acetate, nitrate and ammonium in the supernatants were
118 determined by commercial enzymatic kits (Roche, cat. numb. 1 0716251 035, 1 0176290 035, 1
119 0148261 035, 1 09005658 035 and 1 1112732 035 respectively). All the assays were performed in
120 triplicates and the standard deviations varied between 1 and 5%.

121 Specific consumption rates of glucose, nitrate and ammonium and specific production rates of
122 ethanol and acetic acid were calculated during the exponential phase of growth. The yields of
123 biomass, ethanol and acetic acid were calculated as the total amount of products divided by the total
124 amount of glucose utilized.

125

126 Enzyme activity assays

127 Cell extracts were prepared by extraction with acid-washed glass beads (Sigma) according to
128 Postma et al., [26], and the total amount of extracted proteins was quantified using the Bio-Rad kit
129 no. 500-002 (Bio-Rad). Acetaldehyde dehydrogenase (ACDH) and glucose 6-phosphate
130 dehydrogenase (G6PDH) were assayed according to Postma et al., [26] with the only exception that
131 the concentration of NADH was increased to 4 mM for the assay of NADH-dependent activity of
132 acetaldehyde dehydrogenase. Nitrate reductase assay was performed in 50 mM potassium
133 phosphate buffer pH 7 and 0.2 mM NADH or NADPH. The reaction was started by the addition of
134 10 mM sodium nitrate and the formation of NAD^+ or NADP^+ was followed at 340 nm. Nitrite
135 reductase assay was performed in 50 mM potassium phosphate buffer pH 7, 10 mM MgSO_4 and
136 0.2 mM NADH or NADPH. The reaction was started by the addition of 1 mM sodium nitrite and
137 the formation of NAD^+ or NADP^+ was followed at 340 nm. A unit (U) of enzyme activity is defined

138 as 1 μmol of substrate transformed per minute using an extinction coefficient for NAD(P)H of 6.22
139 $1 \text{ mmol}^{-1} \text{ cm}^{-1}$.

140

141 **Results**

142

143 Nitrate utilization under aerobic conditions

144 In order to obtain a detailed quantitative and qualitative analysis about the effects of utilization of
145 nitrate on the metabolism of glucose and fermentation products in *D. bruxellensis*, batch cultures
146 were performed in bioreactor under strictly controlled aerobic conditions, controlled pH and on
147 synthetic media. The *D. bruxellensis* CBS 2499 strain was chosen because its genome has been
148 sequenced [32, 25], and its glucose metabolism has been characterised under aerobic as well as
149 under anaerobic conditions [13, 28]. In the first series of batch cultures *D. bruxellensis* was
150 cultivated on media containing sodium nitrate as the sole nitrogen source. Under these conditions
151 the growth rate was similar to the ammonium-based one. Interestingly, the main final product of
152 glucose fermentation was acetic acid instead of ethanol, that reached a 3.5 times higher yield and
153 was produced at a 2.5 times higher specific production rate than on ammonium-based ones (Table
154 1). On the other hand, the specific glucose consumption rate as well as the ethanol production rate
155 were both lower on nitrate-based media than on ammonium-based ones (Table 1). The utilization of
156 nitrate determined a slight increase of the biomass yield (Table 1). An analogous redirection of
157 glucose catabolic products was observed also when *D. bruxellensis* was cultivated on media
158 containing a mixture of ammonium and nitrate as nitrogen sources. In these conditions the growth
159 rate was lower than the one observed on media containing ammonium as the sole nitrogen source
160 (Table 1). Nitrate and ammonium were co-assimilated (Fig. 2). Also in this case the consumption of
161 nitrate resulted in a drastic increase in the production of acetic acid, to about a three times higher

162 yield, and in a parallel decrease of ethanol production, to 1/3 of the yield calculated on ammonium
163 sulfate-based media, respectively (Table 1).

164

165 Nitrate utilization under anaerobic conditions

166 *D. bruxellensis* grows under strict anaerobic condition on synthetic media at a very low rate [28].
167 This has been ascribed to a redox imbalance due to its scarce ability to produce glycerol, which
168 plays an important role under anaerobic condition for the reoxidation of NADH produced during the
169 amino acids synthesis [30]. The addition of amino acids to the medium has been shown in fact to
170 help its growth, partially alleviating this problem [6]. Due to the dependence by the nitrate-
171 assimilating enzymes for NAD(P)H (Fig. 1), the utilization of nitrate as nitrogen source could work
172 in the cell as a redox sink. Nitrate metabolism has been shown to greatly facilitate growth on xylose
173 under anaerobic conditions in the fungus *Fusarium oxysporum* [22]. To test this hypothesis, *D.*
174 *bruxellensis* was cultivated under strictly controlled anaerobic conditions in bioreactor on synthetic
175 media containing a mixture of ammonium and nitrate, due to the fact that we observed no growth on
176 plates containing nitrate as the sole nitrogen source (not shown). Under these conditions cells grew
177 at a higher rate than the one observed on ammonium-based medium enriched with amino acids
178 (Table 2). The growth rate was in fact more similar to the one observed under aerobic conditions
179 (Table 1 and 2 for comparison). In contrast to what occurred on ammonium-based media, where no
180 acetic acid production was detected, acetic acid was produced under anaerobic cultivation when
181 nitrate was utilized. Noteworthy, the specific acetic acid production rate corresponded to the
182 specific nitrate consumption rate, indicating that there is a strict correlation between nitrate
183 utilization and acetate production (Table 2). Nevertheless, in this case ethanol was the main product
184 of glucose metabolism and its specific production rate was the highest obtained (Table 1 and 2, for
185 comparison). Also the specific glucose consumption rate was the highest estimated, indicating that
186 the redox unbalance negatively affected glucose metabolism and its fermentative efficiency on

187 ammonium-based media. All these data indicate that the assimilation of nitrate greatly improve the
188 ability to grow under anaerobic conditions, playing like a “valve” to balance the redox potential.

189

190 Analysis of involved enzymatic activities

191 The activity of the enzymes involved in nitrate assimilation was assayed in *D. bruxellensis* cells
192 growing on nitrate-based media under aerobic as well as under anaerobic conditions. Nitrate
193 reductase was found to use either NADPH and NADH *in vitro* as the electron donor (Table 3). The
194 activities were higher in cell extracts from anaerobic growth conditions. The activity of nitrite
195 reductase was undetectable in cell extracts from aerobic cultures, but an extremely low activity was
196 assayed in cells grown under anaerobic conditions, again using either NADPH and NADH as the
197 electron donor (data not shown). In order to understand if nitrate utilization can affect the enzyme
198 activities leading to the increased acetic acid formation, we assayed acetaldehyde dehydrogenase
199 (ACDH). Under aerobic conditions the growth on nitrate-based media resulted in a decreased
200 activity of ACDH (Table 3). Moreover, we observed that its affinity for NADP was higher than for
201 NAD (see Materials and Methods). Interestingly, we found that nitrate utilization under anaerobic
202 conditions determined an increased specific activity of NADP-dependent ACDH, which was in fact
203 higher in nitrate-grown cell extracts than in ammonium-grown ones (Table 3) and, in parallel, a
204 decrease in the NAD-dependent ACDH activity. On the other hand, the activity of glucose 6-
205 phosphate dehydrogenase (G6PDH), which is one of the main sources of NADPH, was lower in
206 nitrate-grown cells (Table 3).

207

208

209 **Discussion**

210

211 The use of nitrate as nitrogen source determines in *D. bruxellensis* deep changes in the distribution
212 of the final fermentation products. This is well evident under aerobic and under strictly anaerobic

213 conditions. Under aerobic conditions acetic acid resulted in fact as the main product of glucose
214 metabolism, at the expense of the ethanol production (Table 1). In *D. bruxellensis* nitrate and nitrite
215 reductases can use, *in vitro*, NADH as well as NADPH as electron donors (Table 3), like most yeast
216 nitrate reductases studied so far [29]. Nitrate assimilatory enzymes and alcohol dehydrogenase
217 (ADH) can then compete for NADH, leading to a reduced ethanol synthesis. As a consequence,
218 acetaldehyde can accumulate and triggers acetic acid formation. Apparently, this is what occurs
219 under aerobic conditions. On the other hand, the stoichiometry of nitrate utilization under anaerobic
220 conditions rather suggested that nitrate reductase could mainly require, *in vivo*, NADPH as the
221 electron donor (Fig. 1). In fact the amount of acetic acid produced in this condition corresponded
222 exactly to the amount of NADPH required in the first nitrate-assimilating step, converting nitrate to
223 nitrite through a NADPH-dependent nitrate reductase (10 mmol of nitrate assimilated and 10 mmol
224 of acetic acid produced). It is noteworthy that acetic acid production has never been found under
225 strict anaerobic conditions in *D. bruxellensis* [28], being its production associated to the oxygen
226 concentration [9]. Furthermore, acetic acid specific production and nitrate specific consumption
227 showed the same rate, corroborating the strict correlation between nitrate utilization and acetate
228 production. This link can indicate that a specific need of NADPH can be satisfied by a NADP-
229 dependent ACDH activity. In agreement with this hypothesis, an increased level of NADP-
230 dependent ACDH activity was found in nitrate-grown cells in comparison to the ammonium-grown
231 cells (Table 3). Since this cofactor is required for cellular biosynthesis, in the case of nitrate
232 assimilation it could become limiting in cell metabolism, and this can in turn stimulate metabolic
233 pathways able to generate it, as acetic acid formation.

234 Another important aspect of the redox balance is linked to NAD/NADH ratio, which is an
235 especially critic step under anaerobic conditions. NADH is in fact generated not only by glycolytic
236 pathway, but also by amino acids synthesis. The most important reaction for reoxidation of this
237 surplus NADH under anaerobic conditions is the production of glycerol (Fig. 1) [30]. In *D.*
238 *bruxellensis* a very low amount of this compound was produced under those conditions [28] and the

239 inefficiency of this pathway has been indicated as the main cause of the “Custers effect” in this
240 species [31]. Nitrate assimilation could then accomplish the role of balancing the redox status. The
241 higher growth rate and higher specific ethanol production rate obtained in this work under anaerobic
242 conditions (Table 2) indicate that nitrate utilization greatly improves the fermentative metabolism in
243 *D. bruxellensis*. In *S. cerevisiae* it has been calculated that 13 mmol of glycerol per g of dry
244 biomass are generated under anaerobic conditions, leading to the reoxidation of 13 mmol of NADH
245 [1]. In *D. bruxellensis* the assimilation of 10 mmol of nitrate to ammonium through a NADH-
246 dependent nitrite reductase could result in the reoxidation of 30 mmol of NADH (Fig. 1), which fits
247 well with the theoretical formation of 26 mmol of NADH generated from the biosynthesis of 2 g of
248 dry biomass produced during its anaerobic growth.

249 In conclusion, nitrate assimilation determines in *D. bruxellensis* an improved ability to grow under
250 anaerobic conditions and enhances its fermentative metabolism, working like a redox “valve” and,
251 in this way, abolishing the “Custers effect”. This offers a new strategy, besides the controlled
252 aeration, for the employment of this yeast in industrial processes.

253

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256

257 **References**

258

259 1. Albers E, Larsson C, Liden G, Niklasson C, Gustafsson L. (1996) Influence of the nitrogen
260 source on *Saccharomyces cerevisiae* anaerobic growth and product formation. Appl Environ
261 Microbiol 62:3187-3195

262 2. Avila J, Pérez MD, Brito N, Gonzàles C and Siverio JM (1995) Cloning and disruption of *YNR1*
263 gene encoding the nitrate reductase apoenzyme of the yeast *Hansenula polymorpha*. FEBS Lett
264 366:137-142

- 265 3. Avila J, Gonzàles C, Brito N, et al. (1998) Clustering of the *YNAI* gene encoding a Zn(II)2Cys6
266 transcriptional factor in the yeast *Hansenula polymorpha* with the nitrate assimilation genes *YNT1*,
267 *YNII* and *YNRI* and its involvement in their transcriptional activation. *Biochem J* 335:647-652
- 268 4. Blomqvist J, Eberhard T, Schnürer J, Passoth V (2010) Fermentation characteristics of *Dekkera*
269 *bruxellensis* strains. *Appl Microbiol Biotechnol* 87:1487-1497
- 270 5. Blomqvist J, South E, Tiukova I, Momeni MH, Hansson H, Ståhlberg J, Horn SJ, Schnürer J,
271 Passoth V (2011) Fermentation of lignocellulosic hydrolysate by the alternative industrial ethanol
272 yeast *Dekkera bruxellensis*. *Lett Appl Microbiol* 53: 73-78
- 273 6. Blomqvist J, Nogué, VN, Gorwa-Grauslund M, Passoth V (2012) Physiological requirements for
274 growth and competitiveness of *Dekkera bruxellensis* under oxygen-limited or anaerobic conditions.
275 *Yeast* 29:265-274
- 276 7. Böer E, Schröter A, Bode R, Piontek M, Kunze G (2009) Characterization and expression
277 analysis of a gene cluster for nitrate assimilation from the yeast *Arxula adenivorans*. *Yeast* 26:83-
278 93
- 279 8. Boulton R, Singleton V, Bisson L, Kunkee R. (eds). (1996) Principles and Practices of
280 Winemaking Chapman Hall, New York
- 281 9. Ciani M, Ferraro L. (1997) Role of oxygen on acetic acid production by *Brettanomyces/Dekkera*
282 in winemaking. *J Sci Food Agric* 75:489-495
- 283 10. Conterno L, Joseph LCM, Arvik TJ, Henick-Kling T & Bisson LF (2006) Genetic and
284 physiological characterization of *Brettanomyces bruxellensis* strains isolated from wine. *Am J Enol*
285 *Vitic* 57:139-147
- 286 11. De Barros Pita W, Leite FC, de Souza Liberal A, Simões DA, Morais MA Jr (2011) The ability
287 to use nitrate confers advantage to *Dekkera bruxellensis* over *S. cerevisiae* and can explain its
288 adaptation to industrial fermentation processes. *Antonie Leeuwenhoek* 100:99-107
- 289 12. Fugelsang KC. (eds). (1996). Wine microbiology. Chapman & Hall Publishers, New York
- 290 13. Galafassi S, Merico A, Pizza F, Hellborg L, Molinari F, Piškur J, Compagno C (2011)

291 *Dekkera/Brettanomyces* yeasts for ethanol production from renewable sources under oxygen-
292 limited and low pH conditions. J Ind Microbiol Biotechnol 38:1079-88

293 14. Garcia-Lugo P, Gonzàles C, Perdomo G, Brito N, Avila J, de la Rosa JM, Siverio JM (2000)
294 Cloning, sequencing and expression of *HaYNRI* and *HaYNII*, encoding nitrate and nitrite
295 reductases in the yeast *Hansenula anomala*. Yeast 16:1099-1105

296 15. Guerrero MG, Vega JM, Losada M (1981) The assimilatory nitrate-reducing system and its
297 regulation. Annu Rev Plant Physiol 32:169-204

298 16. Jeffries TW (1983) Effects of nitrate on fermentation of xylose and glucose by *Pachisolen*
299 *tannophilus*. BIO/Technology 1:503-506

300 17. Leite FC, Basso TO, de Barros Pita W, Gombert AK, Simões DA, de Moraes MA Jr (2012)
301 Quantitative aerobic physiology of the yeast *Dekkera bruxellensis*, a major contaminant in
302 bioethanol production plants. FEMS Yeast Res Sep 21, p S1567 doi: 10.1111/1567-1364.12007

303 18. Liberal ATS, Basilio ACM, Resende AM, Brasileiro BTRV, da Silva-Filho EA, Moraes JOF,
304 Simões DA, Moraes Jr MA (2007) Identification of *Dekkera bruxellensis* as a major contaminant
305 yeast in continuous fuel ethanol fermentation. J Appl Microbiol 102:538-547

306 19. Machin F, Perdomo G, Pèrez MD, Brito N, Siverio JM (2000) Evidence for multiple nitrate
307 uptake systems in *Hansenula polymorpha*. FEMS Microbiol Lett 194:171-174

308 20. Merico A, Sulo P, Piškur J, Compagno C (2007) Fermentative lifestyle in yeasts belonging to
309 the *Saccharomyces* complex. FEBS J 274:976-989

310 21. Oelofse A, Pretorius IS, du Toit M (2008) Significance of *Brettanomyces* and *Dekkera* during
311 winemaking: a synoptic review. S Afr J Enol Vitic 29:128-144

312 22. Panagiotou G, Christakopoulos P, Grotjaer T, Olsson L (2006) Engineering of the redox
313 imbalance of *Fusarium oxysporum* enables anaerobic growth on xylose. Metab Eng 8:474-482

314 23. Passoth V, Blomqvist J, Schnürer J (2007) *Dekkera bruxellensis* and *Lactobacillus vini* from a
315 stable ethanol-producing consortium in a commercial alcohol process. Appl Environ Microbiol
316 73:4354-4356

- 317 24. Pereira LF, Bassi AP, Avansini SH, Neto AG, Brasileiro BT, Ceccato-Antonini SR, de Moraes
318 MA Jr (2012) The physiological characteristics of the yeast *Dekkera bruxellensis* in fully
319 fermentative conditions with cell recycling and in mixed cultures with *Saccharomyces cerevisiae*.
320 *Antonie van Leeuwenhoek* 101:529-539
- 321 25. Piškur J, Ling Z, Marcet-Houben M, Ishchuk OP, Aerts A, LaButti K, Copeland A, Lindquist E,
322 Barry K, Compagno C, Bisson L, Grigoriev IV, Gabaldón T, Phister T (2012) The genome of wine
323 yeast *Dekkera bruxellensis* provides a tool to explore its food-related properties. *Int J Food*
324 *Microbiol* 157:202-209
- 325 26. Postma E, Verduyn C, Scheffers WA, van DijkenJP (1989) Enzymatic analysis of the Crabtree
326 effect in glucose-limited chemostat cultures of *Saccharomyces cerevisiae*. *Appl Env Microbiol*
327 55:468-477
- 328 27. Rossi B, Manasse S, Serrani F, Berardi E (2005) *Hansenula polymorpha* NMR2 and NMR4, two
329 new loci involved in nitrogen metabolite repression. *FEMS Yeast Res* 5:1009-1017
- 330 28. Rozpedowska E, Hellborg L, Ishchuk OP, Orhan F, Galafassi S, Merico A, Woolfit M,
331 Compagno C, Piškur J (2011) Parallel evolution of the make-accumulate-consume strategy in
332 *Saccharomyces* and *Dekkera* yeasts. *Nat Commun* 2:302
- 333 29. Siverio JM. (2002) Assimilation of nitrate by yeasts. *FEMS Microbiol Rev* 26:277-284
- 334 30. van Dijken JP, Scheffers AW (1986) Redox balances in the metabolism of sugars by yeasts.
335 *FEMS Microbiol Rev* 32:199-224
- 336 31. Wijsman MR, van Dijken JP, van Kleeff BH, Scheffers WA (1984) Inhibition of fermentation
337 and growth in batch cultures of the yeast *Brettanomyces intermedius* upon a shift from aerobic to
338 anaerobic conditions (Custers effect). *Antonie Leeuwenhoek* 50(2):183-92
- 339 32. Woolfit M, Rozpedowska E, Piškur J, Wolfe KH (2007) Genome survey sequencing of the wine
340 spoilage yeast *Dekkera (Brettanomyces) bruxellensis*. *Eukaryot Cell* 6(4):721-733
- 341

342 Legend of Tables and Figures.

343

344 **Table 1** Growth parameters during aerobic fermentations on glucose mineral medium with
345 ammonium sulphate (5 g l^{-1}), sodium nitrate (6.43 g l^{-1}) or mixture of ammonium sulphate (5 g l^{-1})
346 and sodium nitrate (1 g l^{-1}) as nitrogen sources. * *data from [28]*

347

348 **Table 2** Growth parameters during anaerobic fermentations on glucose mineral medium with
349 ammonium sulphate (5 g l^{-1}) or mixture of ammonium sulphate (5 g l^{-1}) and sodium nitrate (1 g l^{-1})
350 as nitrogen sources. * *data from [28]*

351

352 **Table 3** Activity of enzymes involved in NAD(P)/NAD(P)H utilization during growth in aerobic or
353 anaerobic conditions, on glucose mineral media with ammonium sulphate (5 g l^{-1}), sodium nitrate
354 (6.43 g l^{-1}) or mixture of ammonium sulphate (5 g l^{-1}) and sodium nitrate (1 g l^{-1}) as nitrogen
355 sources. * *data from [13]*

356

357 **Fig. 1** Schematic representation of pathways involved in glucose metabolism and nitrate utilization
358 with special attention to the steps directly involved in redox balance

359

360 **Fig. 2** Batch fermentation on glucose mineral medium with a mixture of ammonium sulphate (5 g l^{-1})
361 and sodium nitrate (1 g l^{-1}) as nitrogen sources. A: substrate consumption. B: biomass and
362 metabolites production.

363

364

365

Figure
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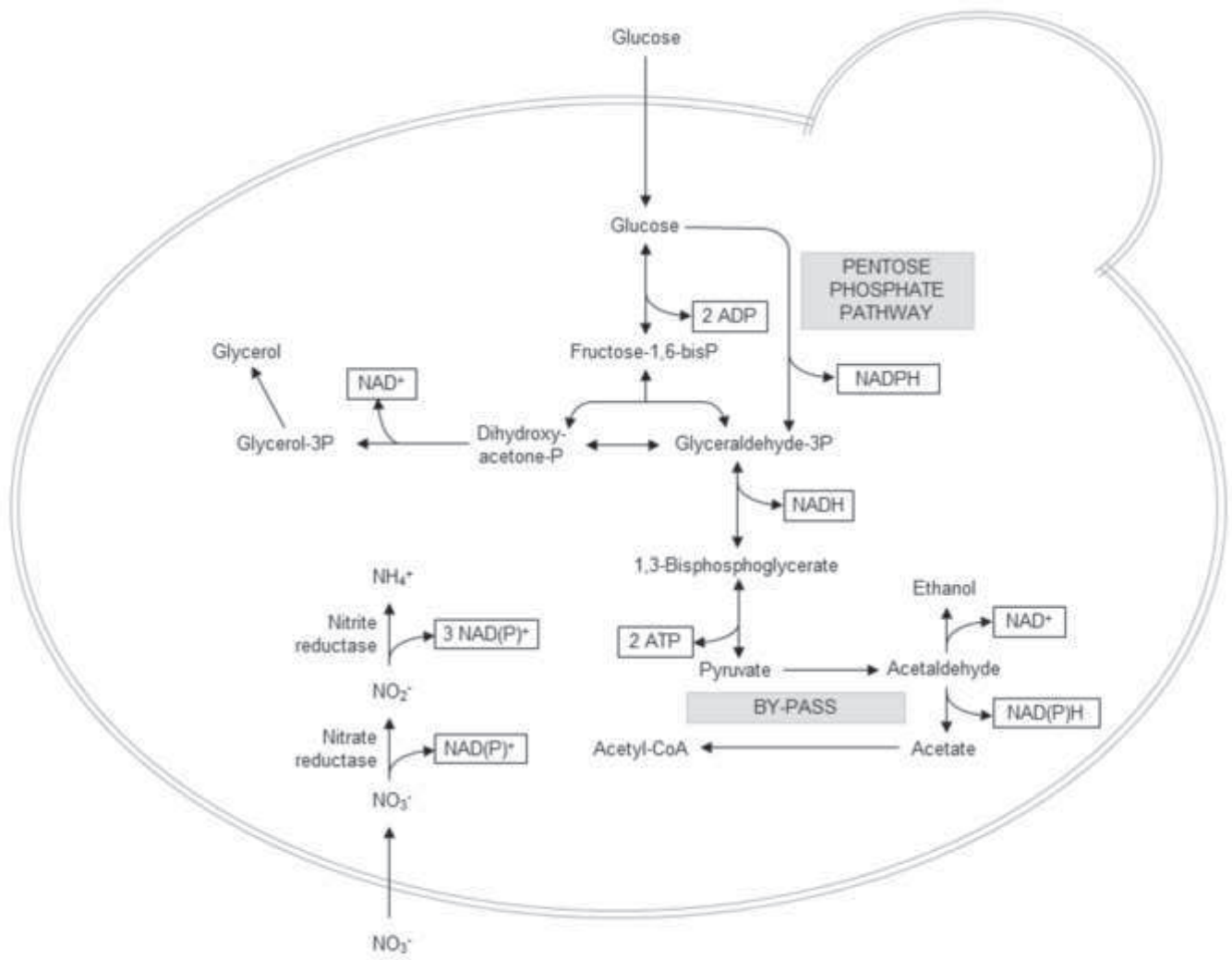
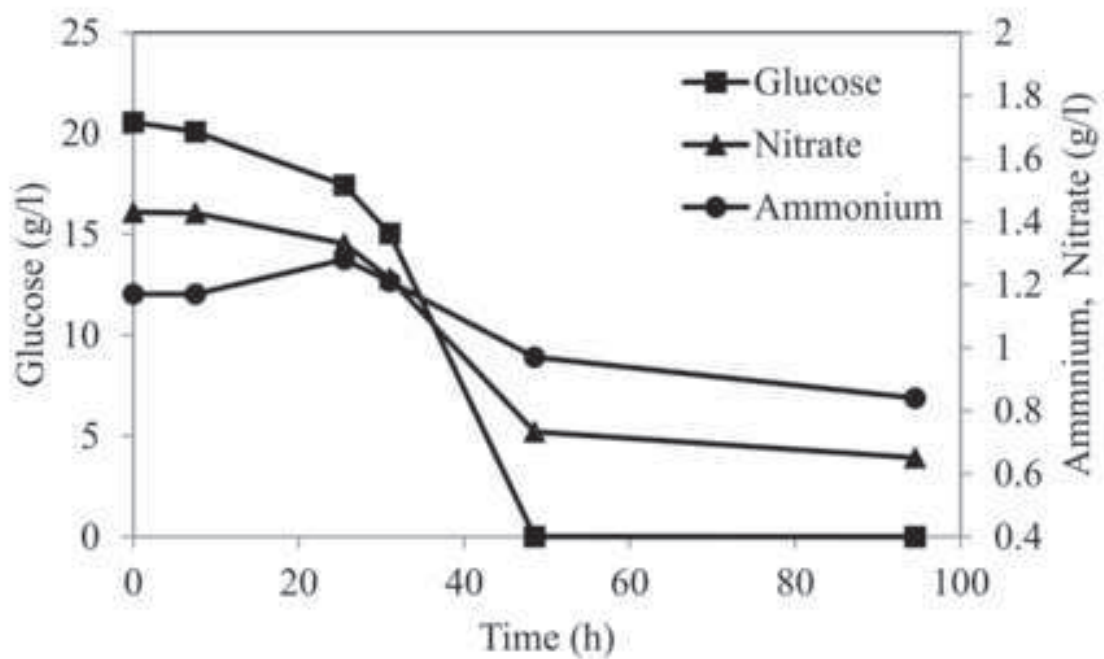


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A



B

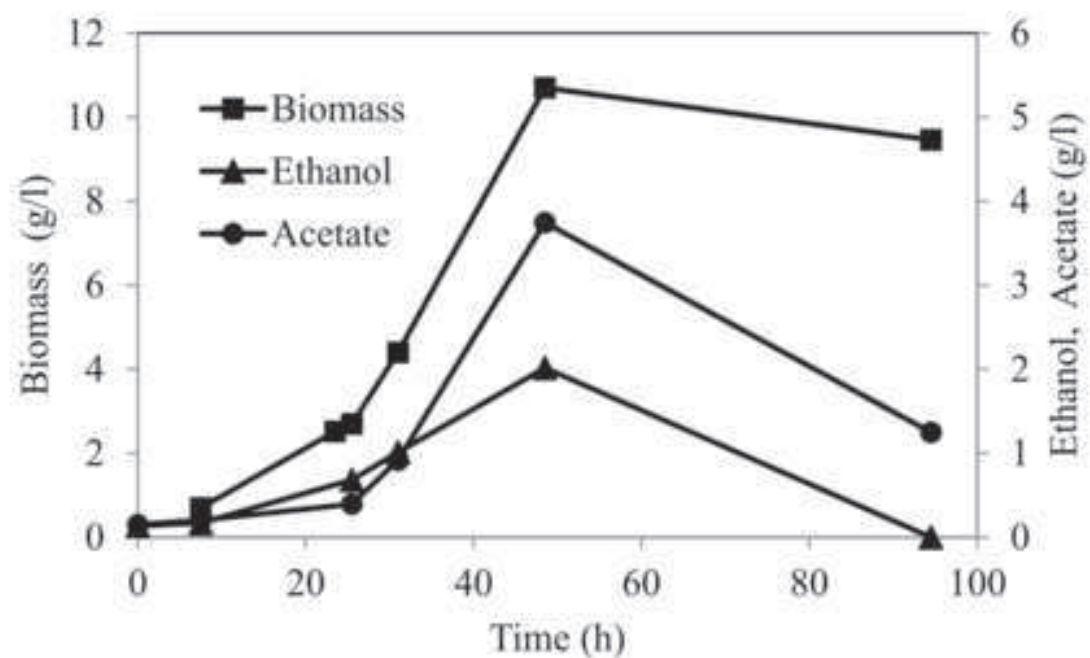


Table 1

	Ammonium*	Nitrate	Mixture
Growth rate (h^{-1})	0.11 - 0.12	0.092 ± 0.006	0.077 ± 0.004
q ($\text{mMg}_{\text{dry weight}} \text{h}^{-1}$)			
Glucose	3.6 - 3.7	2.94 ± 0.5	2.42 ± 0.28
Ethanol	3.9 - 4.4	1.65 ± 0.007	1.30 ± 0.09
Acetate	0.62 - 0.70	1.83 ± 0.009	1.59 ± 0.10
Y ($\text{g g}_{\text{glucose}}^{-1}$)			
Biomass	0.17 - 0.18	0.19 ± 0.004	0.23 ± 0.008
Ethanol	0.320 - 0.335	0.133 ± 0.006	0.138 ± 0.006
Acetate	0.058 - 0.060	0.216 ± 0.006	0.166 ± 0.011

Table 2

	Ammonium with casamino acids *	Ammonium and nitrate w/o casamino acids
Growth rate (h^{-1})	0.070 - 0.075	0.084 ± 0.006
q ($\text{mMg}_{\text{dry weight}} \text{h}^{-1}$)		
Glucose	1.47 - 1.60	4.08 ± 0.26
Ethanol	1.74 - 1.90	5.98 ± 0.42
Acetate	0.02 - 0.02	0.43 ± 0.02
Nitrate	-	0.46 ± 0.02
Y ($\text{g g}_{\text{glucose}}^{-1}$)		
Biomass	0.132 - 0.150	0.10 ± 0.005
Ethanol	0.34 - 0.35	0.35 ± 0.021
Acetate	0	0.033 ± 0.001

Table 3

Enzyme activity (U mg _{protein} ⁻¹)	Cofactor specificity	Aerobiosis		Anaerobiosis	
		Ammonium	Nitrate	Ammonium	Ammonium and nitrate
Acetaldehyde dehydrogenase	NADP ⁺ .	0.23 ±0.007	0.139 ±0.004	0.065 ±0.005	0.115 ±0.003
	NAD ⁺	0.50 ±0.03	0.24 ±0.014	0.13	0.025 ±0.006
Glucose 6 phosphate dehydrogenase	NADPH.	0.21 ±0.09 *	0.10 ±0.007	0.16 ±0.006	0.025 ±0.003
Nitrate reductase	NADH.	-	0.039 ±0.005	-	0.045 ±0.001
	NADPH	-	0.015 ±0.002	-	0.021 ±0.001