## **RESEARCH PAPER**

# Improved 1,3-propanediol Production by *Escherichia coli* from Glycerol Due to Co-expression of Glycerol Dehydratase Reactivation Factors and Succinate Addition

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Abstract Escherichia coli was engineered to produce 1,3-propanediol (1,3-PDO) from glycerol, an inexpensive carbon source. This was done by introducing a synthetic pathway consisting of glycerol dehydratase, glycerol dehydratase reactivation factor, and 1,3-propanediol oxidoreductase isoenzyme. The JM-30BY15AB harboring pQE30/dhaB123, yqhD and pQE15A/gdrA, gdrB produced 1,3-PDO (7.2 g/L) from glycerol, at a level higher than that produced by JM-30BY harboring pQE30/dhaB123, yahD (4.1 g/L). When 10 mM succinate was added to the medium, the titer of 1,3-PDO and the glycerol consumption increased to 9.9 and 23.84 g/L, respectively. In addition, the ratio of NADH to NAD<sup>+</sup> increased by 43%. The titer of 1,3-PDO and glycerol consumption were 145.6 and 86.6% higher, respectively, than those from the control which harbors one vector system without gdrAB and did not include succinate addition. Under fed-batch fermentation conditions, the titer of 1,3-PDO and its conversion yield from glycerol were 13.11 g/L and 0.49 g/g, respectively. This dual-vector system may be a useful approach for the co-expression of recombinant proteins. Further, succinate addition is a promising route for the biotechnological production of NADH-dependent microbial metabolites.

**Keywords:** dual-vector system, engineered *E. coli*, glycerol, *Klebsiella pneumoniae* AJ4, 1,3-propanediol, succinate

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### 1. Introduction

In recent years, global markets for the biodiesel, oleochemical, bioethanol, and soap industries have expanded. The production of glycerol is increasing as a major byproduct of these industries, and this trend has generated a huge surplus of glycerol, which is considered an inexpensive and abundant feedstock for bioprocess industries. Glycerol is a highly reduced carbon source; hence, as a feedstock, it can be converted to various high-value reduced chemicals such as 1,3-propanediol (1,3-PDO) and generate higher product yields than those obtained using glucose [1,2].

1,3-PDO is a chemical with two hydroxyl groups that has a variety of applications in both medicine and industry, especially in the production of biodegradable and biocompatible synthetic polymers such as polytrimethylene terephthalate (PTT) [3,4]. Monomer generation is currently based on chemical production, but much attention has been paid to biological methods that can use renewable feedstock and are friendlier to the environment [5-7]. Many microorganisms such as *Klebsiella*, *Clostridium*, and *Citrobacter* have been studied for their ability to convert glycerol into 1,3-PDO [2,8-12]. However, industrial applications of these microorganisms are restricted because they are either opportunistic pathogens or strict anaerobes.

*E. coli* is generally regarded as a safe, well-studied microorganism for heterologous expression and is easy to culture. However, the bacteria cannot naturally produce 1,3-PDO by glycerol consumption. We constructed recombinant *E. coli* by introducing a synthetic pathway consisting of glycerol dehydratase complex (*dha*B123) and glycerol dehydratase reactivation factors (*gdr*AB) from *Klebsiella pneumoniae* AJ4 [9] and 1,3-propanediol oxidoreductase isoenzyme (*yqh*D) from *E. coli*. The production of 1,3-PDO by

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**Fig. 1.** 1,3-Propanediol production pathway from glycerol in recombinant *E. coli* over-expressing glycerol dehydratase (*dha*B123), glycerol dehydratase reactivation factors (*gdr*AB), and 1,3-propanediol oxidoreductase (*yqh*D). *dha*B<sup>\*</sup> represents the inactivated glycerol dehydratase, which can be reactivated by *gdr*AB in the presence of ATP, coenzyme B<sub>12</sub>, and Mg<sup>2+</sup>. GK, glycerol kinase; G3PDH, glycerol-3-phosphate dehydrogenase; GDH, glycerol dehydrogenase; DHAK, dihydroxyacetone kinase; G3P, glycerol-3-phosphate; DHA, dihydroxyacetone; DHAP, dihydroxyacetonephosphate.

recombinant *E. coli* under aerobic conditions eliminates the large volume of nitrogen gas consumption. This reduces the costs of general 1,3-PDO production. In 1,3-PDO production, *dha*B123 converts glycerol to 3-hydroxypropionaldehyde (3-HPA) in the presence of coenzyme  $B_{12}$ , and *yqh*D catalyzes the conversion of 3-HPA to 1,3-PDO (Fig. 1) [13-16].

Glycerol dehydratase undergoes mechanism-based inactivation by its physiological substrate, glycerol [17,18]. Glycerol dehydratase can be reactivated by a glycerol dehydratase reactivation factors in the presence of ATP, coenzyme B<sub>12</sub>, and Mg<sup>2+</sup> [18,19]. To investigate the effect of *gdr*AB during 1,3-PDO fermentation, we tested three commonly used laboratory *E. coli* strains for their ability to convert glycerol into 1,3-PDO when transformed with either one plasmid (pQE30/*dha*B123, *yqh*D) or two plasmids (pQE30/*dha*B123, *yqh*D and pQE15A/*gdr*A, *gdr*B). Then we identified a suitable strain as a host for the production of 1,3-PDO from glycerol. Finally, the flask-culture experiments of the selected *E. coli* strain JM109 were performed to study its efficiency when tricarboxylic acid (TCA) cycle intermediates were added. Bioreactor fed-batch experiments were carried out to evaluate JM109's potential for increased production of 1,3-PDO from glycerol.

# 2. Materials and Methods

#### 2.1. Bacterial strains and plasmid constructs

All bacterial strains, including engineered strains and plasmids, and primers used in this study are listed in Table 1. *E. coli* XL1-blue, *E. coli* JM109, and *E. coli* W3110 were used as host strains. The plasmids pQE30 and pQE15A were used as expression plasmids. The plasmid pQE15A was constructed by overlap-extension PCR using primers P30 F-P30 R and P15A F-P15A R. A procedure used to replace the ampicillin-resistance and ColE1-origin genes of

| Strains                  | Genotype   | Source                |
|--------------------------|--|-----------------------|
| E.coli XL-blue           | endA1 gyrA96(nal <sup>R</sup> ) thi-1 recA1 relA1 lac glnV44 F'<br>[::Tn10 proAB <sup>+</sup> lacI <sup>q</sup> $\Delta$ (lacZ)M15] hsdR17(r <sub>K</sub> <sup>-</sup> m <sub>K</sub> <sup>+</sup> )                                 | Stratagene            |
| E.coli JM109             | endA1 glnV44 thi-1 relA1 gyrA96 recA1 mcrB <sup>+</sup> $\Delta$ (lac-proAB)<br>e14-[F <sup>+</sup> traD36 proAB <sup>+</sup> lacI <sup>4</sup> $\Delta$ (lacZ)M15] hsdR17(r <sub>K</sub> <sup>-</sup> m <sub>K</sub> <sup>+</sup> ) | NEB                   |
| E.coli W3110             | $F^{-}\lambda^{-}$ rph-1 INV(rrnD, rrnE)   | ATCC27325             |
| Klebsiella pnumoniae AJ4 | Wild type, isolation from soil   | Laboratory collection |
| XL-30BY                  | Recombinant E. coli XL-blue harboring pQE30/dhaB123, yqhD  | This study            |
| XL-30BY15AB              | Recombinant <i>E. coli</i> XL-blue harboring pQE30/ <i>dha</i> B123, <i>yqh</i> D and pQE15A/ <i>gdr</i> A, <i>gdr</i> B   | This study            |
| JM-30BY                  | Recombinant E. coli JM109 harboring pQE30/dhaB123, yqhD  | This study            |
| JM-30BY15AB              | Recombinant <i>E. coli</i> JM109 harboring pQE30/ <i>dha</i> B123, <i>yqh</i> D and pQE15A/ <i>gdr</i> A, <i>gdr</i> B   | This study            |
| W-30BY                   | Recombinant E. coli W3110 harboring pQE30/dhaB123, yqhD  | This study            |
| W-30BY15AB               | Recombinant <i>E. coli</i> W3110 harboring pQE30/ <i>dha</i> B123, <i>yqh</i> D and pQE15A/ <i>gdr</i> A, <i>gdr</i> B   | This study            |
| Plasmids                 | Description  | Source                |
| pQE30                    | ColE1 <i>ori</i> , P <sub>T5</sub> , Amp <sup>R</sup>  | Qiagen                |
| pACYCDuet-1              | P15A <i>ori</i> , P <sub>T7</sub> , Cm <sup>R</sup>  | Novagen               |
| pQE15A                   | pQE30 derivative replaced Amp <sup>R</sup> and ColE1 <i>ori</i> genes with Cm <sup>R</sup> and P15A <i>ori</i> genes of pACYCDuet-1  | This study            |
| р30-ВҮ                   | pQE30 derivative containing <i>dha</i> B123 and <i>yqh</i> D genes   | This study            |
| p15-AB                   | pQE15A derivative containing gdrA, gdrB genes  | This study            |
| Primers                  | Sequence (5' to 3')  | Source                |
| P30 F                    | CCTATAAAAATAGGCGTATCACGAGGC  | This study            |
| P30 R                    | ATTGACTACCGGATAGCTTGGATTCTCACCAATAAAAAACGCC  | This study            |
| P15A F                   | GAGAATCCAAGCTATCCGGTAGTCAATAAACCGGTAAACCA  | This study            |
| P15A R                   | GGGCATGACTAACATGAGAATTACAACTTATAT  | This study            |
| DB S1 F                  | C <u>GAGCTC</u> ATGAAAAGATCAAAACGATTTGCAGTAC   | This study            |
| DB K1 R                  | GG <u>GGTACCT</u> TAGCTTCCTTTACGCAGCTTATG  | This study            |
| YD K1 F                  | GG <u>GGTACC</u> ATGAACAACTTTAATCTGCACACCC   | This study            |
| YD H3 R                  | CCC <u>AAGCTT</u> TTAGCGGGCGGCTTCGTATATA   | This study            |
| GA S1 F                  | C <u>GAGCTC</u> ATGCCGTTAATAGCCGGGATTGAT   | This study            |
| GA X1 R                  | TCC <u>CCCGGG</u> TTAATTCGCCTGACCGGCCAG  | This study            |
| GB X1 F                  | TCC <u>CCCGGG</u> ATGTCGCTTTCACCGCCAGG   | This study            |
| GB H3 R                  | CCC <u>AAGCTT</u> TCAGTTTCTCTCACTTAACGGCAG   | This study            |

Table 1. Bacterial strains, plasmids, and primers used in this study

pQE30 with chloramphenicol-resistance and p15A-origin genes of pACYCDuet-1. The *dha*B123 complex gene from *K. pneumoniae* AJ4 and the *yqh*D gene from *E. coli* were amplified by PCR using primer the sets DB S1 F-DB K1 R and YD K1 F-YD H3 R, respectively. The amplified *dha*B123 fragment was cloned into the *SacI* and *KpnI* sites of pQE30 to generate pQE30/*dha*B123. Then, the amplified *yqh*D fragment was introduced into the *KpnI* and *Hind*III sites of pQE30/*dha*B123 to construct pQE30/*dha*B123, *yqh*D (p30-BY) (Supplementary Fig. 1A). To construct pQE15A/gdrAB (p15-AB) (Supplementary Fig. 1B), the gdrA and gdrB genes from *K. pneumoniae* AJ4 were amplified using the following two pairs of primers, respectively: GA S1 F, GA X1 R; GB X1 F, GB H3 R. The *gdr*A PCR fragment was inserted into the *SacI* and *XmaI* sites of pQE15A to generate pQE15A/*gdr*A. Then, the *gdr*B PCR fragment was cloned into the *XmaI* and *Hind*III sites of pQE15A/*gdr*A to construct pQE15A/*gdr*AB. The constructed plasmids were transformed into *E. coli* strains to generate engineered strains.

# 2.2. Flask cultures and fed-batch fermentation

The 1,3-PDO-producing *E. coli* strains were inoculated into 125-mL baffled Erlenmeyer flasks containing 20 mL Luria-Bertani (LB) broth at 37°C and 180 rpm. The cultures were supplemented with 100 mg/L ampicillin and 25 mg/L chloramphenicol, per requirement. When the cell density at OD<sub>600</sub> reached 1.0, 30 g/L glycerol, 0.2 mM isopropyl  $\beta$ -D-1-thiogalactopyranoside (IPTG), and 10  $\mu$ M coenzyme B<sub>12</sub> were added, and the culture was further incubated at 30°C for 40 h. All strains were analyzed for 1,3-PDO production. To screen the bacterial 1,3-PDO production efficiency with the addition of TCA cycle intermediates, such as malate, succinate, and fumarate; a 5, 10 or 20 mM intermediate was added to the medium at IPTG induction. All flask-culture experiments were performed in triplicate.

Fed-batch fermentation experiments were carried out at 37°C, 400 rpm, and at pH 7.0  $\pm$  0.1 in a 1.5 L bioreactor (BioTron, Seoul, Korea), with an initial working volume of 1 L. The same LB medium as that in the flask culture experiments was used. The same concentrations of IPTG and coenzyme B<sub>12</sub> were added when the cell OD<sub>600</sub> reached at 1.0  $\pm$  0.1. One exception was that the glycerol concentration was decreased to 5 g/L and the culture was further incubated at 30°C. The initial glycerol concentration was 5 g/L and feeding solution was added during 5 ~ 60 h to maintain glycerol concentration within 4 ~ 6 g/L with different feeding rates. The succinate concentration was initially 5 mM and when it decreased to 1  $\pm$  0.5 mM, succinate was fed to increase the concentration to 5  $\pm$  1 mM.

#### 2.3. Analytical methods

The culture samples were centrifuged at  $13,000 \times g$  for 5 min and filtered through a 0.2-µm membrane filter. Organic compounds and carbon sources; including glycerol, 1,3-PDO, acetate, lactate, malate, succinate, and fumarate were measured using a Waters high-performance liquid chromatography system equipped with a refractive index detector, and an organic acid analysis column was used for separation (Aminex HPX-87H, 300 mm  $\times$  7.8 mm, 9  $\mu$ m; Bio-Rad Co., Richmond, CA, USA). The column was eluted with sulfuric acid (4 mM) at a flow rate of 0.4 mL/min at 30°C [20,21]. Cell growth was determined by measuring the optical density at 600 nm using a spectrophotometer.  $OD_{600}$ measurements were converted to the dry weight using a calibration curve to relate  $OD_{600}$  to dry cell weight (DCW). The correlation factor was DCW (g/L) =  $0.45 \times OD_{600}$ . The intracellular redox balance (represented by the ratio of NADH to NAD<sup>+</sup>) was determined by NAD/NADH-Glo<sup>™</sup> assay kit (Promega), according to the manufacturer's instructions.

### 3. Results and Discussion

# 3.1. 1,3-PDO production from glycerol varied with and without p15-AB or strain specificity

The glycerol dehydratase reactivation factors were assumed to serve as a reactivating factor for inactivated glycerol dehydratase [22]. To investigate the effects of *gdr*AB on 1,3-PDO fermentation, the expression plasmids p30-BY and p15-AB were used to construct several engineered *E. coli* strains (Table 1). The glycerol consumption of engineered *E. coli* strains harboring p30-BY alone was lower than *E. coli* strains harboring both p30-BY and p15-AB (Table 2). Due to inactivation of glycerol dehydratase in the fermentation process, it is necessary to co-express the *gdr*AB genes encoding glycerol dehydratase reactivating factor with *dha*B123 and *yqh*D in the engineered *E. coli* to produce 1,3-PDO efficiently. Thus, we constructed the pQE15A plasmid and a dual-vector system in *E. coli*.

Table 2 also shows the appropriate strains used as a host for the production of 1,3-PDO. Under the dual- vector system, we tested three commonly used laboratory E. coli strains, E. coli XL1-blue, E. coli JM109, and E. coli W3110, for their ability to convert glycerol into 1,3-PDO. E. coli JM109 was able to produce the maximum amount of 1,3-PDO (7.2 g/L) compared to that production by E. coli XL1-blue (4.0 g/L) and E. coli W3110 (0.7 g/L) (Table 2). The expression levels and plasmid stability of three laboratory E. coli strains were similar (data not shown). However, the rates of substrate uptake per cell g (dry cell weight) were different. The maximum substrate uptake rate (8.5 g-glycerol/g-DCW) by E. coli JM109 was obtained compared to that rate by E. coli XL1-blue (6.1 g-glycerol/ g-DCW) and E. coli W3110 (5.0 g-glycerol/g-DCW). Therefore, E. coli JM109 was selected for further studies. The results show that glycerol permeability in different E. coli stains may vary significantly and therefore affect production of 1,3-PDO. This research can be applied to other studies where E. coli are used to convert glycerol to alcohols and bio-chemicals.

# **3.2.** The addition of TCA cycle intermediates improved 1,3-PDO production from glycerol

The conversion of glycerol to 1,3-PDO is coupled with the

Table 2. 1,3-PDO production in E. coli strains harboring pQE30/dhaB123, yqhD (p30-BY) with and without pQE15A/gdrA, gdrB (p15-AB)

| Mother strain    | Plasmid(s) | Glycerol consumption (g/L) | 1,3-Propanediol titer (g/L) | Dry cell weight (g/L) |
|------------------|------------|----------------------------|-----------------------------|-----------------------|
| E. coli XL1-blue | 30BY       | $11.12 \pm 0.53$           | $3.10 \pm 0.17$             | $1.73 \pm 0.04$       |
|                  | 30BY15AB   | $11.19 \pm 1.19$           | $4.04\pm0.04$               | $1.84\pm0.01$         |
| E. coli JM109    | 30BY       | $11.49 \pm 0.32$           | $4.06 \pm 0.21$             | $1.93 \pm 0.01$       |
|                  | 30BY15AB   | $17.21 \pm 2.23$           | $7.80\pm0.37$               | $2.03\pm0.08$         |
| E. coli W3110    | 30BY       | $6.20\pm0.42$              | $0.38\pm0.01$               | $1.13 \pm 0.03$       |
|                  | 30BY15AB   | $6.71 \pm 0.63$            | $0.73\pm0.05$               | $1.33 \pm 0.02$       |

conversion of NADH to NAD<sup>+</sup>, requiring the regeneration of NADH for continuous 1,3-PDO production. NADH regeneration requires extra glycerol consumption, which could reduce the conversion of glycerol into 1,3-PDO. In order to overcome these problems, TCA cycle intermediates (malate, succinate, and fumarate) were added to manipulate the intracellular redox balance. By manipulating the ratio of NADH to NAD<sup>+</sup>, production of 1,3-PDO can be improved [23,24].

TCA cycle intermediates were added over a range of  $0 \sim 20$  mM, and the shake-flask fermentation experiments were carried out for 40 h with engineered *E. coli* JM109 (JM-30BY15AB). As shown in Table 3, when TCA

intermediates were added to the medium, the glycerol consumption and 1,3-PDO production increased compared to those observed from control (without the addition of TCA cycle intermediates). Addition of 10 mM succinate caused increased titer of 1,3-PDO (9.9 g/L) compared to that obtained with malate (8.2 g/L) and fumarate (9.4 g/L). Therefore, succinate was selected for manipulating the intracellular redox balance. The optimal succinate concentration was 10 mM.

The time profile of glycerol consumption, 1,3-PDO production, and intracellular redox balance by the engineered *E. coli* JM109 (JM-30BY15AB) was monitored with or without the addition of 10 mM succinate. After 40 h, the

Table 3. Glycerol consumption, 1,3-PDO titer, and dry cell weight of engineered *E. coli* JM109 (JM-30BY15AB) under different TCA cycle intermediate concentrations

|           | Concentration (mM) | Glycerol consumption (g/L) | 1,3-propanediol titer (g/L) | Dry cell weight (g/L) |
|-----------|--------------------|----------------------------|-----------------------------|-----------------------|
| Control   | Non-feeding        | $18.11 \pm 1.23$           | $7.74 \pm 0.17$             | $2.88\pm0.03$         |
| Malate    | 5                  | $18.98 \pm 1.11$           | $7.88\pm0.42$               | $2.86 \pm 0.04$       |
|           | 10                 | $20.95 \pm 0.15$           | $7.96 \pm 0.16$             | $2.99 \pm 0.07$       |
|           | 20                 | $21.20\pm0.83$             | $8.17\pm0.52$               | $2.93\pm0.01$         |
| Succinate | 5                  | $20.14\pm0.67$             | $8.24 \pm 0.49$             | $2.94\pm0.02$         |
|           | 10                 | $23.84\pm0.55$             | $9.91 \pm 0.32$             | $2.97\pm0.01$         |
|           | 20                 | $24.69 \pm 1.05$           | $9.81 \pm 0.46$             | $2.97 \pm 0.03$       |
| Fumarate  | 5                  | $21.14 \pm 1.65$           | $8.62 \pm 0.71$             | $2.84\pm0.12$         |
|           | 10                 | $24.72\pm0.47$             | $9.33 \pm 0.29$             | $3.02 \pm 0.03$       |
|           | 20                 | $25.02\pm0.77$             | $9.46 \pm 0.18$             | $3.01\pm0.01$         |



**Fig. 2.** Time profile of glycerol consumption, 1,3-PDO titer, dry cell weight, ratios of NADH to NAD<sup>+</sup>, and succinate consumption in flask fermentation supplemented with 30 g glycerol/L at 30°C for 40 h. (A and C) 0 mM succinate, (B and D) 10 mM succinate. Error bars indicate one standard deviation, as determined from triplicate experiments. *Closed diamonds*, residual glycerol level; *closed squares*, titer of 1,3-PDO; *closed triangles*, dry cell weight; *open squares*, residual succinate level; *open triangles*, ratio of NADH to NAD<sup>+</sup>.

ratio of NADH to NAD<sup>+</sup> increased by 43% with succinate was addition (Fig. 2). There was an increase of glycerol consumption and 1,3-PDO production by 17 and 38%, respectively, at the time of succinate addition compared to control. Conversion yields of 1,3-PDO from glycerol were similar at 41% regardless of succinate addition. These results show that succinate can be used to increase the ratio of NADH to NAD<sup>+</sup> and to speed up the metabolic flux of 1,3-PDO. Therefore, succinate is a key factor that influences 1,3-PDO production.

# 3.3. 1,3-Propanediol production by the engineered *E. coli* in a bioreactor

Based on the above results, strain JM-30BY15AB exhibited the highest 1,3-PDO titer at 9.9 g/L among the six engineered E. coli. For further optimization of fermentation conditions, a fed-batch experiment was performed in a 1.5 L bioreactor. During 1,3-PDO fermentation, feeding solution was continuously added with different feeding rates to maintain low glycerol concentrations (5  $\pm$  1 g/L). Cell growth and 1,3-PDO production were inhibited when the concentration of glycerol was high [25], so fed-batch experiment with low substrate concentrations were studied. As shown in Fig. 3, the titer of 1,3-PDO and its conversion yield from glycerol were 13.11 g/L and 0.49 g/g, respectively. Under fed-batch fermentation conditions, the titer of 1,3-PDO and its conversion yield were enhanced to 32 and 20%, respectively, compared to those observed under flask culture conditions. This result suggests that fed-batch fermentation is a more efficient way to produce of 1,3-PDO.

Several papers describing construction of *E. coli* capable of producing 1,3-PDO have been published. Skraly *et al.* [14] engineered a recombinant *E. coli* strain that harbored genes from the *dha* operon from *K. pneumoniae*, and produced 6.3 g/L of 1,3-PDO. Rujananon *et al* constructed a recombinant *E. coli* strain by assembling the genes *dha*B1 and *dha*B2 from *Clostridium butyricum* and *yqh*D



**Fig. 3.** Time profile of 1,3-propanediol fed-batch fermentation by recombinant *E.coli* JM109 (JM-30BY15AB) at 30°C for 65 h. *Closed diamonds*, residual glycerol level; *closed squares*, titer of 1,3-PDO; *closed triangles*, dry cell weight; *open squares*, residual succinate level.

from *E. coli* [26]. The current study establishes the optimum conditions for 1,3-PDO production using the response surface methodology, and has produced the maximum values of 1.65 g/L 1,3-PDO production reported in the literature to date. Przystalowska *et al.* [27] constructed an *E. coli* strain harbored six heterologous glycerol catabolism pathway genes from *C. freundii* and *K. pneumoniae*, which used 32.6 g/L of glycerol to produce 10.6 g/L. The engineered *E. coli* JM-30BY15AB did not give satisfying results due to low cell density (2.67 g/L) because the optimization of medium constituents or culture conditions was not performed. If medium constituents and culture conditions were optimized, the cell concentration may increase which could result in a higher titer of 1,3-PDO production.

## 4. Conclusion

In the metabolic pathway of glycerol conversion to 1,3-PDO, the enzyme dhaB123 dehydrates glycerol to 3-HPA and then the enzyme *yqh*D reduces 3-HPA to 1,3-PDO. In this study, we achieved efficient production of 1,3-PDO from glycerol by engineered E. coli JM109 via the coexpression of gdrAB and addition of succinate. Construction of a dual-vector system and succinate addition during flask culture enhanced the glycerol consumption and titer of 1,3-PDO. The glycerol consumption and titer of 1,3-PDO were 86.6 and 145.6% higher, respectively, than those from the control (without co-expression of gdrAB and no addition of succinate), and the engineered E. coli JM-30BY15AB exhibited the highest 1,3-PDO titer at 13.11 g/L under fedbatch fermentation conditions. There are two possible reasons for these results. First, co-expression of gdrAB using a dual-vector system could effectively activate the inactivated glycerol dehydratase. This means that a high level of activated glycerol dehydratase could be achieved during 1,3-PDO fermentation. Second, the ratio of NADH to NAD<sup>+</sup> could be increased by addition of succinate, such that more reducing power is used for converting glycerol to 1,3-PDO. The dual-vector system constructed in this work suggests a useful approach for co-expressing heterologous proteins in E. coli, and succinate addition to the medium is a promising route for the production of NADH-dependent bio-chemicals.

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#### References

- Shams Yazdani, S. and R. Gonzalez (2008) Engineering *Escherichia coli* for the efficient conversion of glycerol to ethanol and co-products. *Metab. Eng.* 10: 340-351.
- Dobson, R., V. Gray, and K. Rumbold (2012) Microbial utilization of crude glycerol for the production of value-added products. *J. Ind. Microbiol. Biotechnol.* 39: 217-226.
- Biebl, H., K. Menzel, A. P. Zeng, and W. D. Deckwer (1999) Microbial production of 1,3-propanediol. *Appl. Microbiol. Biotechnol.* 52: 289-297.
- Saxena, R. K., P. Anand, S. Saran, and J. Isar (2009) Microbial production of 1,3-propanediol: Recent developments and emerging opportunities. *Biotechnol. Adv.* 27: 895-913.
- Gonzalez-Pajuelo, M., J. C. Andrade, and I. Vasconcelos (2005) Production of 1,3-Propanediol by *Clostridium butyricum* VPI 3266 in continuous cultures with high yield and productivity. *J. Ind. Microbiol. Biotechnol.* 32: 391-396.
- Cheng, K. K., J. A. Zhang, D. H. Liu, Y. Sun, M. D. Yang, and J. M. Xu (2006) Production of 1,3-propanediol by *Klebsiella pneu-moniae* from glycerol broth. *Biotechnol. Lett.* 28: 1817-1821.
- Zhu, C. and B. Fang (2013) Application of a two-stage temperature control strategy to enhance 1,3-propanediol productivity by *Clostridium butyricum. J. Chem. Tech. Biotechnol.* 88: 853-857.
- Gonzalez-Pajuelo, M., J. C. Andrade, and I. Vasconcelos (2004) Production of 1,3-propanediol by Clostridium butyricum VPI 3266 using a synthetic medium and raw glycerol. *J. Ind. Microbiol. Biotechnol.* 31: 442-446.
- Hong, E., S. Yoon, J. Kim, E. Kim, D. Kim, S. Rhie, and Y. W. Ryu (2013) Isolation of microorganisms able to produce 1,3-propanediol and optimization of medium constituents for Klebsiella pneumoniae AJ4. *Bioproc. Biosyst. Eng.* 36: 835-843.
- Pflugl, S., H. Marx, D. Mattanovich, and M. Sauer (2014) Heading for an economic industrial upgrading of crude glycerol from biodiesel production to 1,3-propanediol by Lactobacillus diolivorans. *Bioresour. Technol.* 152: 499-504.
- Zheng, Z. M., Q. L. Hu, J. Hao, F. Xu, N. M. Guo, Y. Sun, and D. H. Liu (2008) Statistical optimization of culture conditions for 1,3-propanediol by *Klebsiella pneumoniae* AC 15 via central composite design. *Bioresour. Technol.* 99: 1052-1056.
- Oh, B., J. Seo, M. Choi, and C. Kim (2008) Optimization of culture conditions for 1,3-propanediol production from crude glycerol by *Klebsiella pneumoniae* using response surface methodology. *Biotechnol. Bioproc. Eng.* 13: 666-670.
- Ahrens, K., K. Menzel, A. Zeng, and W. Deckwer (1998) Kinetic, dynamic, and pathway studies of glycerol metabolism by *Klebsiella pneumoniae* in anaerobic continuous culture: III. Enzymes and fluxes of glycerol dissimilation and 1,3-propanediol formation. *Biotechnol. Bioeng.* 59: 544-552.
- 14. Skraly, F. A., B. L. Lytle, and D. C. Cameron (1998) Construction and characterization of a 1,3-propanediol operon. *Appl.*

Environ. Microbiol. 64: 98-105.

- Knietsch, A., S. Bowien, G. Whited, G. Gottschalk, and R. Daniel (2003) Identification and characterization of coenzyme B12dependent glycerol dehydratase- and diol dehydratase-encoding genes from metagenomic DNA libraries derived from enrichment cultures. *Appl. Environ. Microbiol.* 69: 3048-3060.
- Seo, J. W., M. Y. Seo, B. R. Oh, S. Y. Heo, J. O. Baek, D. Rairakhwada, L. H. Luo, W. K. Hong, and C. H. Kim (2010) Identification and utilization of a 1,3-propanediol oxidoreductase isoenzyme for production of 1,3-propanediol from glycerol in Klebsiella pneumoniae. *Appl. Microbiol. Biotechnol.* 85: 659-666.
- Toraya, T. and R. H. Abeles (1980) Inactivation of dioldehydrase in the presence of a coenzyme-B12 analog. *Arch. Biochem. Biophys.* 203: 174-180.
- Kajiura, H., K. Mori, T. Tobimatsu, and T. Toraya (2001) Characterization and mechanism of action of a reactivating factor for adenosylcobalamin-dependent glycerol dehydratase. *J. Biol. Chem.* 276: 36514-36519.
- Honda, S., T. Toraya, and S. Fukui (1980) *In situ* reactivation of glycerol-inactivated coenzyme B12-dependent enzymes, glycerol dehydratase and diol dehydratase. *J. Bacteriol.* 143: 1458-1465.
- Ma, B. B., X. L. Xu, G L. Zhang, L. W. Wang, M. Wu, and C. Li (2009) Microbial production of 1,3-propanediol by *Klebsiella pneumoniae* XJPD-Li under different aeration strategies. *Appl. Biochem. Biotechnol.* 152: 127-134.
- Xue, X., W. Li, Z. Li, Y. Xia, and Q. Ye (2010) Enhanced 1,3propanediol production by supply of organic acids and repeated fed-batch culture. *J. Ind. Microbiol. Biotechnol.* 37: 681-687.
- Toraya, T. and K. Mori (1999) A reactivating factor for coenzyme B12-dependent diol dehydratase. J. Biol. Chem. 274: 3372-3377.
- Zhang, Y., Z. Huang, C. Du, Y. Li, and Z. Cao (2009) Introduction of an NADH regeneration system into *Klebsiella oxytoca* leads to an enhanced oxidative and reductive metabolism of glycerol. *Metab. Eng.* 11: 101-106.
- Ma, Z., X. Shentu, Y. Bian, and X. Yu (2013) Effects of NADH availability on the *Klebsiella pneumoniae* strain with 1,3-propanediol operon over-expression. *J. Basic Microbiol.* 53: 348-354.
- Zhu, M. M., P. D. Lawman, and D. C. Cameron (2002) Improving 1,3-propanediol production from glycerol in a metabolically engineered *Escherichia coli* by reducing accumulation of snglycerol-3-phosphate. *Biotechnol. Progr.* 18: 694-699.
- Rujananon, R., P. Prasertsan, and A. Phongdara (2014) Biosynthesis of 1,3-propanediol from recombinant *E. coli* by optimization process using pure and crude glycerol as a sole carbon source under two-phase fermentation system. *World J. Microbiol. Biotechnol.* 30: 1359-1368.
- Przystalowska, H., J. Zeylanda, D. Szymanowska-Powatowska, M. Szalata, R. Stomski, and D. Lipinski (2015) 1,3-Propanediol production by new recombinant *Escherichia coli* containing genes from pathogenic bacteria. *Microbiol. Res.* 171: 1-7.