Physiological Characterization of a Novel Wild-Type Yarrowia lipolytica Strain Grown on Glycerol: Effects of Cultivation Conditions and Mode on Polyols and Citric Acid Production

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Abstract: A new yeast wild-type Yarrowia lipolytica isolate presented efficient growth on glycerol. During flask cultures, nitrogen limitation led to the secretion of sugar-alcohols as the major metabolites of the process (mannitol, arabitol and erythritol), whereas insignificant quantities of citrate were synthesized. Although in some instances high initial glycerol concentrations were employed (∼150 g/L), remarkable glycerol assimilation and polyol secretion was observed. Total polyols ∼ 52 g/L (conversion yield on glycerol consumed = 0.43 g/g) was recorded in the flask experiments. The sugar-alcohol production bioprocess was successfully simulated with the aid of a modified Velhust–Aggelis model that fitted very well with the experimental data, while optimized parameter values seemed to be quite consistent. In bioreactor trials, a noticeable metabolic shift towards citric acid production was observed, while simultaneously insignificant polyol quantities were produced. In fed-batch bioreactor experiments, a total citric acid quantity ∼ 102 g/L was recorded—one of the highest in the literature for wild-type Y. lipolytica strains. This metabolic transition was due to higher oxygen saturation into the medium that occurred in the bioreactor experiments compared with the flasks. Cellular lipids produced in the bioreactor trial contained higher concentrations of unsaturated fatty acids compared with those produced in flasks.

Keywords: citric acid; erythritol; glycerol; mannitol; sugar-alcohols; Yarrowia lipolytica

1. Introduction

The constant and continuous decrease in non-renewable energy feedstocks and the concomitant rise in the price of crude oil have resulted in the increment of scientific interest concerning the discovery and study of “renewable” types of energy sources. Biodiesel (viz. alky-esters of fatty acids esterified with low-molecular weight alcohols) and bioethanol are considered as the most important and interesting types of biofuels [1–5]. The synthesis of the above-mentioned biofuels is connected with the production of concentrated glycerol-containing waters, which are the principal side-products of these processes. Moreover, besides bioethanol production, important quantities of glycerol-containing
water can also be generated as side products in various alcoholic beverages' production facilities; for instance, during the typical ethanol fermentation process, ethanol is separated via distillation. The liquid fraction of the remaining material (the so-called “thin stillage”) contains c. 2% \( w/v \) of glycerol [6]. Additionally, liquid waste streams containing relatively elevated levels of glycerol are generated in the various oleochemical facilities employing transformations of vegetable or animal fats (i.e., soap-production facilities) [2,3]. Finally, even larger quantities of glycerol-containing water can be envisaged to be generated into the near future, due to the very high intra-cellular accumulation of glycerol as a storage material (in quantities up to c. 85% \( w/w \)) in several algal species like *Dunaliella* sp. In some of the above-mentioned cases, quantities of glycerol in water reaching levels up to 7.8 M (equivalent to c. 720 g of glycerol per L of water) may be generated [7].

This entire situation has inevitably negatively affected the price of glycerol feedstocks in the market volume, which presents a constant decrease. In 2007, a landmark year for the process of biodiesel production, in several EU countries including but not limited to Germany, concentrated glycerol-containing water derived from the biodiesel manufacture process, was considered as a “typical industrial waste-water”. Its price was c. 0.0 USD/kg (as opposed to pure glycerol which presented a price of c. 1.20 USD/kg that year). Thus, crude glycerol was employed only for biogas production [8]. In the years that followed, the situation has been relatively ameliorated by taking into consideration both the price and the processes of the purification of glycerol; especially in countries with an advanced technological and scientific level, the industrial production of biodiesel was accompanied (and is currently accompanied) by the side production of relatively highly purified industrial glycerol feedstocks (e.g., purity of \( \approx 85–90\% w/w \), sometimes >90% \( w/w \)). This feedstock can be used directly as the starting material for the chemical synthesis of epichlorohydrin or it can be used in various pharmaceutical applications after further purification processes are carried out [1,4].

In any case though, the appearance of glycerol as a side-product in the various agro-food, biofuel and industrial activities mentioned above, is constantly rising. Therefore, the conversion of this low-value compound into several types of biotechnological products currently attracts enormous attention [3–5,9,10]. Wild-type, mutant (i.e., acetate negative ones) and genetically engineered strains of the non-conventional polymorphic yeast *Yarrowia* (synonyms: *Candida—Saccharomycopsis—Endomycopsis* *lipolytica*) have been successfully used in recent years as microbial cell factories, amenable to convert glycerol into a plethora of valuable interesting compounds, including but not limited to microbial oils, organic acids (i.e., citric acid, succinic acid, acetic acid), yeast mass, sugar alcohols (i.e., mannitol, erythritol, arabitol), etc. [3,10–15].

Recent investigations by our research team have indicated that after an initial screening study, a wild-type novel isolate of the non-conventional polymorphic yeast *Y. lipolytica*, namely strain LMBF-Y46, has been revealed as being capable to produce amounts of polyols (mainly mannitol), during a screening study in shake-flask experiments on biodiesel-derived glycerol [16]. It was decided to further quantitatively study the physiological behavior of the above-mentioned strain, using a nitrogen-limited media based on crude or pure glycerol, and also in cultures with increased initial concentrations of glycerol, in shake flasks. Moreover, a scaled-up laboratory-scale stirred tank bioreactor production system was produced. Considerations concerning the yeast physiological behavior were comprehensively assessed and discussed.

2. Materials and Methods

2.1. Microorganism, Media and Culture Conditions

The yeast strain used in the current study was a new indigenous wild-type *Y. lipolytica* strain, coded LMBF Y-46, isolated from gilt-head (sea) bream (*Sparus aurata*) fish [17], having been previously demonstrated as capable of producing significant amounts of yeast biomass and mannitol in a first screening-type study [16]. The maintenance of the strain as well as the salt composition of the basal medium used for the fermentations, have both previously been described [18,19]. The nitrogen sources
employed were (NH$_4$)$_2$SO$_4$ (containing c. 21%, w/w, of nitrogen) and yeast extract (containing c. 7%, w/w, of nitrogen and c. 8%, w/w, of carbon). Initially, (NH$_4$)$_2$SO$_4$ at 1.5 g/L and yeast extract at 1.0 g/L were employed; at this stage, a relatively low initial glycerol concentration (Gly$_0$) was employed (~40 g/L). Thereafter, and in order to enhance the production of yeast biomass, in addition to potentially enhancing glycerol assimilation, higher initial nitrogen concentrations were employed; in this second set of cultures, (NH$_4$)$_2$SO$_4$ at an initial concentration of 2.0 g/L and yeast extract at an initial concentration of 3.0 g/L were employed [18]. Additionally, higher initial concentrations of glycerol were used (up to c. 150 g/L). The carbon sources used in all trials were either industrial (biodiesel-derived) crude glycerol (Agroinvest SA; Acharia, Fithiotis prefecture, Greece) with a purity of c. 81% w/w [18] or pure glycerol (Prolabo Chemicals, Fontenaysous bois, France, purity c. 98% w/w). The impurities in industrial glycerol were composed of 11–12% w/w water, 5–6% w/w potassium salts, 1% w/w free-fatty acids and <0.2% w/w methanol. The initial pH of all media was 6.0 ± 0.1.

Shake-flask trials were conducted batch-wise in Erlenmeyer non-baffled flasks of 250 mL filled with 50 ± 1 mL of culture medium, previously sterilized (at $T = 115^\circ$C; 20 min) and inoculated with 1 mL of exponential pre-culture containing 1–3 $\times 10^6$ cells (the initial cell mass concentration—$X_0$ was ≈0.10 g/L) [18,19]. The pre-culture was carried out in liquid-modified YPD medium, in which dextrose had been replaced by glycerol (the pre-culture medium contained yeast extract at 10.0 g/L, peptone at 10.0 g/L and pure glycerol at 20.0 g/L). Pre-cultures were incubated in an orbital shaker (for 24 ± 2 h at 180 ± 5 rpm, $T = 28 ± 1^\circ$C) after inoculation of a loop by a freshly prepared slant of the strain. After inoculation by the pre-culture, flasks were incubated at $T = 28 ± 1^\circ$C at 180 ± 5 rpm in a rotary shaker (New Brunswick Innova 43 Incubator Shaker; NH, USA). In the experiments performed in shake-flask mode, a constant drop in the pH value of the medium was observed throughout the culture time, and the pH value was always maintained in the flasks at values of ≥4.5 ± 0.1 by aseptically adding 2M NaOH (for more details see: Papanikolaou et al. [19]). In most of the culture steps and for most of the shake-flask experiments performed, the medium pH remained at >4.8, being found within the range 4.8–5.8.

Batch and pulsed fed-batch bioreactor experiments were also performed, in a bench-top 3 L bioreactor (Applikon Biotechnology; Delft, The Netherlands), containing 1.9 L of previously sterilized fermentation medium (at $T = 115^\circ$C; 45 min) and inoculated with 100 mL of 24 h exponential Y. lipolytica pre-culture (see previously for the pre-culture details). Culture conditions in the bioreactor experiments were as in Papanikolaou et al. [18]: incubation temperature $T = 28 ± 1^\circ$C; agitation = 800 ± 5 rpm; aeration set between 0.1 and 2.0 vvm in order to maintain a dissolved oxygen tension (DOT) in the vessel above 20%, during all culture phases, thus full aerated conditions were imposed (see Palaiogeorgou et al. [20]); pH = 6.0 ± 0.2 regulated with automatic addition of concentrated NaOH (5 M). For the pulsed fed-batch bioreactor experiment, pulses of concentrated and previously sterilized glycerol quantities were added into the vessel at various incubation times, and all fermentation parameters were the same as in the batch bioreactor trial.

2.2. Analyses

In order to proceed to the various analyses, the whole content of the flasks (~46–48 mL) or a sample ±20 mL of the bioreactor experiments was collected. In the case of the shake-flask trials, due to slight water evaporation, the volume of the liquid medium in the flasks after harvesting was 47 ± 1 mL. Certainly, the more the culture duration was elevated, the more water evaporation of the medium occurred. Nevertheless, the volume of the medium was never lower than 46 mL. This volume was always corrected to 50 mL and the whole content of the flask was used in the performed analyses. Regardless of the volume of the sample, yeast cells were harvested by centrifugation, and yeast total biomass quantity was determined through its dry cell weight (DCW, g/L) [16,21]. Non-consumed glycerol (Gly), citric acids (citric + iso-citric acid) (CA), mannitol (MI), arabitol (Ara) and erythritol (Ery) were determined with the aid of HPLC analysis [18,22]. In several fermentation points, iso-citric was assayed according to Papanikolaou et al. [23]. Cellular lipids were extracted from the yeast DCW
according to the modified “Folch” method [16,24], while total microbial oil was recovered and trans-esterified according to Gardeli et al. [24]. The produced fatty acid methyl-esters (FAMEs) were suspended in hexane and analyzed by GC analysis in a Varian CP-3800 chromatograph equipped with flame ionization detector (Agilent Technologies, Santa Clara, CA, USA), in which an Agilent J&W Scientific DB23 capillary column (model n.123–2332, 30.0 m × 0.32 mm, film thickness 0.25 μm) (Agilent Technologies, Santa Clara, CA, USA) was used. Helium gas was used as a carrier gas with a column flow rate of 2.0 mL/min. The set up conditions were as follows: initial oven temperature was set at $T = 150 \, ^\circ\text{C}$, held for 18 min, subsequently ramped to $T = 185 \, ^\circ\text{C}$ at a rate of 5 °C/min and held for 2 min. Then the oven temperature was moved to $T = 210 \, ^\circ\text{C}$ at a flow rate of 5 °C/min and held for 2 min, then increased to $T = 240 \, ^\circ\text{C}$ at 10 °C/min. The injector and flame ionization detector temperatures were set at $T = 260 \, ^\circ\text{C}$ and $T = 270 \, ^\circ\text{C}$, respectively. Individual FAMEs were identified by comparison of their retention times with external standard (Supelco 37 Component FAME Mix, CRM47885) retention times. The amounts of individual FAMEs identified were expressed in % of the total fatty acid areas chromatograms identified.

In the case of the shake-flask cultures and for various fermentation points, the DOT was determined according to the procedure proposed in Palaiogeorgou et al. [20]. In all of the cases in which DOT was measured (measurements were conducted at the phase in which polyols were accumulated into the medium), and for all of the $\text{Gly}_{t_0}$ quantities set into the medium, DOT values were between 20% and 50% v/v. Moreover, in several of the shake-flask cultures performed, ammonium ion concentration into the medium was determined with an ammonium selective electrode (SA 720, Orion; USA), according to previously indicated procedure [25].

In all the kinetic experiments both in shake-flask and bioreactor trials, each experimental point presented in the figures and the tables is the mean value of two measurements deriving from two independent experiments (SE in most of the experimental points was ≤18%).

Nomenclature: $X$: Biomass concentration (dry cell weight—DCW, g/L); $\text{Gly}$: Glycerol concentration (g/L); $t$: reaction time (h); $\text{CA}$: Citric acids (citric + iso-citric acid) concentration (g/L); $\text{MI}$: Mannitol concentration (g/L); $\text{Ara}$: Arabitol concentration (g/L); $\text{Ery}$: Erythritol concentration (g/L); $\text{Pol}$: Total polyols concentration (g/L); $g_{\text{pol}}$: specific polyols formation rate, g of polyols per g of biomass per h; $Y_{\text{Pol/Gly}}$: yield coefficient for total polyols production with respect to glycerol consumed, $g_{\text{pol}}/g_{\text{Gly}}$ (g of total polyols produced per g of glycerol consumed); $Y_{\text{CA/Gly}}$: yield coefficient for citric acid production with respect to glycerol consumed, $g_{\text{CA}}/g_{\text{Gly}}$ (g of citric acid produced per g of glycerol consumed); $Y_{\text{CA/Gly}}$: yield coefficient for DCW production with respect to glycerol consumed, $g_{\text{CA}}/g_{\text{Gly}}$ (g of DCW produced per g of glycerol consumed); $r_{\text{CA}}$: rate of citric acid production ($g_{\text{CA}}$ per L per h); $r_{\text{Gly}}$: rate of glycerol consumption ($g_{\text{Gly}}$ per L per h); $r_{\text{DCW}}$: rate of biomass production ($g_{\text{X}}$ per L per h); $\mu$: specific biomass production rate (h⁻¹), calculated in batch cultures by the formula $\mu_{\text{max}} = \ln\left(\frac{X_1}{X_0}\right) / t_{10} - t_0$, where $X_1$ was the first experimental point after inoculation and $t_{10}$ was the respective time; $X_0$ was the initial DCW quantity and $t_0 = 0$ h; $P_{\text{CA}}$: Citric acids volumetric productivity ($g_{\text{CA}}$ per L per h); indexes 0 and cons refer to the initial and the consumed quantity of the elements in the experiments carried out; index max refers to the maximum value of a particular parameter in the cultures.

3. Results and Discussion

3.1. Metabolic Characterization of Yarrowia lipolytica LMBF Y-46: Effect of Different Glycerol Sources under Nitrogen Limitation

In the first part of this report, it was desirable to evaluate the performances of $Y$. lipolytica LMBF Y-46 in shake-flask experiments in media in which pure or crude glycerol ($\text{Gly}_{t_0} \approx 40 \, g/L$) had been employed. Initial nitrogen concentration used in the medium was $(\text{NH}_4)_2\text{SO}_4$ at 1.5 g/L and yeast extract at 1.0 g/L (initial molar ratio ≈ 48 moles/moles). The kinetics of both experiments are shown in Figure 1a–d. For both cases, DCW production almost reached a plateau ($X = 10–11 \, g/L$) at $t = 120$ h after inoculation. Onset of the secretion of polyols had been given earlier ($t \approx 50–55$ h
were synthesized, in notably lower quantities, were erythritol and to lesser extent arabitol. For both points, cultures were performed under nitrogen-limited conditions, providing evidence that the onset of polyols and DCW that seemed not to differ by the utilization of the pure or the crude substrate. For both fermentations, \( r_{\text{Gly}} \) was \( 0.24 \text{ g/L/h} \) (Figure 1b). This result, together with the observations concerning the production of polyols and DCW that seemed not to differ between the two fermentations carried out, indicate that the impurities of the substrate did not seem to have any important impact upon the physiological

Figure 1. Kinetics of dry yeast cell mass production (\( X, \text{g/L} \)) (a), glycerol assimilation (\( \text{Gly}, \text{g/L} \)) (b), total polyols synthesis (\( \text{Pol}, \text{g/L} \)) and (c) lipid in dry cell weight (\%, \text{w/w}), (d) by \text{Yarrowia lipolytica LMBF Y-46}, during growth on pure and biodiesel-derived glycerol in shake-flask experiments under nitrogen-limited conditions. Culture conditions: growth in 250 mL conical flasks at 180 ± 5 rpm, initial pH = 6.0 ± 0.1, pH ranging between 4.5 and 6.1, incubation temperature \( T = 28 ± 1 ^\circ \text{C} \). Each experimental point is the mean value of two independent determinations (SE ≤ 18%). The assimilation of glycerol for both trials was almost linear (see kinetics of remaining into the medium glycerol in Figure 1b) and the rate of glycerol consumption (\( r_{\text{Gly}} \)), calculated after linear regression of the experimental data of remaining in the medium glycerol vs. time for both trials, seemed unaffected by the utilization of the pure or the crude substrate. For both fermentations, \( r_{\text{Gly}} \) was \( 0.24 \text{ g/L/h} \) (Figure 1b). This result, together with the observations concerning the production of polyols and DCW that seemed not to differ between the two fermentations carried out, indicate that the impurities of the substrate did not seem to have any important impact upon the physiological
behavior of the strain. In agreement with these results, for a number of wild-type microbial strains (i.e., Klebsiella oxytoca, Mortierella ramanniana, some strains of the species Y. lipolytica) cultivated on glycerol deriving from various origins, the impurities of the feedstock did not seem to have any impact upon the physiological behavior of the mentioned microorganisms [22,23,26,27]. In contrast, in other cases (i.e., Clostridium butyricum, C. pasteurianum, Meyerozyma caribbica, Rhodolorula minuta, other strains belonging to the species Y. lipolytica, etc.), glycerol impurities seemed to (drastically) modify the cellular metabolism of the implicated microorganisms [28–34]. Therefore, the impurities found in the crude glycerol feedstock did not seem to have a clear positive or negative effect upon the production of dry yeast biomass and metabolites by Y. lipolytica yeast. However, in some cases it must be pointed out that the impurities found in the medium seemed to enhance the production of the metabolites by various yeast strains including Y. lipolytica [29,32–34]. From the above-mentioned analysis, therefore, it seems that the biosynthesis of metabolites by Y. lipolytica strains cultivated on glycerol, appears to be strain dependent in relation to the origin of the feedstock implicated in the bioprocess performed.

According to theory, polyols are synthesized as secondary anabolic compounds with their biosynthesis being coupled with nitrogen limitations in the growth medium [3,10,13–15,35–37]. Though, the current investigation, as well as other limitations in the literature [18,21] indicate that the total biomass biosynthesis may occur simultaneously with the biosynthesis of polyols into the medium. In fact, in the current investigation it was seen that typically after nitrogen depletion from the medium (at t ≈ 50–55 h after inoculation), polyols were secreted, while an almost linear profile of polyols biosynthesis occurred between 50 and 150 h after inoculation (see Figure 1c). On the other hand, in spite of nitrogen depletion that, as stated occurred at t ≈ 50–55 h after inoculation (at that fermentation step X concentration was ≈7.0 g/L), some supplementary biomass production occurred between 50–55 h and t = 120 h after inoculation; as indicated, DCW production plateaued at t = 120 h, and when biomass concentration reached its plateau X concentration was ≈10–11 g/L. This supplementary total biomass production that occurred at the late growth stages did not indicate that cells “fattened” as fermentation proceeded; in fact, insignificant quantities of total lipids were accumulated inside the yeast cells for both trials performed, despite the fact that cultures were performed under nitrogen limitation (Figure 1d). The kinetic results of the current investigation indicate that somehow elevated lipid quantities were synthesized at the relatively early growth stage of the culture, with lipid in DCW values decreasing as the fermentation proceeded (Figure 1d) This is in accordance with various literature reports in which Y. lipolytica strains (wild-type or mutant ones) had been cultured batch-wise in glycerol-based media [18,38–40]. The total dry biomass production increase that occurred after nitrogen had been depleted from the culture (viz. between 50–55 h and 150 h after inoculation), could suggest the accumulation of intra-cellular polysaccharides inside the cells occurring after nitrogen deprivation from the medium. This assumption is in accordance with the reports of several microorganisms like other Y. lipolytica strains or other yeast species/genera like Apiotrichum curvatum, Rhodospiridium toruloides, Metschnikowia sp. [16,37,41–43].

3.2. Metabolic Characterization of Yarrowia lipolytica LMBF Y-46: Effect of Glycerol Concentration in Media with Constant Initial Nitrogen in Shake-Flask Trials

In the next approach, and in order to more profoundly quantify the metabolism of Y. lipolytica LMBF Y-46, another set of shake-flask cultures was carried out. In this set, a higher initial nitrogen concentration was set into the medium (ammonium sulfate and yeast extract at an initial concentration of 2.0 and 3.0 g/L, respectively) in order to further “boost” the production of the total biomass, and potentially to increase the consumption rate of glycerol. Variable Gly0 concentrations with higher initial substrate quantities than the previous experiment, were tested (Gly0 quantities adjusted between ≈70 and ≈150 g/L). It was also decided to utilize the pure substrate as a carbon source (pure glycerol therefore was employed instead of crude). This decision was taken in order to ascertain that all potential changes observed in the quantitative cellular metabolism in these experiments were due only to the increment in Gly0 concentration (and also to the increase in the initial C/N ratio) employed
into the medium and not to the potential accumulation of crude glycerol impurities into the medium, specifically at the higher Gly<sub>0</sub> concentrations imposed. The obtained results of this set of experiments are shown in Table 1. As was anticipated, higher total DCW quantities were produced when compared with the previous experiment, since increased initial nitrogen concentrations were employed in the former case. Likewise, glycerol consumption was equally linear for this series of trials; linear regression of the experimental data of remaining glycerol concentration vs. incubation time for the points in which non-negligible glycerol quantity remained un consumed into the medium, revealed a glycerol consumption rate value (r<sub>Gly</sub>, g/L/h) (see also Section 3.1). This value noticeably decreased as the Gly<sub>0</sub> concentration increased in the medium. However, the r<sub>Gly</sub> value was always higher than in the previous set of experiments in which a lower initial nitrogen concentration was employed; in all trials and irrespective of the Gly<sub>0</sub> concentration in the medium, linear regression of the remaining glycerol (Gly<sub>f</sub>) f(t) revealed an r<sup>2</sup> value ≥ 0.95 (linear regressions not presented) providing evidence of the linearity of consumption of glycerol by Y. lipolytica, with r<sub>Gly</sub> ranging between 0.54 g/L/h (Gly<sub>0</sub> ≈ 70 g/L) and 0.37 g/L/h (Gly<sub>0</sub> ≈ 150 g/L).

Table 1. Quantitative data of cultures of Yarrowia lipolytica LMBF Y-46 on glycerol-based media in which various initial glycerol concentrations (Gly<sub>0</sub>) were added with nitrogen being added to the medium as ammonium sulfate (2.0 g/L) and yeast extract (3.0 g/L). Culture conditions: growth in 250 mL conical flasks (at 180 rpm, initial pH = 6.0 ± 0.1, pH ranging between 4.4 and 6.1, incubation temperature T = 28 ± 1 °C). Culture details when the maximum concentration of mannitol (Ml, g/L) (a), erythritol (Ery, g/L) (b), arabitol (Ara, g/L) (c) and yeast dry cell biomass (X, g/L) (d) was achieved. Pol is the sum of polyols (in g/L) at the given fermentation times. Each experimental point presented is the mean value of two independent determinations.

| Gly<sub>0</sub> (g/L) | Time (h) | Gly<sub>cons</sub> (g/L) | r<sub>Gly</sub> (g/L/h) | X (g/L) | Ml (g/L) | Ery (g/L) | Ara (g/L) | Pol (g/L) | Y<sub>pol/Gly</sub> (g/g) |
|---------------------|---------|-----------------------|----------------------|--------|---------|----------|---------|--------|-------------|
| ≈70                 | a, b, c | 134                   | 65.0 ± 4.5           | 0.54   | 16.1 ± 1.5 | 9.2 ± 1.4 | 7.3 ± 1.2 | 4.1 ± 0.7 | 20.6 ± 3.3 | 0.32 |
|                     | d       | 183                   | 65.6 ± 3.9           | 22.4 ± 2.0 | 6.1 ± 0.8 | 3.1 ± 0.6 | 1.9 ± 0.4 | 11.1 ± 1.8 | 0.17 |
| ≈100                | a, b, c, d | 183                  | 95.4 ± 7.1           | 0.52   | 15.0 ± 1.9 | 13.8 ± 2.2 | 17.1 ± 2.5 | 7.3 ± 1.2 | 38.2 ± 5.9 | 0.40 |
| ≈120                | a, b, c, d | 250                  | 109.2 ± 9.0          | 0.45   | 13.5 ± 2.1 | 15.7 ± 2.4 | 20.6 ± 2.8 | 7.1 ± 1.1 | 43.4 ± 6.3 | 0.40 |
| ≈150                | a, b, c, d | 280                  | 120.5 ± 12.8         | 0.37   | 11.1 ± 1.8 | 13.1 ± 2.0 | 24.1 ± 4.0 | 14.1 ± 2.1 | 51.3 ± 8.1 | 0.43 |

Despite the high concentration of glycerol in some runs (i.e., in the experiment with the highest Gly<sub>0</sub> concentration into the medium this quantity was ≈155 g/L), the microorganism presented an appreciable DCW production, providing evidence again of the suitability of this substrate for strains of this species [8,11,16,29,44–48]. Though, the DCW<sub>max</sub> quantity somehow decreased with the Gly<sub>0</sub> concentration increase in the medium, indicating the potential inhibition exerted due to the high Gly<sub>0</sub> quantity added into the medium (in accordance with reports for other Y. lipolytica strains; see i.e., Papanikolaou et al. [8,18,21]). In all cases, and despite the fact that the pH in the medium remained always in values ≥4.5 ± 0.1, carbon catabolism was mostly directed towards the synthesis of polyols, while indeed low quantities of organic acids (mostly citric acid) were found into the medium; CA concentrations were ≤3.0 g/L for all culture phases and all Gly<sub>0</sub> concentrations rested. In most cases in the literature in which Y. lipolytica strains have grown under nitrogen limitation on glycerol-based media and medium pH remaining >4.5, this event normally favored the biosynthesis and secretion into the medium of citric acid. In contrast, relatively low quantities of polyols were simultaneously synthesized [8,23,40,46,48,49]. However, in agreement with the behavior of the strain LMBF Y-46 that mostly produced polyols instead of citrate, despite the maintenance of pH in values ranging between 4.5–6.1, in recent years, some reports have indicated similar physiological behaviors of other wild-type Y. lipolytica strains cultivated on slightly acidic conditions (medium pH between 4.5–6.5) with glycerol employed as the substrate, in which mostly polyols have been synthesized [15,16,18,36,37]. As indicated, the glycerol uptake rate noticeably decreased with Gly<sub>0</sub> increment into the medium. Despite significant extension of the culture time (i.e., up to 280 h that corresponds to ≈11.5 days) for the
high Gly$_0$ concentration media, glycerol was not fully consumed (see Table 1). Nevertheless, by taking into consideration that in high Gly$_0$ concentration media in which glycerol had not been completely assimilated (Table 1) glycerol uptake was always constant, it is strongly possible that further incubation would decrease the content of glycerol even more and increase the Pol biosynthesis in these cultures.

Y. lipolytica LMBF Y-46 growth on glycerol in shake flasks with various Gly$_0$ concentrations into the medium was performed under full aerobic conditions; DOT was measured mainly at the phase in which polyols were secreted into the medium, and irrespective of the Gly$_0$ quantities added into the medium, DOT values ranged between 20% and 50% v/v. On the other hand, the more the Gly$_0$ quantity into the medium increased, the more Pol concentration increased (Table 1), in accordance with results indicated for other wild-type Y. lipolytica strains [18,21,22]. Furthermore, in the experiment with the somehow lower Gly$_0$ concentration added into the medium (Gly$_0$ ≈ 70 g/L), when the glycerol concentration from the medium was exhausted, polyols re-consumption and new yeast cell biomass occurred (Table 1). Re-consumption of extra-cellular and previously synthesized polyols in favor of the biosynthesis of new cell material (or potentially the production of other metabolites, like citric acid) has been observed in batch cultures performed by various wild-type Y. lipolytica strains [20–22,35,38]. Moreover, as stated, when higher Gly$_0$ concentrations were employed, in absolute values (viz. g/L), enhanced production of polyols occurred. It can be hypothesized that in the trials with high Gly$_0$ concentrations, either the increment of carbon excess or the increased osmotic pressure into the medium due to high Gly$_0$ quantities [2,4,10,13] enhanced Pol production in Y. lipolytica LMBF Y-46. On the other hand, the total conversion yield of the sum of polyols (Pol, g/L) produced per unit of glycerol consumed (Y$_{pol/Gly}$, g/g) originated from linear regression of all of the experiments points of Pol produced f (glycerol consumed) for the batch-flask runs at Gly$_0$ ≈ 100, 120 and 150 g/L, seemed to be constant and ≈0.43 g/g (Figure 2).

![Figure 2](image-url)  
**Figure 2.** Total polyols (Pol, g/L) produced vs. glycerol consumed (g/L) in shake-flask experiments during growth on pure glycerol by Yarrowia lipolytica LMBF Y-46, at elevated initial glycerol concentrations (Gly$_0$ ≈ 100, 120 and 150 g/L) under nitrogen-limited conditions. Culture conditions: growth on 250 mL conical flasks at 180 ± 5 rpm, initial pH ≈ 6.0 ± 0.1, pH ranging between 4.5 and 6.1, incubation temperature T = 28 ± 1 °C. Each experimental point is the mean value of two independent determinations (SE ≤ 18%).

In the above-mentioned figure, the results concerning the production of polyols in the trial in which Gly$_0$ ≈ 70 g/L were not incorporated, since in the latter case notably lower total polyols quantities, accompanied by remarkably depleted yield Y$_{pol/Gly}$ values and higher yeast dry biomass quantities compared with the experiments where Gly$_0$ ≈ 100, 120 and 150 g/L, were recorded (see Table 1). Moreover, as in the previously mentioned flask cultures (“effect of different glycerol sources under...
nitrogen limitation”), in this second set of experiments ($Gly_0 \approx 70, 100, 120$ and $150$ g/L) ammonium nitrogen was depleted from the medium at $t \approx 50–55$ h after inoculation; in the second set of trials, the initial $NH_4^+$ concentration was $= 550$ mg/L, while $NH_4^+$ concentration at $t \approx 50–55$ h after inoculation became $= 25 \pm 10$ mg/L, and therefore, nitrogen limitation, accompanied by polyols secretion was noted. In the early stages of the cultivation ($t \leq 50$ h after inoculation), thus, no polyols biosynthesis was observed. Nevertheless, and despite the fact that at the early culture steps glycerol did not contribute to polyols production, the representation of $Pol$ produced $f$ (glycerol consumed) and the linear regression performed for the batch-flask runs carried out makes sense, as most of the experimental points were found after nitrogen depletion where glycerol assimilation mostly contributed to the synthesis of polyols, whereas the correlation coefficient of the regression ($r^2 = 0.91$) is quite satisfactory (Figure 2).

The metrics of polyols produced by yeasts cultivated on glycerol-based media and belonging to *Yarrowia* and *Candida* clades, and their comparisons with the current investigation, are presented in Table 2.

### Table 2. Metrics of production of polyols by several yeasts belonging to *Yarrowia* and *Candida* clades during growth on glycerol in several fermentation modes. Comparison with the current study.

| Strain                  | Ery (g/L) | Ml (g/L) | Ara (g/L) | Pol (g/L) | $Y_{Pol Gly}$ (g/g) | Cultivation Type     | Reference     |
|-------------------------|-----------|----------|-----------|-----------|---------------------|----------------------|---------------|
| 1.23 e                   | 93.5      | 34.0     | -         | 127.5     | 0.43                | Fed-batch reactor    | Rymowicz et al. [46] |
| Wratislavia 1.31 *       | 132.0     | 23.0     | -         | 155.0     | 0.52                | Fed-batch reactor    | Rymowicz et al. [46] |
| Wratislavia K1 *         | 170.0     | 12.0     | -         | 182.0     | 0.60                | Fed-batch reactor    | Rymowicz et al. [46] |
| CCY-29-26-3 b            | 23.0      | 2.6      | 2.3       | 27.9      | 0.40                | Batch flasks         | Tomaszewska et al. [29] |
| CCY-29-26-4 b            | 26.7      | 1.0      | 2.2       | 29.9      | 0.40                | Batch flasks         | Tomaszewska et al. [29] |
| A-15 e                   | 71.0      | 8.0      | 1.8       | 80.8      | 0.50                | Batch reactor        | Tomaszewska et al. [29] |
| A UV1 *                  | 63.0      | 8.8      | 9.2       | 81.0      | 0.50                | Batch reactor        | Tomaszewska et al. [29] |
| Wratislavia K1 *         | 80.0      | 2.6      | 0.3       | 82.9      | 0.51                | Batch reactor        | Tomaszewska et al. [29] |
| Wratislavia K1 *         | 135.5     | 3.9      | 0.1       | 139.5     | 0.58                | Repeated-batch reactor  | Mirończuk et al. [31] |
| Wratislavia K1 *         | 208.0     | 0.2      | -         | 208.2     | 0.41                | Repeated-batch reactor  | Mirończuk et al. [31] |
| Wratislavia 1.31 *       | 26.2      | 16.8     | 3.7       | 46.7      | 0.36                | Batch reactor        | Tomaszewska et al. [48] |
| Wratislavia AWCG7 *      | 25.7      | 17.1     | 2.7       | 45.5      | 0.30                | Batch reactor        | Tomaszewska et al. [48] |
| Wratislavia K1 *         | 40.7      | 15.1     | 2.9       | 58.7      | 0.40                | Batch reactor        | Tomaszewska et al. [48] |
| MK1 *                    | 79.5      | 2.7      | 0.4       | 82.6      | 0.55                | Batch reactor        | Mirończuk et al. [50] |
| MK1 *                    | 138.8     | 3.3      | 0.3       | 142.4     | 0.69                | Repeated-batch reactor  | Mirończuk et al. [50] |
| M1 *                     | 177.3     | 2.2      | -         | 179.5     | 0.67                | Repeated-batch reactor  | Mirończuk et al. [50] |
| CBS01046 4               | 44.6      | 5.2      | 10.5      | 60.3      | 0.53                | Batch reactor        | Rakicka et al. [51] |
| CBS4855 4                | 33.4      | 7.6      | 6.8       | 47.8      | 0.43                | Batch reactor        | Rakicka et al. [51] |
| CBS11013 4               | 35.4      | 0.6      | 8.9       | 44.9      | 0.41                | Batch reactor        | Rakicka et al. [51] |
| FCY 214 *                | 79.4      | n.i.     | n.i.      | 79.4      | 0.48                | Batch reactor        | Carly et al. [35] |
| FCY 218 *                | 80.6      | n.i.     | n.i.      | 80.6      | 0.53                | Batch reactor        | Carly et al. [35] |
| HA 829 4                 | = 4       | = 28     | = 6       | = 38      | n.i.                | Batch reactor        | Egermeier et al. [36] |
| HA 1251 4               | = 4       | = 32     | = 5       | = 41      | n.i.                | Batch reactor        | Egermeier et al. [36] |
| ACA YC 5029 4            | 33.6      | 28.9     | -         | 62.5      | 0.45                | Batch flasks         | Papanikolaou et al. [18] |
| ACA YC 5030 4            | 35.5      | 32.1     | -         | 67.6      | 0.49                | Batch flasks         | Papanikolaou et al. [18] |
| ALB 4                    | 56.7      | 12.6     | 6.0       | 75.3      | 0.49                | Fed-batch reactor    | Rakicka et al. [52] |
| ALB pADUTGUT1 *          | 82.2      | 11.0     | 7.5       | 100.7     | 0.67                | Fed-batch reactor    | Rakicka et al. [52] |
| ACA-DC 5029 4            | 65.8      | 6.5      | 3.4       | 75.7      | 0.44                | Batch flasks         | Sarris et al. [53] |
| ACA-DC 5029 4            | 15.6      | 10.5     | 3.4       | 29.5      | 0.39                | Batch flasks         | Sarris et al. [49] |
| MS3-S *                  | 69.9      | 10.1     | -         | 80.0      | 0.80                | Batch reactor        | Liu et al. [54] |
| MS3-S *                  | 72.5      | 10.0     | -         | 82.5      | 0.82                | Batch reactor        | Liu et al. [54] |

LMBF Y-46 24.1 13.1 14.1 51.3 0.43 Batch flasks Present study

*: Mutant or genetically modified *Yarrowia lipolytica*; e: Wild-type *Yarrowia lipolytica*; f: *Candida oslonensis*; 4: *Candida hollandica*; 4: *Yarrowia divulgata*; 4: Fermentations in which the medium pH remained >4.5 throughout the culture (in all other trials, medium pH was 3.0–3.5); n.i.: Not indicated in the manuscript; In most cases, the presented $Y_{Pol Gly}$ values of the various literature sources, were already indicated in the relevant manuscripts.

With the exception of a scarce number of reports [50,52,54], the yield $Y_{Pol Gly}$ has rarely been >0.60 g/g and in most cases it ranged between 0.40 and 0.55 g/g, in accordance with the results of the current submission. The majority of the studies dealing with the production of polyols by *Y. lipolytica* strains, has been carried out with the use of mutant (i.e., acetate negative ones) or genetically engineered strains, that have been reported to produce, in many cases, relatively or even much higher $Pol$ quantities compared to the present study (see Table 2). On the other hand, strain LMBF Y-46 is wild-type novel yeast isolate, originated from fish-origin products [17], that has not been previously studied in detail,
especially concerning its physiological behavior. Moreover, for almost all of the reports indicated in Table 2, trials had been performed in pH values ≈ 3.0–3.5, whereas in the current investigation the pH value was ≈ 2.45, therefore, it is attainable to obtain higher Pol quantities with the use of the LMBF Y-46 strain after further culture optimization, specifically as regards growth in lower pH values.

3.3. Metabolic Characterization of Yarrowia lipolytica LMBF Y-46: Scale-Up in Laboratory-Scale Bioreactor

In the next step of this investigation, the bioprocess of glycerol fermentation by Y. lipolytica LMBF Y-46 was scaled-up in a bench-top laboratory-scale bioreactor (active process volume = 2.0 L) which allowed fine pH control and dissolved oxygen measurement. Trials with Gly_0 concentration adjusted to ≈ 100 g/L were performed ((NH₄)₂SO₄ at 2.0 g/L and yeast extract at 3.0 g/L were set into the medium). The choice of Gly_0 = 100 g/L was based on the results indicated in Table 1; precisely, in flask cultures with this Gly_0 concentration, glycerol was almost completely consumed by the microorganism in a not excessively high incubation time (c. 180 h), while equally, DCW_max concentration was substantially high not being seriously influenced by the inhibition exerted due to high Gly_0 concentrations into the medium (see Table 1). The obtained results in the equivalent trials between the shake-flask and the bioreactor culture are depicted in Table 3, while the evolution of DCW and glycerol in the two experiments is presented in Figure 3a,b. From the obtained results it can be indicated that drastically different kinetic and physiological profiles were recorded between the two experiments; in the bioreactor trial, much higher total biomass production occurred. This biomass value was obtained significantly more rapidly than the flask culture. On the other hand, the μ_max value between the shake-flask and the bioreactor trial (for the calculation see Section 2.2) was not substantially different between the two experiments (μ_max ≈ 0.16 h⁻¹ for both performed cultures). In contrast, the biomass production rate r_X (≡ ΔX/Δt in g/L/h; calculation performed when X obtained its maximum value), was drastically higher in the bioreactor experiment compared with the shake-flask one (viz. 0.506 vs. 0.08 g/L/h). Moreover, glycerol consumption was drastically higher in the bioreactor trial compared with the flask culture (see Table 3 and Figure 3b). As previously, fitting of the experimental data through linear regression of the remaining glycerol concentration f (fermentation time) demonstrated linear glycerol consumption rate r_Gly (albeit, r² value was only 0.90 for the bioreactor experiment vs. 0.99 for the shake-flask trial; see Figure 3b) that was 0.52 g/L/h for the flask experiment and >3-fold time greater for the bioreactor culture (the T_Gly value for the latter case was = 1.85 g/L/h).

**Table 3.** Quantitative data of cultures of Yarrowia lipolytica LMBF Y-46 on glycerol-based media in which initial glycerol concentration (Gly_0) = 100 g/L was added (nitrogen was added as ammonium sulfate at 2.0 g/L and yeast extract at 3.0 g/L) in shake-flask and batch bioreactor experiments. Culture conditions: for the flask experiment as in Table 1. For the bioreactor experiment: agitation 800 ± 5 rpm, pH = 6.0 ± 0.1, aeration fluctuating from 0.1 up to 2.0 vvm in order to have a DOT ≥ 10% qTV; incubation temperature T = 28 ± 1 °C, active volume = 2.0 L. Culture details when the maximum concentration of polyols (Pol, g/L) (a), citric acid (CA, g/L) (b) and yeast dry cell biomass (X, g/L) (c) was achieved. MI, Ery and Ara is the concentration of mannitol, erythritol and arabitol, respectively (in g/L), at the given fermentation times. Each experimental point presented is the mean value of two independent determinations.

| Gly₀ (g/L) | Time (h) | Gly₀_final (g/L) | X (g/L) | MI (g/L) | Ery (g/L) | Ara (g/L) | CA (g/L) | Pol (g/L) | Y_CA/Gly (g/g) | Y_Pol/Gly (g/g) |
|------------|----------|-----------------|---------|----------|-----------|-----------|----------|-----------|----------------|-----------------|
| 100 Flasks | a, c     | 183             | 95.4 ± 7.1 | 15.0 ± 1.9 | 13.8 ± 2.2 | 17.1 ± 2.5 | 7.3 ± 1.2 | -         | 38.2 ± 5.9    | -               |
|           |          |                 |         |          |           |           |          |           |                | 0.40            |
| Bioreactor | b        | 65              | 103.5 ± 9.9 | 19.0 ± 3.4 | 0.5 ± 0.1 | -         | -         | -         | 42.4 ± 5.5    | 0.5 ± 0.1       |
|           |          |                 |         |          |           |           |          |           |                | 0.34            |

**Table 3.**
which mostly citric acid was produced occurred in the bioreactor experiment. Specifically, a
quantity ranged between 5–9% of the total citric acid concentration determined.

In the above-mentioned fermentation point, iso-citric acid corresponded to 8.0% of the
final fermentation products between the two cultures; in the shake-flask trial (and
contrast, the biomass production rate \( r_X \) (Figure 3), fed-batch (i.e., 0.40–0.93 g/L/h [55]),
and continuous (between 0.70 and 1.30 g/L/h [60]) bioreactor operations (in the cited batch and fed-batch cultures, \( P_{CA_{\text{max}}} \) was calculated as in the present study, whereas in the continuous cultures, \( P_{CA} = CA \cdot D \), where \( CA \) was the concentration at the steady-state and \( D \) was the dilution rate imposed). It is noted
that in the shake-flask citric acid production experiments, the achieved productivities (calculated as previously stated) are, in general, much lower (0.07–0.25 g/L/h) [8,19,23,40,49,53,61,62].

The completely different metabolic profile observed between the shake-flask and the batch
bioreactor trial should be attributed to the different agitation/aeration regime between the two cultures
and more specifically to the higher level of oxygenation that occurred in the bioreactor experiment; for all of the fermentation steps in which noticeable biosynthesis of polyols occurred in the shake-flask trials, DOT values in the flasks always ranged between 20% and 50% \( \psi_{O_2} \). On the contrary, in the bioreactor experiment, with the exception of the very early fermentation step (\( t = 20 \) h) in which DOT \( \approx 20.0\% \psi_{O_2} \), rapidly afterwards, DOT level was always \( >40\% \), and at the end of the trial, when glycerol had been
consumed and the maximum \( CA \) concentration was recorded (=42.4 g/L), the DOT value was \( >90\% \psi_{O_2} \) (kinetics of DOT evolution not presented). On the other hand, this differentiation in the metabolic responses between flask and bioreactor cultures did not seem to be seriously attributed to the different pH values observed between these two types of cultures; pH is certainly a crucial culture parameter that may influence the final spectrum of products for \( Y. \text{lipolytica} \) cultures. It has been demonstrated that the very same strain cultivated under nitrogen limitation, can produce almost exclusively citric acid at medium pH values = 5.0–6.0, while metabolic shift and almost exclusive production of polyols may
Occur when cultures are performed at pH values $= 3.0–3.5$ \cite{23,46,48,63}. In the current investigation, the differences in the pH values between the flask (pH mostly found within 4.8–5.8) and the reactor (pH $= 6.0 \pm 0.1$) experiments were not so high, thus we assume that pH was not the principal factor affecting the differences in the metabolism observed between the flask and the reactor trials.

In a very similar physiological pattern with the strain LMBF Y-46, the wild-type food-derived strain ACA YC 5029, when cultivated in shake-flask trials (equally in pH $> 4.5$) produced mainly polyols (erythritol and mannitol). Minimal quantities of citric acid were synthesized $\cite{18}$. In contrast, in well-agitated and aerated bioreactors (800 rpm, aeration between 0.1 and 2.0 vvm in order to always have a cascade of DOT $\geq 10\%$, v/fb—significantly higher DOT values than 10\% v/f were finally recorded) mainly citric acid was synthesized, whereas simultaneously insignificant biosynthesis of polyols occurred $\cite{18}$. In agreement with these results and according to the literature, CA biosynthesis is strongly influenced by the DOT values in the culture medium; Ferreira et al. $\cite{64}$, performing culture optimization of the strains Y. lipolytica W29 and CBS 2073, observed a clear positive effect of the oxygen mass transfer rate (OTR) upon the biosynthesis of CA for these yeast strains during glycerol fermentation in shake-flask experiments. Likewise, in batch bioreactor experiments with the strain W29 growing on glycerol, CA production was clearly favored in trials presenting higher oxygen volumetric mass transfer coefficients ($k_{La}$); therefore, increase in the initial $k_{La}$ value from 7 h$^{-1}$ to 55 h$^{-1}$ led to a 7.8-fold increase in the final concentration of citric acid in this strain $\cite{65}$. In accordance with the mentioned results, for the acetate negative Y. lipolytica mutants Wratislavia 1.31 and Wratislavia AWG7 cultivated in batch glycerol-based bioreactor experiments, remarkable agitation into the medium (800 rpm) and cascade aeration ensuring DOT $> 40\%$ v/f throughout the culture clearly favored CA in these strains; for instance, under this aeration regime the strain Wratislavia 1.31 achieved a $CA_{max} = 92.8$ g/L, with yield $Y_{CAgly} \approx 0.63$ g/g $\cite{66}$. Concluding, it may be indicated that a variety of external factors like medium pH, salt addition and aeration may critically influence the physiological and kinetic behavior of Y. lipolytica yeast $\cite{10–12,14,15}$. However, such high discrepancies in the final spectrum of metabolites between shake-flask and aerated/agitated batch-bioreactor experiment that have been observed in the current investigation, is indeed, of notable importance.

Given that Y. lipolytica LMBF Y-46 was revealed as capable at rapidly assimilating glycerol and efficiently converting it into citric acid in bioreactor experiments, a pulsed fed-batch bioreactor trial (like the one performed by Förster et al. $\cite{57}$ and Tan et al. $\cite{67}$), in which culture conditions (viz. aeration, agitation, pH) were the same as in the previous batch, was performed (see Figure 4a–c). Significant glycerol uptake occurred, whereas noticeable DCW quantities were synthesized relatively early ($X_{max} \approx 21$ g/L) with these values being drastically depleted afterwards due to significant cell autolysis (final DCW quantity when the culture stopped $\approx 11$ g/L; Figure 4a). Despite significant cell autolysis, yeast biomass maintained its viability, since simultaneously with biomass decrease noticeable glycerol uptake occurred and remarkable quantities of CA were synthesized. When the culture stopped, a noticeable CA quantity ($= 101.1$ g/L) with a simultaneous quite high CA volumetric productivity ($=0.61$ g/L/h) were recorded (Figure 4b), with a global conversion of citric acid produced per unit of glycerol consumed ($Y_{CAgly}$) being $= 0.46$ g/g (Figure 4c). In the above-mentioned fermentation point, iso-citric acid corresponded to 7.8\% w/w of CA formed ($= 7.9$ g/L). Although at the early steps of the bioreactor culture ($t = 0–30$ h after inoculation), the assimilation of glycerol did not contribute importantly to CA formation (see Figure 4b), finally the representation of CA produced ($f$ (glycerol consumed)) and the linear regression performed for the whole fed-batch culture carried presents interest, as for most of the experimental points glycerol assimilation mostly contributed to the synthesis of CA, whereas the correlation coefficient of the regression ($r^2 = 0.97$) presented a very satisfactory value (Figure 4c). Moreover, CA production was connected with a high DOT in the medium (Figure 4a), whereas interestingly and in accordance with a number of reports indicated in the literature $\cite{40,68}$, CA production linearly increased with the time ($r^2 = 0.97$), resulting in a global $r_{CA} = 0.58$ g/L/h (Figure 4b). Metrics of citric acid production by wild-type, mutant and genetically modified Y. lipolytica strains cultivated in bioreactor experiments with several types of carbon sources
employed as substrates, in various fermentation configurations and their comparisons with the current investigation, are depicted in Table 4.

Figure 4. Kinetics of dry yeast cell mass evolution (X, g/L) and dissolved oxygen tension (% v/v) vs. time (a), glycerol evolution (Gly, g/L) and citric acid production (CA, g/L) vs. time (b), and citric acid produced (CA, g/L) vs. glycerol consumed (g/L) (c) by Yarrowia lipolytica LMBF Y-46, during growth on pure glycerol in pulsed fed-batch bioreactor experiments under nitrogen-limited conditions. Culture conditions as in Table 3. Each experimental point is the mean value of two independent determinations (SE ≤ 18%).

The maximum CA quantity obtained in the current investigation (=101.2 g/L) compares favorably with most cultures of wild-type Y. lipolytica strains, while this value is also higher compared with that obtained for various mutants of genetically engineered strains (see Table 4). Concerning the conversion yield \( Y_{CAGly} \) values achieved in the batch and fed-batch bioreactor operations of the present study (=0.41–0.46 g/g), they are slightly lower compared with the relevant studies in which other Y. lipolytica strains were grown on glycerol \([60,63,66,67]\). The growth of Y. lipolytica strains on other substrates that present higher carbon percentage and reductance degree compared with glycerol (i.e., oils or ethanol), naturally is accompanied by higher conversion yields \([56,71,77]\) compared with the present study. On the other hand, there are a number of reports in which mutants of genetically modified strains have attained greater \( C_{A_{max}} \) quantities compared with the current submission (i.e., CA quantities ranging between 102–140 g/L; \([56,57,67,72,77]\)), whereas in a scarce number of investigations, indeed tremendous \( C_{A_{max}} \) quantities (i.e., 161 and 175 g/L) have been reported \([73,74]\). On the other hand, in all the above-mentioned cases, mutant or genetically engineered strains have been implicated in these bioprocesses, whereas the strain LMBF Y-46, as previously indicated, is a food-originated wild-type new isolate that has not previously been systematically studied. Moreover, optimization of the CA production bioprocess via the above-strain as regards medium pH, agitation, oxygen uptake rate etc., can further increase the achieved CA production. Likewise, further incubation in the already realized fed-batch experiment, could further enhance the \( C_{A_{max}} \) quantity achieved by the strain LMBF Y-46 (see Figure 4b).
Table 4. Metrics of production of citric acid by yeasts belonging to the species *Yarrowia* (*Candida—Saccharomycopsis—Endomycopsis* *lipolytica*), during growth on various substrates in bioreactor experiments. Comparisons with the current study.

| Strain               | Citric Acid (g/L) | Substrate | Yield (g/g) | Bioreactor Configuration | Reference                |
|----------------------|-------------------|-----------|-------------|--------------------------|--------------------------|
| NRRL Y-7576 *        | 51.5              | Glucose   | 0.71        | Fed-batch                | Klasson et al. [69]      |
| Y-1995 *            | 78.5              | Glucose   | 0.79        | Fed-batch                | Kane and Sims [70]       |
| ATCC 20346           | 69.0              | Glucose   | 0.52        | Fed-batch                | Moresi [68]              |
| N1 *                | 120.0             | Ethanol   | 0.85        | Fed-batch                | Kamzolova et al. [71]    |
| 187I *              | 135.1             | Rapseoil  | 1.55        | Fed-batch                | Kamzolova et al. [56]    |
| Wratislavia AWG7 *   | 88.1              | Crude glycerol | 0.46  | Batch                    | Rymowicz et al. [44]     |
| Wratislavia K1 *    | 75.7              | Crude glycerol | 0.40  | Batch                    | Rymowicz et al. [44]     |
| H222-S4(p67ICL1)75* | 91.0 *            | Sucrose   | 0.53        | Fed-batch                | Förster et al. [57]      |
| H222-S4(p67ICL1)75* | 133.0 *           | Sucrose   | 0.78        | Fed-batch                | Förster et al. [57]      |
| A-101–1.22 *        | 119.1 *           | Crude glycerol | 0.64  | Fed-batch                | Rymowicz et al. [47]     |
| A-101–1.22 *        | 115.6 *           | Crude glycerol | 0.68  | Repeated batch           | Rymowicz et al. [47]     |
| A-101 *             | 69.3              | Glucose   | 0.45        | Batch                    | Rychwiska et al. [53]    |
| A-101 *             | 66.5              | Pure glycerol | 0.44  | Batch                    | Rychwiska et al. [53]    |
| A-101 *             | 66.8              | Crude glycerol | 0.43  | Batch                    | Rychwiska et al. [53]    |
| Wratislavia 1.31 *  | 76.4              | Glucose   | 0.52        | Batch                    | Rychwiska et al. [53]    |
| Wratislavia 1.31 *  | 63.9              | Pure glycerol | 0.40  | Batch                    | Rychwiska et al. [53]    |
| Wratislavia 1.31 *  | 82.0              | Crude glycerol | 0.53  | Batch                    | Rychwiska et al. [53]    |
| Wratislavia 1.31 *  | 126.0 *           | Crude glycerol | 0.63  | Fed-batch                | Rychwiska et al. [72]    |
| Wratislavia AWG7 *  | 157.5 *           | Crude glycerol | 0.58  | Fed-batch                | Rychwiska et al. [72]    |
| Wratislavia AWG7 *  | 160.5 *           | Crude glycerol | 0.81  | Repeated batch           | Rychwiska and Rymowicz [73] |
| Wratislavia 1.31 *  | 124.9 *           | Crude glycerol | 0.59  | Fed-batch                | Rychwiska and Rymowicz [73] |
| N15 *              | 98.0              | Pure glycerol | 0.70  | Fed-batch                | Kamzolova et al. [61]    |
| NG40/UV7 *          | 175.0             | Rapseoil  | 1.50        | Fed-batch                | Kamzolova et al. [74]    |
| Wratislavia AWG7 *  | 63.3              | Pure glycerol | 0.67  | Continuous               | Rychwiska et al. [60]    |
| Wratislavia 1.31 *  | 92.8              | Pure glycerol | 0.63  | Batch                    | Rychwiska et al. [66]    |
| SWJ-1b              | 84.0              | Inulin    | 0.89        | Batch                    | Liu et al. [75]          |
| NG40/UV7 *          | 115.0             | Pure glycerol | 0.64  | Fed-batch                | Morgunov et al. [59]     |
| NG40/UV7 *          | 112.0             | Crude glycerol | 0.90  | Fed-batch                | Morgunov et al. [59]     |
| Wratislavia AWG7 *  | 85.7              | Crude glycerol | 0.52  | Batch                    | Tomaszewskas et al. [48] |
| Wratislavia K1 *    | 65.0              | Crude glycerol | 0.43  | Batch                    | Tomaszewskas et al. [48] |
| SWJ-1b              | 101.6             | Glucose   | 0.89        | Fed-batch                | Tan et al. [67]          |
| CBS 6114 *          | +55 *             | Pure glycerol | n.i. | Batch                    | Egermeier et al. [36]    |
| H222 *              | +50 *             | Pure glycerol | n.i. | Batch                    | Egermeier et al. [36]    |
| DSM 1345 *          | +52 &             | Pure glycerol | n.i. | Batch                    | Egermeier et al. [36]    |
| VKM Y-2373 *        | 80–85             | Glucose   | 0.70–0.75   | Batch                    | Kamzolova and Morgunov [76] |
| ACA YC 5029 *       | 39.0              | Crude glycerol | 0.42  | Batch                    | Papanikolaou et al. [19] |
| NG40/UV5 *          | 140.0             | Rapseoil  | 1.50        | Fed-batch                | Morgunov et al. [77]     |
| NG40/UV5 *          | 108.8             | Glucose   | 0.80        | Fed-batch                | Morgunov et al. [77]     |
| NG40/UV5 *          | 87.0              | Crude glycerol | 0.64  | Fed-batch                | Morgunov et al. [77]     |
| K57 *              | 72.1              | Glucose   | 0.77        | Batch                    | Carsana et al. [62]      |
| AJD pADUTGut1/2 *   | 75.8 &              | Crude glycerol | 0.51  | Batch                    | Rzechonka et al. [63]    |
| LMBF Y-46 *         | 42.4 &             | Pure glycerol | 0.41  | Batch                    | Present study            |
| LMBF Y-46 *         | 101.3 &            | Pure glycerol | 0.46  | Fed-batch                | Present study            |

*: Mutant or genetically modified *Yarrowia lipolytica*; 5: Wild-type *Yarrowia lipolytica*; 6: Total citric acid (citric + iso-citric acid, in which iso-citric acid ranged between 5–9% of total citrate); 8: Total citric acid (citric + iso-citric acid, in which iso-citric acid is >50% of total citrate); n.i.: Not indicated in the manuscript.

3.4. Modeling the Production of Polyols by *Yarrowia lipolytica* LMBF Y-46 Growing on Glycerol in Shake-Flask Experiments

The production of polyols by *Y. lipolytica* LMBF Y-46 in shake-flask experiments in which increasing Gly concentrations were adjusted into the medium while the initial nitrogen concentration remained constant (see kinetic results in Table 1) was numerically modeled. Modeling *Y. lipolytica* growth on glycerol and the accompanied total polyols production, was based on the equations and the numerical procedure used in describing the lipid production by *Cryptococcus curatus*, grown on commercial-type xylose [78], originated from the modeling of citric acid production by another *Y. lipolytica* strain growing batch-wise in glucose-enriched olive mill wastewater-based media [79]. The governing equations and the associated kinetic parameters related to yeast DCW production, glycerol consumption and polyols accumulation into the medium for the experiments with the three higher glycerol concentrations employed, that is, for 100, 120 and 150 g/L nominal concentrations, are presented on Table 5. In the
present work, a Verhulst-type equation [79] in which the glycerol decreasing ratio, \( \text{Gly/Gly}_0 \), according to the approach proposed by Aggelis and colleagues [80–82], was used in order to quantitatively describe the specific growth rate (\( \mu \)). Note that the treatment of biomass concentration at the beginning of the experiments (i.e., \( X_0 > 0 \)) as an adjustable parameter (Table 5) highly improved the quality of the fitting. Comparisons between experimental data and theoretical concentration values are depicted in Figure 5a–c, demonstrating a very satisfactory fitting for all runs performed.

**Table 5.** Governing equations and associated kinetic parameters related to biomass growth, glycerol consumption and polyols production during *Yarrowia lipolytica* culture on different glycerol concentrations.

| Equation | Number of Data Points | Glycerol Concentration (g/L) |
|----------|------------------------|------------------------------|
| \( \frac{dX}{dt} = \mu X \) | 1 | 100 | 120 | 150 |
| | | 11 | 12 | 13 |
| \( \mu = \mu_{\text{max}} \left( 1 - \frac{X}{X_{\text{max}}} \right) \) | 2 | \( \frac{dX}{dt} \) = \( \mu_{\text{max}} (\text{gX/L}) \) | \( X_0 (\text{gX/L}) \) | SSE | \( \mu_{\text{max}} \) (h\(^{-1}\)) | \( X_{\text{max}} \) (gX/L) |
| | | | | | 2.26 | 2.22 | 2.23 |
| | | | | | 13.09 | 9.48 | 8.03 |
| | | | | | 0.942 | 0.937 | 0.935 |
| \( \frac{d\text{Pol}}{dt} = q_{\text{Pol}} X \) | 3 | \( \frac{dX}{dt} \) = \( q_{\text{Pol}} (g\text{Pol}/(gXh)) \) | \( X_0 (g\text{Pol}/L) \) | SSE | \( q_{\text{Pol}} \) (gPol/(gXh)) |
| | | | | | | | | | 40.14 | 48.86 | 162.87 |
| \( \frac{d\text{Pol}}{dt} = q_{\text{Pol}} \left( 1 - \frac{\text{Pol}}{\text{Pol}_{\text{max}}} \right) \) | 4 | \( \frac{dX}{dt} \) = \( q_{\text{Pol}} (g\text{Pol}/(gXh)) \) | \( X_{\text{max}} (g\text{Pol}/L) \) | SSE | \( q_{\text{Pol}} \) (gPol/(gXh)) |
| | | | | | | | | | 127.02 | 59.43 | 68.33 |
| | | | | | | | | | 0.951 | 0.974 | 0.981 |
| \( \frac{-d\text{Gly}}{dt} = \mu \frac{1}{Y_{\text{X/Gly}}} X + q_{\text{Pol}} \frac{1}{Y_{\text{X/Gly}}/X} X \) | 5 | \( \frac{dX}{dt} \) = \( \mu (g\text{Gly}/g\text{X}) \) | \( X_{\text{max}} (g\text{Gly}/L) \) | SSE | \( Y_{\text{X/Gly}} (g\text{X}/g\text{Gly}) \) | \( Y_{\text{Pol/Gly}} (g\text{Pol}/g\text{Gly}) \) |
| | | | | | | | | | 0.2160 | 0.2749 | 0.1210 |
| | | | | | | | | | 0.9997 | 0.6575 | 1.0586 |
| | | | | | | | | | 445.2 | 187.1 | 1027.8 |
| | | | | | | | | | 0.954 | 0.984 | 0.956 |

SSE: Sum of squared errors.

The kinetic parameter values are comparable to the theoretical and experimental values recorded in the literature; specifically, the optimized parameter value for \( \mu_{\text{max}} \) in the current investigation (>0.03 h\(^{-1}\)), presented slightly lower values than that reported for other optimized \( \mu_{\text{max}} \) values for various *Y. lipolytica* strains cultivated batch-wise and producing a plethora of secondary metabolites (optimized \( \mu_{\text{max}} \) values ranged between 0.16 and 0.44 h\(^{-1}\)) [68,69,79,82]. Optimized \( q_{\text{Pol}} \) values are found in the same order of magnitude with experimental values reported for other *Y. lipolytica* strains cultured on glycerol in batch experiments (0.002–0.04 gPol/gXh; see: Tomaszewska et al. [29]; [48]). The same can be stressed concerning the optimized \( Y_{\text{X/Gly}} \) values reported in the current investigation (Table 5), which presented remarkable similarities with optimized values achieved from application of several types of numerical modes for other *Y. lipolytica* strains cultured on glycerol (=0.29–0.30 g/g) [82]. Equally, the above-mentioned optimized \( Y_{\text{X/Gly}} \) values (see Table 5) present important similarities with experimentally achieved \( Y_{\text{X/Gly}} \) values for various *Y. lipolytica* strains cultured on crude glycerol in batch experiments (0.11–0.30 g/g) [18,22,29,48].

Optimized \( Y_{\text{Pol/Gly}} \) values (ranging between 0.66 and 1.06 gPol/gGly—see Table 5) were somehow overestimated when compared to the experimental data (see Table 1 and Figure 2). The biochemical patterns of polyols formation from glycerol fermentation, have not yet been completely elucidated [10,13], therefore, it is not known which the maximum theoretical conversion yield of glycerol into mannitol/erythritol and arabitol is. On the other hand, as previously stressed, with the exception of a scarce number of reports [50,52,54], experimental values of the \( Y_{\text{Pol/Gly}} \) yield have rarely been reported to be >0.60 g/g, therefore, the optimized \( Y_{\text{Pol/Gly}} \) parameter values seem to be overestimated. Finally, the rather high \( \text{Pol}_{\text{max}} \) value of 162.87 g/L obtained for the data of Gly0 \( \approx 150 \) g/L, is attributed to the fact that glycerol was not totally consumed and polyols formation was still rising for the duration of the experiment (Figure 5) making predictions of \( \text{Pol}_{\text{max}} \) ambiguous.
Figure 5. Experimental data and fitted values showing the evolution of biomass (X), glycerol (Gly) and total polyols (Pol) concentration during growth of *Yarrowia lipolytica* on glycerol at: ≈100 g/L (a); ≈120 g/L (b); ≈150 g/L (c), initial substrate concentrations in shake-flask experiments under nitrogen-limited conditions. Culture conditions as in Table 3. Each experimental point is the mean value of two independent determinations (SE ≤ 18%).

The proposed model was not proved to be adequate to describe the data for the experiment with the 70 g/L initial glycerol concentrations. A close look at the experimental data (Figure 6) reveals a continuous increase in biomass, above 20 g/L values, even after glycerol depletion, at the expense of polyols. Such a behavior requires a different, rather unique modeling approach that was beyond the scope of the present work.

3.5. Cellular Lipids of *Yarrowia lipolytica* LMBF Y-46

Total cellular lipids were extracted and quantified for both the shake-flask and the bioreactor experiments, and the kinetics of the evolution of lipids in DCW are depicted in Figure 7a,b. In all cases, and as it had previously been demonstrated (Figure 1b), the microorganism under investigation did not produce remarkable quantities of total lipids in yeast biomass, irrespective of the culture configuration (flask or bioreactor experiments). In fact, as it has been demonstrated in several previous studies dealing with wild-type and genetically engineered *Y. lipolytica* strains cultured in batch experiments in both shake-flask and bioreactor trials, the oleaginous character of this species is strain dependent;
in fact, in culture conditions of nitrogen starvation, in which carbon is found in the form of glucose, glycerol or similarly metabolized compounds (that favors the so-called “de novo” cellular lipid accumulation process; [5,83]), excess carbon is directed either towards the synthesis and secretion of low-molecular weight compounds (i.e., citric acid, polyols) or it is channeled towards the accumulation of storage lipids [5,9,83]. In Y. lipolytica, the above-indicated regulation is difficult, and it has been demonstrated that it depends on the availability of extra-cellular nitrogen (for example, Ochoa-Estopier and Guillouet [84] have demonstrated that a C/N ratio of 11.7 yields lipid accumulation, while a ratio of 47.6 favors CA production using D-stat continuous cultivation techniques), the utilization of double- or multiple-limited media (for example in multiple-limited media and during growth on glucose or glycerol batch-wise, significant lipid quantities have been produced by Y. lipolytica yeasts that in simple nitrogen-limited ones produced mostly CA [8,23,85]), etc. Adaptive laboratory evolution strategies or genetic engineering approaches have resulted in the formation of Y. lipolytica strains that are capable of producing high amounts of cellular lipids and limited amounts of citric acid [43,86–88].

![Graph](image)

**Figure 6.** Kinetics of dry yeast cell mass production (X, g/L), glycerol assimilation (Gly, g/L) and total polyols production (Pol, g/L) during Yarrowia lipolytica growth on initial glycerol adjusted to ≈70 g/L in shake-flask experiments under nitrogen-limited conditions. Culture conditions as in Table 3. Each experimental point is the mean value of two independent determinations (SE ≤ 18%).

In a number of cases, Y. lipolytica strains cultured in nitrogen-limited glucose- or glycerol-based media, produced sequentially lipid and extra-cellular low molecular-weight metabolites (citric acid and polyols); in the first steps of the culture, lipid accumulation was somehow induced, and then lipid in DCW values were depleted with time, even though significant substrate quantities remained unconsumed into the medium. At that period, secretion of citric acid or polyols, in high quantities, was carried out [18,22,38–40,85]. Exactly the same feature as regards lipid production and secretion of polyols was observed for the strain LMBF Y-46 in shake-flask experiments; somehow elevated lipid DCW values (i.e., 13–15% w/w) were recorded at the earlier growth phases of the strain cultured on glycerol in flasks, irrespective of the Gly0 concentration set into the medium, and these values were significantly reduced when the culture proceeded (Figure 7a). The onset of secretion of polyols into the medium was carried out at c. t = 50 ± 3 h, occurring simultaneously with lipid DCW value depletion. In contrast, different kinetic profiles were recorded for the bioreactor experiment (Figure 7b). Compared with the equivalent flask experiment (Gly0 ≈ 100 g/L), and despite the higher oxygenation that occurred in the fermentation vessel, the microorganism produced noticeably lower lipid quantities in the bioreactor experiment (see Figure 7b, bioreactor experiment, fermentation time t = 0–66 h, in which lipid DCW values were ≈5% w/w). After the two consecutive pulses with concentrated glycerol into the medium, lipid in yeast mass quantities somehow increased (to values ≈ 10–12% w/w). This total lipid quantity increase occurred simultaneously with cell autolysis and substantial decrease in dry biomass concentration (Figure 4a), indicating once more, that despite cell autolysis, Y. lipolytica biomass was quite active at the late fermentation steps in the bioreactor culture.
with the flask cultures. Moreover, the first experiment point in the bioreactor trial that somehow presented
(i.e., arachidic—C20:0, behenic—C22:0 and lignoceric—C24:0) aliphatic chain saturated FAs have also
These concentrations decreased as fermentations proceeded [23,38,39,49]. This could be attributed
to the lower DOT values in these cultures at the early growth steps of the balanced growth phase of
Y. lipolytica. Finally, in accordance with the FA composition analysis results presented in the
current investigation, strains of the oleaginous yeast

Figure 7. Kinetics of intra-cellular lipids produced per unit of dry cell weight (%) w/w of Yarrowia lipolytica LMBF Y-46 during growth on pure glycerol, at various initial glycerol concentrations (Gly0) imposed under nitrogen-limited conditions in shake-flask experiments (a) and shake-flask vs. pulsed fed-batch bioreactor experiments (in the latter case, for the shake-flask experiment, Gly0 concentration was ≈100 g/L). (b) Culture conditions as in Tables 1 and 3. Each point is the mean value of two independent determinations (SE ≤ 18%).

Total cellular lipids were converted into their corresponding FAMEs, and cellular fatty acid (FA) composition analysis was performed for the late growth phase in the case of the shake-flask cultures, and throughout the culture, for the fed-batch bioreactor experiment (Table 6). In general, FA composition of total lipids of the strain LMBF Y-46 presented important similarities with those reported for various other Y. lipolytica strains cultivated in batch flask and bioreactor cultures, during growth on glycerol or similarly metabolized compounds, under nitrogen limitation [8,16,18,19,21,22,38–40,49,53]. Specifically, the yeast lipids were mainly composed of the FAs oleic (C18:1), palmitic (C16:0), palmitoleic (C16:1), stearic (C18:0) and linoleic (C18:2). No quantities of other poly-unsaturated FAs (i.e., α-linolenic acid—C18:3) were detected inside the yeast lipids in accordance with the literature [5,9,83]. Small but not negligible quantities of lower (i.e., capric—C10:0, lauric—C12:0 and myristic—C14:0) or higher (i.e., arachidic—C20:0, behenic—C22:0 and lignoceric—C24:0) aliphatic chain saturated FAs have also been detected in the yeast lipids (Table 6). An interesting result that was observed through the cellular lipid analysis, referred to the fact that cellular lipids were somehow more saturated (mostly more rich in the FAs C16:0 and C18:0) in the shake-flask experiments compared with those obtained in the bioreactor cultures. In the latter case, cellular lipids were notably richer in the FAs C18:1 and C18:2. This could reflect the fact that bioreactor cultures were conducted under higher aeration compared with the flasks. Moreover, the first experimental point in the bioreactor trial that somehow presented higher concentrations of the cellular palmitic and stearic acids (t = 20.5 h) compared with all other experimental points in this bioreactor experiment (see Table 6), could be attributed to the fact that at this very fermentation point, the DOT value (in %, v/v) was the lowest one in the relevant bioreactor trial (at t = 20.5 h, DOT value into the reactor vessel was ≈20% v/v, and thereafter it increased; see Figure 4a). Similarly with this result, higher concentrations of the cellular FAs C16:0 and C18:0 have been noted in the beginning of other batch cultures performed by several Y. lipolytica strains on glucose or glycerol. These concentrations decreased as fermentations proceeded [23,38,39,49]. This could be attributed to the lower DOT values in these cultures at the early growth steps of the balanced growth phase of

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simple nitrogen-limited ones produced mostly CA [8,23,85], etc. Adaptive laboratory evolution

had been cultured in air-lift bioreactor

mediated compounds, under nitrogen limitation [8,16,18,21,22,38,40,49,53]. Specifically, the yeast lipids were mainly composed of the FAs oleic (C18:1), palmitic (C16:0), palmitoleic (C16:1), stearic (C18:0) and linoleic (C18:2). No quantities of other poly-unsaturated FAs (i.e., α-linolenic acid—C18:3) were detected inside the yeast lipids in accordance with the literature [5,9,83]. Small but not negligible quantities of lower (i.e., capric—C10:0, lauric—C12:0 and myristic—C14:0) or higher (i.e., arachidic—C20:0, behenic—C22:0 and lignoceric—C24:0) aliphatic chain saturated FAs have also been detected in the yeast lipids (Table 6). An interesting result that was observed through the cellular lipid analysis, referred to the fact that cellular lipids were somehow more saturated (mostly more rich in the FAs C16:0 and C18:0) in the shake-flask experiments compared with those obtained in the bioreactor cultures. In the latter case, cellular lipids were notably richer in the FAs C18:1 and C18:2. This could reflect the fact that bioreactor cultures were conducted under higher aeration compared with the flasks. Moreover, the first experimental point in the bioreactor trial that somehow presented higher concentrations of the cellular palmitic and stearic acids (t = 20.5 h) compared with all other experimental points in this bioreactor experiment (see Table 6), could be attributed to the fact that at this very fermentation point, the DOT value (in %, v/v) was the lowest one in the relevant bioreactor trial (at t = 20.5 h, DOT value into the reactor vessel was ≈20% v/v, and thereafter it increased; see Figure 4a). Similarly with this result, higher concentrations of the cellular FAs C16:0 and C18:0 have been noted in the beginning of other batch cultures performed by several Y. lipolytica strains on glucose or glycerol. These concentrations decreased as fermentations proceeded [23,38,39,49]. This could be attributed to the lower DOT values in these cultures at the early growth steps of the balanced growth phase of Y. lipolytica yeast. Finally, in accordance with the FA composition analysis results presented in the current investigation, strains of the oleaginous yeast A. curvatum had been cultured in air-lift bioreactor experiments, where somehow low DOT (and, thus, low OTR) values had been imposed, and while DCW production and cellular lipid biosynthesis were not seriously curtailed, storage lipids were noticeably enriched in the saturated FAs compared with the experiments presenting higher OTRs [89].
Table 6. Fatty acid composition of the cellular lipids produced by *Yarrowia lipolytica* LMBF Y-46 cultivated on glycerol in shake-flask, batch and fed-batch bioreactor experiments. Each experimental point presented is the mean value of two independent determinations. Culture conditions as in Tables 1 and 3.

| Culture Type/Time | Shake-Flasks | Fatty Acid Composition of Yeast Lipids (% w/w) |
|-------------------|--------------|-----------------------------------------------|
|                   | ≤C12:0 *     | C14:0 | C16:0 | C16:1 | C18:0 | C18:1 | C18:2 | C20:0 | ≥C22:0 $^\dagger$ | Other $^x$ |
| Gly0 ≈ 70 g/L, t ≈ 180 h | 2.5 ± 0.3 | 2.3 ± 0.4 | 21.1 ± 1.7 | 5.1 ± 0.6 | 15.6 ± 2.0 | 30.2 ± 3.5 | 11.5 ± 1.3 | 0.8 ± 0.1 | 2.2 ± 0.3 | 8.7 ± 0.9 |
| Gly0 ≈ 100 g/L, t ≈ 160 h | 0.8 ± 0.1 | 0.7 ± 0.1 | 22.2 ± 3.5 | 5.5 ± 0.4 | 18.5 ± 2.4 | 32.3 ± 4.0 | 12.2 ± 2.0 | 1.2 ± 0.2 | 2.5 ± 0.4 | 4.1 ± 0.3 |
| Gly0 ≈ 100 g/L, t ≈ 180 h | 1.3 ± 0.2 | 0.9 ± 0.2 | 17.5 ± 2.5 | 3.6 ± 0.5 | 25.3 ± 2.4 | 28.4 ± 4.1 | 8.8 ± 1.0 | 5.9 ± 1.1 | 3.0 ± 0.4 | 5.3 ± 0.6 |
| Gly0 ≈ 120 g/L, t ≈ 160 h | 1.8 ± 0.3 | 1.0 ± 0.1 | 19.9 ± 2.0 | 4.6 ± 0.7 | 18.9 ± 3.0 | 30.8 ± 5.0 | 9.0 ± 1.5 | 4.2 ± 0.9 | 2.0 ± 0.3 | 7.8 ± 1.4 |
| Gly0 ≈ 120 g/L, t ≈ 190 h | 0.2 ± 0.1 | 0.4 ± 0.1 | 21.7 ± 2.4 | 6.6 ± 0.8 | 15.9 ± 2.5 | 36.2 ± 5.2 | 12.8 ± 2.0 | 1.1 ± 0.1 | 2.9 ± 0.5 | 2.2 ± 0.4 |
| Gly0 ≈ 150 g/L, t ≈ 190 h | 3.9 ± 0.2 | 2.4 ± 0.2 | 19.5 ± 1.9 | 5.1 ± 0.4 | 18.0 ± 3.2 | 30.0 ± 5.0 | 10.7 ± 1.8 | 2.0 ± 0.2 | 2.0 ± 0.4 | 6.4 ± 0.9 |

| Culture Type/Time | Bioreactor | Fatty Acid Composition of Yeast Lipids (% w/w) |
|-------------------|------------|-----------------------------------------------|
|                   | ≤C12:0 *   | C14:0 | C16:0 | C16:1 | C18:0 | C18:1 | C18:2 | C20:0 | ≥C22:0 $^\dagger$ | Other $^x$ |
| t = 20.5 h        | 1.9 ± 0.4  | 3.0 ± 0.4 | 21.1 ± 2.5 | 7.7 ± 0.8 | 17.0 ± 1.8 | 30.5 ± 3.9 | 17.7 ± 2.6 | - | - | 1.1 ± 0.3 |
| t = 41.5 h        | -          | 1.2 ± 0.2 | 18.4 ± 2.0 | 6.8 ± 0.9 | 12.0 ± 2.0 | 37.3 ± 4.0 | 16.2 ± 3.0 | - | 3.0 ± 0.4 | 5.1 ± 0.3 |
| t = 44.5 h        | 1.5 ± 0.4  | 0.7 ± 0.3 | 16.5 ± 2.0 | 8.7 ± 1.0 | 7.7 ± 1.0 | 39.0 ± 5.1 | 15.4 ± 2.9 | 0.5 ± 0.1 | 2.6 ± 0.4 | 7.4 ± 1.2 |
| t = 66.0 h        | 2.5 ± 0.2  | 1.2 ± 0.2 | 16.4 ± 2.1 | 7.3 ± 0.9 | 9.2 ± 1.4 | 36.2 ± 6.1 | 14.9 ± 3.0 | 0.5 ± 0.2 | 2.2 ± 0.4 | 9.9 ± 1.5 |
| t = 67.0 h        | 1.7 ± 0.1  | 0.8 ± 0.1 | 16.3 ± 2.2 | 8.1 ± 1.2 | 8.1 ± 1.3 | 39.7 ± 5.8 | 16.7 ± 3.1 | 0.4 ± 0.1 | 2.8 ± 0.3 | 5.4 ± 0.9 |
| t = 89.0 h        | 0.7 ± 0.1  | 0.6 ± 0.1 | 16.7 ± 2.3 | 9.0 ± 1.4 | 8.1 ± 1.6 | 40.3 ± 6.0 | 16.1 ± 3.2 | 0.4 ± 0.1 | 2.7 ± 0.3 | 5.4 ± 1.0 |
| t = 91.0 h        | 2.2 ± 0.3  | 1.1 ± 0.2 | 17.0 ± 1.9 | 8.4 ± 1.7 | 8.4 ± 1.5 | 37.4 ± 5.1 | 16.2 ± 3.0 | 0.3 ± 0.1 | 1.5 ± 0.2 | 7.5 ± 1.3 |
| t = 97.0 h        | 0.9 ± 0.1  | 0.7 ± 0.2 | 16.3 ± 2.5 | 8.5 ± 1.8 | 8.0 ± 1.6 | 40.2 ± 6.1 | 16.1 ± 2.7 | 0.5 ± 0.2 | 2.8 ± 0.3 | 6.0 ± 1.0 |
| t = 122.5 h       | 0.9 ± 0.1  | 0.5 ± 0.2 | 16.1 ± 2.0 | 9.5 ± 1.8 | 7.0 ± 1.0 | 41.1 ± 4.9 | 16.9 ± 2.0 | 0.7 ± 0.2 | 2.5 ± 0.3 | 4.8 ± 1.2 |
| t = 164.0 h       | 1.1 ± 0.2  | 0.8 ± 0.4 | 15.0 ± 1.9 | 9.5 ± 1.5 | 6.0 ± 1.0 | 43.0 ± 4.4 | 17.9 ± 3.1 | 0.8 ± 0.1 | 1.9 ± 0.3 | 4.0 ± 0.8 |

$^*$: C12:0 and C10:0; $^\dagger$: C22:0 and C24:0; $^x$: C8:0, C11:0, C20:1, C17:0, C15:0.
4. Concluding Remarks

Y. lipolytica LMBF Y-46 significantly resisted high Gly0 concentrations in the medium (up to \( \cong 150 \text{ g/L} \)), demonstrating simultaneously remarkable DCW synthesis and glycerol uptake during shake-flask cultures. Although medium pH remained \( \cong 4.5 \), cultures were accompanied by an interesting production of polyols (Pol \( \cong 52 \text{ g/L} \)). In aerated bioreactor experiments, a metabolic shift towards citric acid production was observed, while simultaneously insignificant polyol quantities were synthesized. A maximum total citric acid quantity of \( \cong 102 \text{ g/L} \), one of the highest in the literature for wild-type Y. lipolytica strains, was recorded in a successful fed-batch experiment with glycerol employed as the substrate. The metabolic transition from the biosynthesis of polyols (shake-flask trials) towards citric acid (bioreactor trials) was attributed to the higher oxygen saturation in the medium that occurred in the latter case. In the bioreactor experiments, cellular lipids containing higher concentrations of unsaturated FAs were noted. The current study therefore, provided strong evidence that a new wild-type food-derived Y. lipolytica strain can successfully be implicated as a microbial cell factory in order to convert glycerol, an abundant low-cost material, into added-value metabolic compounds of importance for the food and chemical industries.

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