

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

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

Keywords *Dekkera bruxellensis* · Nitrate metabolism · Custers effect · Ethanol production

Introduction

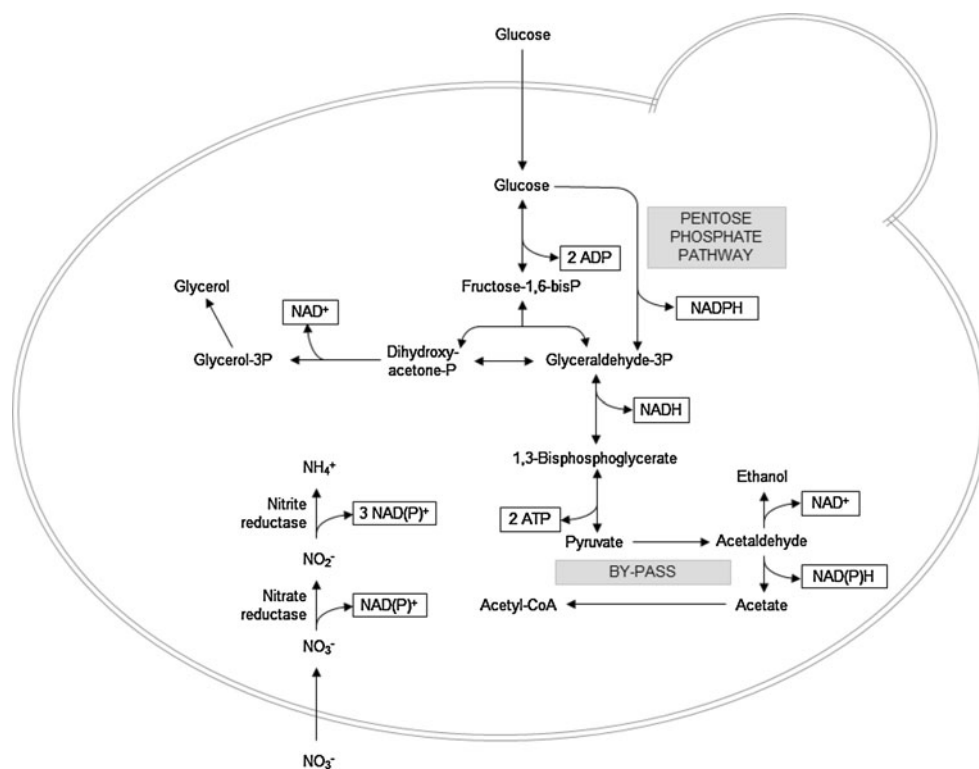
Nitrate is one of the most abundant nitrogen sources in nature. In the biosphere, nitrate assimilation is the major

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

Dekkera bruxellensis is often associated with wine production and lambic beer fermentation and it may contribute in a positive or negative way to the flavor development [8, 12, 21]. *D. bruxellensis* has also been reported to contaminate distilleries producing fuel ethanol, especially in continuous fermentation systems where it has been observed that this yeast can outcompete *S. cerevisiae* [18, 23]. Although *D. bruxellensis* and *S. cerevisiae* are considered as two phylogenetically very distant relatives, they share several peculiar traits, such as

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



the ability to produce ethanol under aerobic conditions, high tolerance towards ethanol and acid, and the ability to grow without oxygen [13, 17, 28]. Apparently, these traits have evolved in parallel in both groups, but the molecular mechanisms involved may be different [28]. *D. bruxellensis* can use nitrate as the sole nitrogen source [10], whereas *S. cerevisiae* cannot. Woolfit et al. [32] reported the presence of five genes encoding for nitrate assimilation pathway in *D. bruxellensis*. Recently, it has been shown that this metabolic characteristic can render *D. bruxellensis* able to overcome *S. cerevisiae* populations in industrial fermentations [11]. The presence of these skills together with a wide range of carbon sources utilization by this yeast species has been leading to the idea that *D. bruxellensis* could become a new industrially relevant ethanol-producing organism [4, 5, 13, 24]. In order to better characterize how nitrate utilization affects carbon metabolism and the yields of fermentation products, we investigated this trait under well-controlled aerobic and anaerobic conditions and in well-defined media. Our experiments showed that utilization of nitrate determines in *D. bruxellensis* a different pattern of fermentation products, in comparison to the one obtained by ammonium utilization. We have also demonstrated that nitrate assimilation abolishes the “Custers effect” under anaerobic conditions, improving its growth and fermentative metabolism.

Materials and methods

Yeast strains

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

Media and growth conditions

Aerobic batch cultivations were performed in shake-flasks and in a Biostat-Q system bioreactor (B-Braun) with a working volume of 0.8 l. The temperature was set at $30\text{ }^{\circ}\text{C}$, the stirring speed at 500 rpm, and the pH, measured by Mettler Toledo pH electrode, was adjusted to 5.0 by automatic addition of 2 M KOH. The dissolved oxygen concentration (more than 30 % of air saturation) was measured by Mettler Toledo polarographic oxygen probe. The medium used was a defined synthetic mineral medium as reported by Merico et al. [20] with the only exceptions that nitrogen sources were: ammonium sulfate, 5.0 g l^{-1} ; sodium nitrate, 6.43 g l^{-1} ; mixtures of ammonium sulfate, 5.0 g l^{-1} and sodium nitrate, 1 g l^{-1} , as specified. The media for anaerobic cultures were supplemented with

uracil, 50 mg l⁻¹; ergosterol, 10 mg l⁻¹; and Tween 80, 420 mg l⁻¹.

The anaerobic batch cultivations were performed in a Biostat-Q system. The bioreactor was flushed with nitrogen (<3 ppm O₂) with a flow of 0.1–0.31 l min⁻¹. The stirring was kept constant at 500 rpm. Norprene tubes (Cole-Palmer, General Control, Milan, Italy) were used to minimize the diffusion of oxygen into the bioreactor. All the cultivations were repeated at least two times.

Anaerobic plate test

The plates for anaerobic test were performed on mineral medium (glucose, 20 g l⁻¹; Yeast Nitrogen Base without amino acids and ammonium sulfate, 1.7 g l⁻¹; agarose, 20 g l⁻¹) supplemented with uracil (50 mg l⁻¹), ergosterol (10 mg l⁻¹) and Tween 80 (420 mg l⁻¹) and nitrogen sources (ammonium sulfate, 5.0 g l⁻¹; sodium nitrate 6.43 g l⁻¹; mixtures of ammonium sulfate, 5.0 g l⁻¹ and sodium nitrate, 1 g l⁻¹). Cells grown on liquid YPD until the exponential phase were harvested and suspended in distilled water. Approximately 500 cells were spotted on the respective plates and were grown anaerobically for 2 weeks. The anaerobic environment was established using Anaerocult A system (Merck, cat. no. 1138290001 and 116387) and the strength of the anaerobiosis was checked with Anaerotest strips (Merck, cat. no. 115112), ensuring an oxygen content below 1 ppm O₂. The assembly was prepared and the growth was followed according to Merck instructions. Each plate included the positive and negative controls, *S. cerevisiae* and *K. lactis*, respectively.

Biomass and metabolites quantification

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

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

Enzyme activity assays

Cell extracts were prepared by extraction with acid-washed glass beads (SIGMA) according to Postma et al. [26], and the total amount of extracted proteins was quantified using the Bio-Rad kit no. 500-002 (Bio-Rad, Hercules, CA, USA). Acetaldehyde dehydrogenase (ACDH) and glucose 6-phosphate dehydrogenase (G6PDH) were assayed according to Postma et al. [26] with the only exception that the concentration of NADH was increased to 4 mM for the assay of NADH-dependent activity of acetaldehyde dehydrogenase. Nitrate reductase assay was performed in 50 mM potassium phosphate buffer pH 7 and 0.2 mM NADH or NADPH. The reaction was started by the addition of 10 mM sodium nitrate and the formation of NAD⁺ or NADP⁺ was followed at 340 nm. Nitrite reductase assay was performed in 50 mM potassium phosphate buffer pH 7, 10 mM MgSO₄, and 0.2 mM NADH or NADPH. The reaction was started by the addition of 1 mM sodium nitrite and the formation of NAD⁺ or NADP⁺ was followed at 340 nm. A unit (U) of enzyme activity is defined as 1 μ mol of substrate transformed per minute using an extinction coefficient for NAD(P)H of 6.22 l mmol⁻¹ cm⁻¹.

Results

Nitrate utilization under aerobic conditions

In order to obtain a detailed quantitative and qualitative analysis about the effects of utilization of nitrate on the metabolism of glucose and fermentation products in *D. bruxellensis*, batch cultures were performed in a bioreactor under strictly controlled aerobic conditions, controlled pH, and on synthetic media. The *D. bruxellensis* CBS 2499 strain was chosen because its genome has been sequenced [32, 25], and its glucose metabolism has been characterized under aerobic as well as under anaerobic conditions [13, 28]. In the first series of batch cultures, *D. bruxellensis* was cultivated on media containing sodium nitrate as the sole nitrogen source. Under these conditions, the growth rate was similar to the ammonium-based one. Interestingly, the main final product of glucose fermentation was acetic acid instead of ethanol, which reached a 3.5 \times higher yield and was produced at a 2.5 \times higher specific production rate than on ammonium-based ones (Table 1). On the other hand, the specific glucose consumption rate as well as the ethanol production rate were both lower on nitrate-based media than on ammonium-based ones (Table 1). The utilization of nitrate determined a slight increase of the biomass yield (Table 1). An analogous redirection of glucose catabolic products was observed also when *D. bruxellensis* was cultivated on

media containing a mixture of ammonium and nitrate as nitrogen sources. In these conditions the growth rate was lower than the one observed on media containing ammonium as the sole nitrogen source (Table 1). Nitrate and ammonium were co-assimilated (Fig. 2). Also in this case the consumption of nitrate resulted in a drastic increase in

the production of acetic acid, to about a three times higher yield, and in a parallel decrease of ethanol production, to one-third of the yield calculated on ammonium sulphate-based media, respectively (Table 1).

Nitrate utilization under anaerobic conditions

Dekkera bruxellensis grows under strict anaerobic condition on synthetic media at a very low rate [28]. This has been ascribed to a redox imbalance due to its scarce ability to produce glycerol, which plays an important role under anaerobic condition for the reoxidation of NADH produced during the amino acids synthesis [30]. The addition of amino acids to the medium has been shown in fact to help its growth, partially alleviating this problem [6]. Due to the dependence by the nitrate-assimilating enzymes for NAD(P)H (Fig. 1), the utilization of nitrate as nitrogen source could work in the cell as a redox sink. Nitrate metabolism has been shown to greatly facilitate growth on xylose under anaerobic conditions in the fungus *Fusarium oxysporum* [22]. To test this hypothesis, *D. bruxellensis* was cultivated under strictly controlled anaerobic conditions in bioreactor on synthetic media containing a mixture of ammonium and nitrate, due to the fact that we observed no growth on plates containing nitrate as the sole nitrogen source (not shown). Under these conditions, cells grew at a higher rate than the one observed on ammonium-based medium enriched with amino acids (Table 2). The growth rate was in fact more similar to the one observed under aerobic conditions (Tables 1 and 2 for comparison). In contrast to what occurred on ammonium-based media, where no acetic acid production was detected, acetic acid was produced under anaerobic cultivation when nitrate was utilized. Noteworthy, the specific acetic acid production

Table 1 Growth parameters during aerobic fermentations on glucose mineral medium with ammonium sulphate (5 g l^{-1}), sodium nitrate (6.43 g l^{-1}), or a mixture of ammonium sulphate (5 g l^{-1}) and sodium nitrate (1 g l^{-1}) as nitrogen sources

	Ammonium ^a	Nitrate	Mixture
Growth rate (h^{-1})	0.11–0.12	0.092 ± 0.006	0.077 ± 0.004
q ($\text{mM g}_{\text{dry weight}} \text{h}^{-1}$)			
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

^a Data from [28]

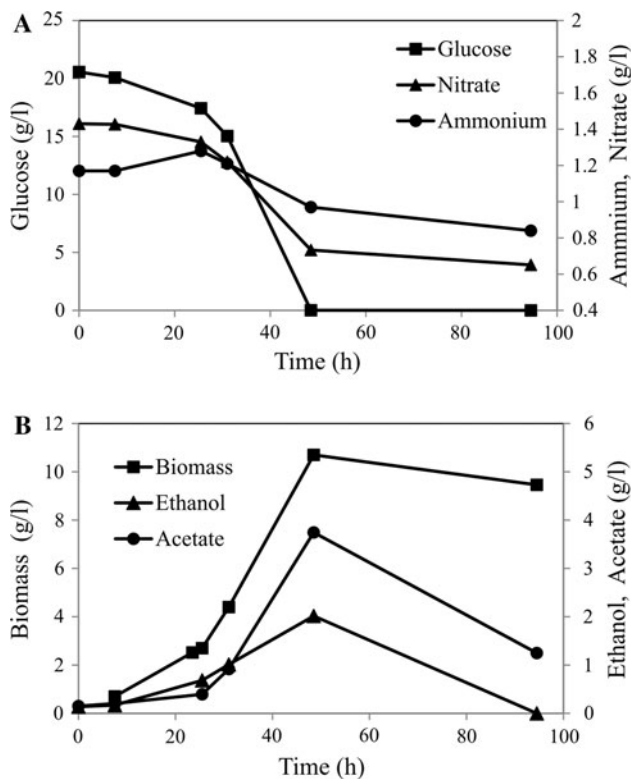


Fig. 2 Batch fermentation on glucose mineral medium with a mixture of ammonium sulphate (5 g l^{-1}) and sodium nitrate (1 g l^{-1}) as nitrogen sources. **a** Substrate consumption. **b** Biomass and metabolites production

Table 2 Growth parameters during anaerobic fermentations on glucose mineral medium with ammonium sulphate (5 g l^{-1}) or mixture of ammonium sulphate (5 g l^{-1}) and sodium nitrate (1 g l^{-1}) as nitrogen sources

	Ammonium with casamino acids ^a	Ammonium and nitrate w/o casamino acids
Growth rate (h^{-1})	0.070–0.075	0.084 ± 0.006
q ($\text{mM g}_{\text{dry weight}} \text{h}^{-1}$)		
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

^a Data from [28]

rate corresponded to the specific nitrate consumption rate, indicating that there is a strict correlation between nitrate utilization and acetate production (Table 2). Nevertheless, in this case, ethanol was the main product of glucose metabolism and its specific production rate was the highest obtained (Tables 1 and 2, for comparison). Also, the specific glucose consumption rate was the highest estimated, indicating that the redox unbalance negatively affected glucose metabolism and its fermentative efficiency on ammonium-based media. All these data indicate that the assimilation of nitrate greatly improves the ability to grow under anaerobic conditions, playing like a “valve” to balance the redox potential.

Analysis of involved enzymatic activities

The activity of the enzymes involved in nitrate assimilation was assayed in *D. bruxellensis* cells growing on nitrate-based media under aerobic as well as under anaerobic conditions. Nitrate reductase was found to use either NADPH and NADH in vitro as the electron donor (Table 3). The activities were higher in cell extracts from anaerobic growth conditions. The activity of nitrite reductase was undetectable in cell extracts from aerobic cultures, but an extremely low activity was assayed in cells grown under anaerobic conditions, again using either NADPH and NADH as the electron donor (data not shown). In order to understand if nitrate utilization can affect the enzyme activities leading to the increased acetic acid formation, we assayed acetaldehyde dehydrogenase (ACDH). Under aerobic conditions, the growth on nitrate-based media resulted in a decreased activity of ACDH (Table 3). Moreover, we observed that its affinity for NADP was higher than for NAD (see “Materials and methods”). Interestingly, we found that nitrate utilization under anaerobic conditions determined an increased specific activity of NADP-dependent ACDH, which was in fact higher in nitrate-grown cell

extracts than in ammonium-grown ones (Table 3) and, in parallel, a decrease in the NAD-dependent ACDH activity. On the other hand, the activity of glucose 6-phosphate dehydrogenase (G6PDH), which is one of the main sources of NADPH, was lower in nitrate-grown cells (Table 3).

Discussion

The use of nitrate as nitrogen source determines in *D. bruxellensis* deep changes in the distribution of the final fermentation products. This is well evident under aerobic and under strictly anaerobic conditions. Under aerobic conditions, acetic acid resulted in fact as the main product of glucose metabolism, at the expense of the ethanol production (Table 1). In *D. bruxellensis*, nitrate and nitrite reductases can use, in vitro, NADH as well as NADPH as electron donors (Table 3), like most yeast nitrate reductases studied so far [29]. Nitrate assimilatory enzymes and alcohol dehydrogenase (ADH) can then compete for NADH, leading to a reduced ethanol synthesis. As a consequence, acetaldehyde can accumulate and trigger acetic acid formation. Apparently, this is what occurs under aerobic conditions. On the other hand, the stoichiometry of nitrate utilization under anaerobic conditions rather suggested that nitrate reductase could mainly require, in vivo, NADPH as the electron donor (Fig. 1). In fact, the amount of acetic acid produced in this condition corresponded exactly to the amount of NADPH required in the first nitrate-assimilating step, converting nitrate to nitrite through a NADPH-dependent nitrate reductase (10 mmol of nitrate assimilated and 10 mmol of acetic acid produced). It is noteworthy that acetic acid production has never been found under strict anaerobic conditions in *D. bruxellensis* [28], being its production associated to the oxygen concentration [9]. Furthermore, acetic acid-specific production and nitrate-specific consumption showed the

Table 3 Activity of enzymes involved in NAD(P)/NAD(P)H utilization during growth in aerobic or anaerobic conditions, on glucose mineral media with ammonium sulphate (5 g l⁻¹), sodium nitrate (6.43 g l⁻¹), or mixture of ammonium sulphate (5 g l⁻¹) and sodium nitrate (1 g l⁻¹) as nitrogen sources

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
	NADPH	0.21 ±0.09 ^a	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

^a Data from [13]

same rate, corroborating the strict correlation between nitrate utilization and acetate production. This link can indicate that a specific need of NADPH can be satisfied by a NADP-dependent ACDH activity. In agreement with this hypothesis, an increased level of NADP-dependent ACDH activity was found in nitrate-grown cells in comparison to the ammonium-grown cells (Table 3). Since this cofactor is required for cellular biosynthesis, in the case of nitrate assimilation it could become limiting in cell metabolism, and this can in turn stimulate metabolic pathways able to generate it, as acetic acid formation.

Another important aspect of the redox balance is linked to NAD/NADH ratio, which is an especially critical step under anaerobic conditions. NADH is in fact generated not only by the glycolytic pathway but also by amino acid synthesis. The most important reaction for reoxidation of this surplus NADH under anaerobic conditions is the production of glycerol (Fig. 1) [30]. In *D. bruxellensis*, a very low amount of this compound was produced under those conditions [28] and the inefficiency of this pathway has been indicated as the main cause of the “Custers effect” in this species [31]. Nitrate assimilation could then accomplish the role of balancing the redox status. The higher growth rate and higher specific ethanol production rate obtained in this work under anaerobic conditions (Table 2) indicate that nitrate utilization greatly improves the fermentative metabolism in *D. bruxellensis*. In *S. cerevisiae*, it has been calculated that 13 mmol of glycerol per gram of dry biomass are generated under anaerobic conditions, leading to the reoxidation of 13 mmol of NADH [1]. In *D. bruxellensis*, the assimilation of 10 mmol of nitrate to ammonium through a NADH-dependent nitrite reductase could result in the reoxidation of 30 mmol of NADH (Fig. 1), which fits well with the theoretical formation of 26 mmol of NADH generated from the biosynthesis of 2 g of dry biomass produced during its anaerobic growth.

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

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References

- Albers E, Larsson C, Liden G, Niklasson C, Gustafsson L (1996) Influence of the nitrogen source on *Saccharomyces cerevisiae* anaerobic growth and product formation. *Appl Environ Microbiol* 62:3187–3195
- Avila J, Pérez MD, Brito N, Gonzàles C, Siverio JM (1995) Cloning and disruption of *YNRI* gene encoding the nitrate reductase apoenzyme of the yeast *Hansenula polymorpha*. *FEBS Lett* 366:137–142
- Avila J, Gonzàles C, Brito N et al (1998) Clustering of the *YNAI* gene encoding a Zn(II)2Cys6 transcriptional factor in the yeast *Hansenula polymorpha* with the nitrate assimilation genes *YNT1*, *YNI1* and *YNRI* and its involvement in their transcriptional activation. *Biochem J* 335:647–652
- Blomqvist J, Eberhard T, Schnürer J, Passoth V (2010) Fermentation characteristics of *Dekkera bruxellensis* strains. *Appl Microbiol Biotechnol* 87:1487–1497
- Blomqvist J, South E, Tiukova I, Momeni MH, Hansson H, Ståhlberg J, Horn SJ, Schnürer J, Passoth V (2011) Fermentation of lignocellulosic hydrolysate by the alternative industrial ethanol yeast *Dekkera bruxellensis*. *Lett Appl Microbiol* 53:73–78
- Blomqvist J, Nogué VN, Gorwa-Grauslund M, Passoth V (2012) Physiological requirements for growth and competitiveness of *Dekkera bruxellensis* under oxygen-limited or anaerobic conditions. *Yeast* 29:265–274
- Böer E, Schröter A, Bode R, Piontek M, Kunze G (2009) Characterization and expression analysis of a gene cluster for nitrate assimilation from the yeast *Arxula adeninivorans*. *Yeast* 26:83–93
- Boulton R, Singleton V, Bisson L, Kunkee R (eds) (1996) Principles and practices of winemaking. Chapman & Hall, New York
- Ciani M, Ferraro L (1997) Role of oxygen on acetic acid production by *Brettanomyces/Dekkera* in winemaking. *J Sci Food Agric* 75:489–495
- Conterno L, Joseph LCM, Arvik TJ, Henick-Kling T, Bisson LF (2006) Genetic and physiological characterization of *Brettanomyces bruxellensis* strains isolated from wine. *Am J Enol Vitic* 57:139–147
- De Barros Pita W, Leite FC, de Souza Liberal A, Simões DA, Morais MA Jr (2011) The ability to use nitrate confers advantage to *Dekkera bruxellensis* over *S. cerevisiae* and can explain its adaptation to industrial fermentation processes. *Antonie Leeuwenhoek* 100:99–107
- Fugelsang KC (ed) (1996) Wine microbiology. Chapman & Hall, New York
- Galafassi S, Merico A, Pizza F, Hellborg L, Molinari F, Piškur J, Compagno C (2011) *Dekkera/Brettanomyces* yeasts for ethanol production from renewable sources under oxygen-limited and low pH conditions. *J Ind Microbiol Biotechnol* 38:1079–1088
- García-Lugo P, Gonzàles C, Perdomo G, Brito N, Avila J, de la Rosa JM, Siverio JM (2000) Cloning, sequencing and expression of *HaYNRI* and *HaYNI1*, encoding nitrate and nitrite reductases in the yeast *Hansenula anomala*. *Yeast* 16:1099–1105
- Guerrero MG, Vega JM, Losada M (1981) The assimilatory nitrate-reducing system and its regulation. *Annu Rev Plant Physiol* 32:169–204
- Jeffries TW (1983) Effects of nitrate on fermentation of xylose and glucose by *Pachisolen tannophilus*. *Nature Biotechnol* 1:503–506
- Leite FC, Basso TO, de Barros Pita W, Gombert AK, Simões DA, de Morais MA Jr (2012) Quantitative aerobic physiology of the yeast *Dekkera bruxellensis*, a major contaminant in bioethanol production plants. *FEMS Yeast Res* Sep 21, p S1567 doi: 10.1111/1567-1364.12007
- Liberal ATS, Basílio ACM, Resende AM, Brasileiro BTRV, da Silva-Filho EA, Morais JOF, Simões DA, Morais MA Jr (2007) Identification of *Dekkera bruxellensis* as a major contaminant yeast in continuous fuel ethanol fermentation. *J Appl Microbiol* 102:538–547
- Machin F, Perdomo G, Pérez MD, Brito N, Siverio JM (2000) Evidence for multiple nitrate uptake systems in *Hansenula polymorpha*. *FEMS Microbiol Lett* 194:171–174

20. Merico A, Sulo P, Piškur J, Compagno C (2007) Fermentative lifestyle in yeasts belonging to the *Saccharomyces* complex. *FEBS J* 274:976–989
21. Oelofse A, Pretorius IS, du Toit M (2008) Significance of *Brettanomyces* and *Dekkera* during winemaking: a synoptic review. *S Afr J Enol Vitic* 29:128–144
22. Panagiotou G, Christakopoulos P, Grotjaer T, Olsson L (2006) Engineering of the redox imbalance of *Fusarium oxysporum* enables anaerobic growth on xylose. *Metab Eng* 8:474–482
23. Passoth V, Blomqvist J, Schnürer J (2007) *Dekkera bruxellensis* and *Lactobacillus vini* from a stable ethanol-producing consortium in a commercial alcohol process. *Appl Environ Microbiol* 73:4354–4356
24. Pereira LF, Bassi AP, Avansini SH, Neto AG, Brasileiro BT, Ceccato-Antonini SR, De Morais MA Jr (2012) The physiological characteristics of the yeast *Dekkera bruxellensis* in fully fermentative conditions with cell recycling and in mixed cultures with *Saccharomyces cerevisiae*. *Antonie Van Leeuwenhoek* 101:529–539
25. Piškur J, Ling Z, Marcet-Houben M, Ishchuk OP, Aerts A, LaButti K, Copeland A, Lindquist E, Barry K, Compagno C, Bisson L, Grigoriev IV, Gabaldón T, Phister T (2012) The genome of wine yeast *Dekkera bruxellensis* provides a tool to explore its food-related properties. *Int J Food Microbiol* 157:202–209
26. Postma E, Verduyn C, Scheffers WA, van Dijken JP (1989) Enzymatic analysis of the Crabtree effect in glucose-limited chemostat cultures of *Saccharomyces cerevisiae*. *Appl Environ Microbiol* 55:468–477
27. Rossi B, Manasse S, Serrani F, Berardi E (2005) *Hansenula polymorpha* *NMR2* and *NMR4*, two new loci involved in nitrogen metabolite repression. *FEMS Yeast Res* 5:1009–1017
28. Rozpedowska E, Hellborg L, Ishchuk OP, Orhan F, Galafassi S, Merico A, Woolfit M, Compagno C, Piškur J (2011) Parallel evolution of the make-accumulate-consume strategy in *Saccharomyces* and *Dekkera* yeasts. *Nat Commun* 2:302
29. Siverio JM (2002) Assimilation of nitrate by yeasts. *FEMS Microbiol Rev* 26:277–284
30. van Dijken JP, Scheffers AW (1986) Redox balances in the metabolism of sugars by yeasts. *FEMS Microbiol Rev* 32:199–224
31. Wijsman MR, van Dijken JP, van Kleeff BH, Scheffers WA (1984) Inhibition of fermentation and growth in batch cultures of the yeast *Brettanomyces intermedius* upon a shift from aerobic to anaerobic conditions (Custers effect). *Antonie Leeuwenhoek* 50(2):183–192
32. Woolfit M, Rozpedowska E, Piškur J, Wolfe KH (2007) Genome survey sequencing of the wine spoilage yeast *Dekkera (Brettanomyces) bruxellensis*. *Eukaryot Cell* 6(4):721–733