

RESEARCH ARTICLE

Production of 3-Hydroxypropionic Acid via the Propionyl-CoA Pathway Using Recombinant *Escherichia coli* Strains

Hui Luo^{1*}, Dafeng Zhou¹, Xiaohui Liu¹, Zhihua Nie^{2,3}, Diego Leandro Quiroga-Sánchez¹, Yanhong Chang^{2,3*}

1 Department of Biological Science and Engineering, University of Science and Technology Beijing, Beijing, 100083, China, **2** Department of Environmental Engineering, University of Science and Technology Beijing, Beijing, 100083, China, **3** Key Laboratory of Educational Ministry for High Efficient Mining and Safety in Metal Mine, University of Science and Technology Beijing, Beijing, 100083, China

* luohui@ustb.edu.cn (HL); yhchang@ustb.edu.cn (YC)



CrossMark
click for updates

OPEN ACCESS

Citation: Luo H, Zhou D, Liu X, Nie Z, Quiroga-Sánchez DL, Chang Y (2016) Production of 3-Hydroxypropionic Acid via the Propionyl-CoA Pathway Using Recombinant *Escherichia coli* Strains. PLoS ONE 11(5): e0156286. doi:10.1371/journal.pone.0156286

Editor: Ivan A. Berg, University of Freiburg, GERMANY

Received: March 22, 2015

Accepted: May 11, 2016

Published: May 26, 2016

Copyright: © 2016 Luo et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the paper and its Supporting Information files.

Funding: This work was supported by the National Science foundation of China Grant number: 21276023, <http://www.nsfc.gov.cn/>. HL received the funding. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

Abstract

Our study aimed to produce the commercially promising platform chemical 3-hydroxypropionic acid (3-HP) via the propionyl-CoA pathway in genetically engineered *Escherichia coli*. Recombinant *E. coli* Ec-P overexpressing propionyl-CoA dehydrogenase (PACD, encoded by the *pacd* gene from *Candida rugosa*) under the T7 promoter produced 1.33 mM of 3-HP in a shake flask culture supplemented with 0.5% propionate. When propionate CoA-transferase (PCT, encoded by the *pct* gene from *Megasphaera elsdenii*) and 3-hydroxypropionyl-CoA dehydratase (HPCD, encoded by the *hpcd* gene from *Chloroflexus aurantiacus*) were expressed along with PACD, the 3-HP titer of the resulting *E. coli* Ec-PPH strain was improved by 6-fold. The effect of the cultivation conditions on the 3-HP yield from propionate in the Ec-PPH strain was also investigated. When cultured at 30°C with 1% glucose in addition to propionate, 3-HP production by Ec-PPH increased 2-fold and 12-fold compared to the cultivation at 37°C (4.23 mM) or without glucose (0.68 mM). Deletion of the *ygfH* gene encoding propionyl-CoA: succinate CoA-transferase from Ec-PPH (resulting in the strain Ec-ΔY-PPH) led to increase of 3-HP production in shake flask experiments (15.04 mM), whereas the strain Ec-ΔY-PPH with deletion of the *prpC* gene (encoding methylcitrate synthase in the methylcitrate cycle) produced 17.76 mM of 3-HP. The strain Ec-ΔY-ΔP-PPH with both *ygfH* and *prpC* genes deleted produced 24.14 mM of 3-HP, thus showing an 18-fold increase in the 3-HP titer in compare to the strain Ec-P.

Introduction

3-Hydroxypropionic acid (3-HP; C₃H₆O₃; MW 90.08) is one of the most valuable platform chemicals due to its great potential in the synthesis of novel polymer materials and other derivatives, such as 1,3-propanediol, malonic acid, acrylonitrile, and acrylamide, all of which are used in large quantities in industry [1–3]. Compared to the present costly multi-step chemical

processes, the biosynthesis of 3-HP by microorganisms is considered to be an effective alternative for its mild operation conditions and reduced environmental loads [4, 5].

It was reported that 3-HP could be accumulated as an end product from glycerol, glucose or acrylic acid in a variety of microorganisms [6–8]. Several distinct enzymatic routes have been proposed for microbial 3-HP production [9–14], among them, the modified β -oxidation pathway (also known as the propionyl-CoA pathway: propionic acid → propionyl-CoA → acryloyl-CoA → 3-hydroxypropionyl-CoA → 3-HP) that likely exists in some microorganisms that naturally produce 3-HP [15]. In *E. coli* grown on glucose propionyl-CoA can be derived from succinyl-CoA [16], and a *Candida rugosa* mutant is able to produce 40.2 g/l of 3-HP, presumably via the propionyl-CoA pathway. This is the highest titer obtained from non-engineered microorganisms in the literature. This route is therefore proposed to have great potential for the bioengineering of *E. coli* to produce 3-HP from propionyl-CoA pathway [15]. However, the biological production of 3-HP via the propionyl-CoA pathway in *E. coli* has not been studied.

Only a few key enzymes need to be introduced to establish 3-HP production via the propionyl-CoA pathway in *E. coli* (Fig 1). These enzymes are propionate CoA-transferase (PCT, EC:2.8.3.1) [17–19], catalyzing the CoA transfer from acetyl-CoA to propionate, propionyl-CoA dehydrogenase (PACD, EC:1.3.8.7) [15, 20, 21], which subsequently converts propionyl-CoA to acryloyl-CoA, and 3-hydroxypropionyl-CoA dehydratase (HPCD, EC:4.2.1.17) [22, 23], converting acryloyl-CoA to 3-hydroxypropionyl-CoA. Finally, 3-hydroxypropionyl-CoA (3-HP-CoA) can be converted to 3-HP by the action of either PCT or 3-hydroxypropionyl-CoA hydrolase [17, 18]. In this schematic biosynthesis of 3-HP, competitive pathways such as the methylcitrate cycle (MC cycle) and tricarboxylic acid cycle (TCA cycle) are also indicated (Fig 1).

Propionate and its derivative propionyl-CoA can be metabolized by *E. coli* (propionyl-CoA metabolism pathway in KEGG Database: pathway ID ko00640) [24]. To increase the yield of

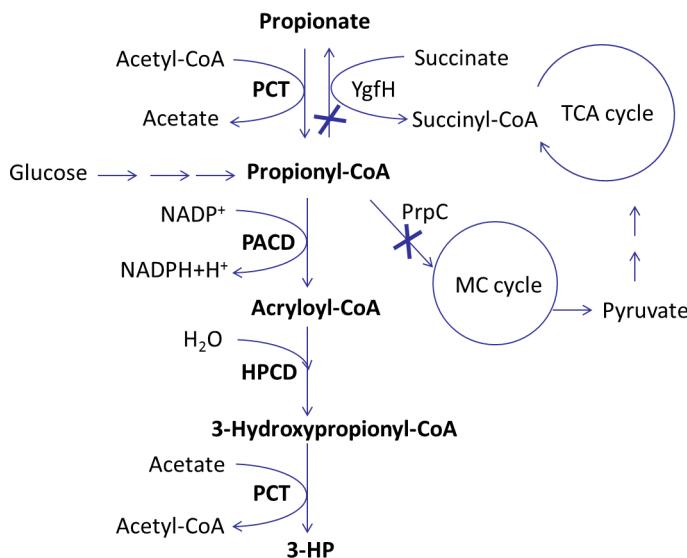


Fig 1. Metabolic pathway involved in the conversion of propionate to 3-hydroxypropionate in recombinant *E. coli* strains. The enzymes in bold are overexpressed, while disrupted pathway steps are indicated by the bold “x” symbols. MC cycle: methylcitrate cycle; TCA cycle: tricarboxylic acid cycle; PCT: propionate CoA-transferase; PACD: propionyl-CoA dehydrogenase; HPCD: 3-hydroxypropionyl-CoA dehydratase; YgfH: propionyl-CoA: succinate-CoA transferase; PrpC: methylcitrate synthase.

doi:10.1371/journal.pone.0156286.g001

3-HP on propionate, these competing propionate metabolic pathways should be down-regulated or even disrupted. Enzymes such as propionyl-CoA: succinate CoA-transferase (*YgfH*) [25, 26] and methylcitrate synthase (*PrpC*) [24, 27, 28] are responsible for the propionate metabolism in *E. coli*, and deleting them could result in an increased yield of 3-HP [26, 28, 29].

In this work, the genes of PCT, HPCD and PACD were obtained from different bacterial species, and a recombinant *E. coli* containing the propionyl-CoA pathway for 3-HP production was constructed. The effects of the culture conditions (e.g., temperature and carbon source) on 3-HP accumulation were also tested. To improve the 3-HP yield on propionate by preventing the degradation of propionyl-CoA in the cells, the *ygfH* and *prpC* genes in the host strain were deleted, and the 3-HP productivity of the engineered strains was evaluated.

Materials and Methods

Materials

The genomic DNA isolation kit and the TA cloning vector pBS-T were purchased from Tiangen Co., Ltd. (Beijing, China). The RNApure yeast kit and cDNA synthesis kit were purchased from Aidlab Biotechnologies Co., Ltd. (Beijing, China). The DNA gel purification kit was purchased from Promega (USA). The restriction and DNA-modifying enzymes were obtained from Takara Co., Ltd. (Dalian, China). The primers were synthesized by Sunbiotech Co., Ltd. (Beijing, China). 3-HP was purchased from Tokyo Chemical Industry Co., Ltd. Propionic acid and all other chemicals were purchased from Biodee Co., Ltd. (Beijing, China) as analytical grade.

Cloning of *pct*, *pacd* and *hpcd* genes

The bacterial strains and the plasmids used in this study are summarized in [Table 1](#). Lysogeny broth (LB) medium was used for the routine genetic engineering, protein expression and culture maintenance. A final concentration of 50 mg/l kanamycin and 25 mg/l chloramphenicol were added to the medium before inoculation. Gene manipulations were carried out using standard methods [30]. To express *pacd*, the pACYCDuet-1 vector with the T7 promoter was adopted. The *pct* and *hpcd* genes were cloned into the pET-28a vector. Details on the construction of the plasmids are shown in [Fig 2](#).

To construct a recombinant expression vector pACYCDuet-PACD, the full-length cDNA encoding the PACD of *Candida rugosa* was cloned by RT-PCR using the Aidlab Biotechnologies kit according to the manufacturer's instructions, and the coding region of PACD (GenBank number GU338397) was amplified by PCR with the primers listed in [S1 Table](#). The *Nde* I, *Xho* I-digested PCR product was finally incorporated into the MCS-II of the pACYCDuet-1 vector.

The *pct* gene (GenBank: HE576794.1) was obtained from the isolated genomic DNA of *Megasphaera elsdenii* by PCR. The 1554 bp PCR fragment was first ligated in the pBS-T vector by means of TA cloning, and its sequence was confirmed by Sunbiotech Co., Ltd (Beijing). The *pct* gene was then released by the flanking *Nde*I and *Bam*HI sites and sub-cloned into the pET-28a vector to generate pET-PCT.

To obtain the HPCD-overexpressing plasmid pET-HPCD, the *hpcd* gene (encoding HPCD) was amplified from the genomic DNA of *Chloroflexus aurantiacus* and successively constructed in the pBS-T and pET-28a vectors using the same method as for *pct*. The T7-HPCD fragment was then digested via *Bgl*II and *Eco*RI sites from pET-HPCD, and the pET-PCT fragment missing the sequence between the *Bam*HI (the isocaudomer of *Bgl*II) and *Eco*RI sites was incorporated with it, thereby obtaining pET-PH.

The plasmid pACYCDuet-PACD was transformed into *E. coli* BL21(DE3) individually and combined with pET-PH, and the recombinant strains Ec-P and Ec-PPH were obtained for protein expression and 3-HP production, respectively.

Table 1. Bacterial strains and plasmids used in the study.

Strains and plasmids	Description	Source
Strains		
<i>Candida rugosa</i>	Source for <i>pacd</i>	CICC, China
<i>Megasphaera elsdenii</i>	Source for <i>pct</i>	CICC, China
<i>Choroflexus aurantiacus</i>	Source for HPCD gene <i>hpcd</i>	CAS, China
<i>E. coli</i> TOP 10	Cloning host	Novagen
<i>E. coli</i> BL21(DE3)	Expression host	Novagen
<i>E. coli</i> JM109(DE3)	Expression host	Tiangen
<i>Ec</i> -Δ <i>Y</i>	<i>E. coli</i> JM109(DE3) Δ <i>ygfH</i>	This study
<i>Ec</i> -Δ <i>P</i>	<i>E. coli</i> JM109(DE3) Δ <i>prpC</i>	This study
<i>Ec</i> -Δ <i>Y</i> -Δ <i>P</i>	<i>E. coli</i> JM109(DE3) Δ <i>ygfH</i> ,Δ <i>prpC</i>	This study
<i>Ec</i> -P	Recombinant <i>E. coli</i> BL21(DE3) harboring plasmid pACYCDuet-PACD	This study
<i>Ec</i> -PPH	Recombinant <i>E. coli</i> BL21(DE3) harboring plasmid pACYCDuet-PACD and pET-PH	This study
<i>Ec</i> -Δ <i>Y</i> -PPH	Recombinant <i>E. coli</i> JM109(DE3) Δ <i>ygfH</i> harboring plasmid pACYCDuet-PACD and pET-PH	This study
<i>Ec</i> -Δ <i>P</i> -PPH	Recombinant <i>E. coli</i> JM109(DE3) Δ <i>prpC</i> harboring plasmid pACYCDuet-PACD and pET-PH	This study
<i>Ec</i> -Δ <i>Y</i> -Δ <i>P</i> -PPH	Recombinant <i>E. coli</i> JM109(DE3) Δ <i>ygfH</i> ,Δ <i>prpC</i> harboring plasmid pACYCDuet-PACD and pET-PH	This study
Plasmids		
pBS-T	<i>lacZ</i> a, TA cloning vector, T7/T3 promoter, Amp ^r	Tiangen
pET-28a	<i>lacI</i> , expression vector, T7 promoter, PBR322-ori, kanamycin (Kan) ^r	Novagen
pACYCDuet-1	<i>lacI</i> , expression vector, T7 promoter, P15A-ori; 2 sets of MCS, MCS I-His ₆ -N, MCS II-S-tag-N, chloramphenicol (Cm) ^r	Novagen
pKD46	repA101(Ts), ara-gamma-Beta-exo, onR101, ampicillin(Amp) ^r	Reference [31]
pKD13	FTP, Kan ^r	Reference [31]
pCP20	Rep(Ts), cl857λ(Ts), Amp ^r , Cm ^r , FLP	Reference [31]
pBS-T-PCT	<i>pct</i> in pBS-T vector, Amp ^r	This study
pBS-T-HPCD	<i>hpcd</i> in pBS-T vector, Amp ^r	This study
pET- PCT	<i>pct</i> in pET28a vector, Kan ^r	This study
pET-HPCD	<i>hpcd</i> in pET28a vector, Kan ^r	This study
pET-PH	<i>pct</i> and <i>hpcd</i> in pET28a vector, Kan ^r	This study
pACYCDuet-PACD	<i>pacd</i> in pACYCDuet-1 vector, Cm ^r	This study

doi:10.1371/journal.pone.0156286.t001

Gene deletion of *YgfH* and *PrpC*

The *ygfH* and *prpC* gene deletions of *Ec*-PPH were performed by directly disrupting the target gene via kanamycin marker insertion using Red recombination, followed by the kanamycin resistance cassette excision mediated through the flippase recombination enzyme harbored by pCP20 [31]. The primers used for inserting the kanamycin cassette into the *ygfH* and *prpC* genes are listed in S1 Table. The *ygfH*, *prpC*, and both *ygfH* and *prpC* gene deletion mutants are referred to as *Ec*-Δ*Y*-PPH, *Ec*-Δ*P*-PPH, and *Ec*-Δ*Y*-Δ*P* -PPH, respectively.

Protein expression and gel electrophoresis

To verify that the recombinant strains had been properly constructed, the parent strain *E. coli* BL21(DE3) and two recombinant strains (*Ec*-P and *Ec*-PPH) were cultured in LB medium and induced at 25°C with 50 μM IPTG at OD₆₀₀ of 0.8. The cells were harvested after 6 h induction and centrifuged at 10,000 g at 4°C for 15 min. The cell pellets were washed twice and then

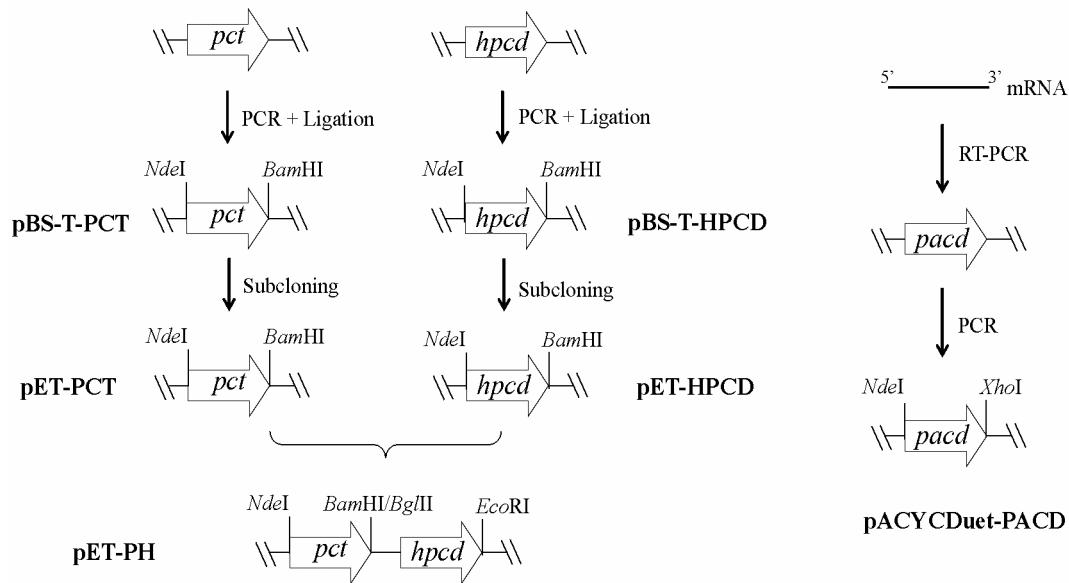


Fig 2. Construction of plasmids pET-PH and pACYCDuet-PACD.

doi:10.1371/journal.pone.0156286.g002

resuspended in 100 mM potassium phosphate buffer (pH 7.0). The cells were lysed by sonication and then centrifuged at 10,000 g for 20 min. The supernatants were subjected to SDS-PAGE to examine protein expression [30]. Coomassie Brilliant Blue R250 was used for protein staining.

Cultivation of recombinant *E. coli* strains

Unless indicated otherwise, shake flask cultivation was carried out with a 20 mL working volume in 100-ml Erlenmeyer flasks at 30°C in an incubator shaker at 200 rpm. The media used in this study were MI medium, containing glucose 1%, peptone 1%, yeast extract 0.5%, NaCl 1% and MgSO₄·7H₂O 0.025%, pH 7.0, and MII medium, which is MI medium without the glucose. Kanamycin (Kan) at 50 mg/l was added to the strain Ec-P. For Ec-PPH, kanamycin and chloramphenicol (Cm) were added at 50 mg/l and 25 mg/l, respectively. The recombinant strains were cultured in the above mediums and induced at OD₆₀₀ of 0.8 with 50 μM IPTG. After 6 h induction, 0.5% propionate was added. Samples were withdrawn periodically to determine the concentrations of 3-HP and residual propionate.

Analytical methods

The propionate and 3-HP concentrations were assayed by high performance liquid chromatography (HPLC). The supernatants obtained by the centrifugation of the culture samples at 10,000 g for 10 min were filtered through a 0.22 μm filter and isocratically eluted through a Venusil MP C18 column (4.6 mm×250 mm, 5 μm particle size, Agela Technologies, Inc.) with methanol-10 mM H₃PO₄ (5:95, v/v) as the mobile phase. The flow rate and the detection wavelength were set to 0.8 ml/min and UV 208 nm, respectively. Pure 3-HP and propionate were assayed by HPLC independently to determine their retention time to be able to identify them later in the chromatogram of each sample. The samples of three independent cultures were assayed, and the standard deviation of the measurements was less than 5% for propionate and 3-HP.

Results

Expression of PCT, HPCD and PACD in recombinant *E. coli*

To construct a pathway as shown in Fig 1, three enzymes corresponding to the formation of the intermediates propionyl-CoA, acryloyl-CoA and 3-hydroxypropionyl-CoA are required.

E. coli possesses genes encoding PrpE (propionyl-CoA synthetase) [32, 33] and enoyl-CoA hydratase [34, 35], which are functionally similar to PCT and HPCD, respectively. It does, however, lack a gene encoding PACD. Therefore, introducing PACD alone may already be enough to establish the pathway in *E. coli*.

It was reported that the overexpression of propionyl-CoA synthetase (PrpE) in *E. coli* could increase the formation of propionyl-CoA [28]. However, the activation of propionate to its CoA ester by PrpE requires two ATPs [35]. Thus, employing a CoA transferase could be preferable for the production of propionyl-CoA, as no additional ATP would be required (Fig 1).

Enoyl-CoA hydratase catalyzes the hydration of α,β -unsaturated CoA thioesters during β -oxidation of fatty acids [36]. There might be low substrate specificity for the enoyl-CoA hydratase of *E. coli* for acryloyl-CoA, which is not a native metabolite in the host cell. Hence, the gene *hpcd* from *Chloroflexus aurantiacus* [22], encoding an enoyl-CoA hydratase specific for acryloyl-CoA, was cloned and expressed in the recombinant *E. coli*.

Therefore, to produce 3-HP via the propionyl-CoA pathway, the recombinant *E. coli* strains *Ec-PPH* (expressing PACD, PCT and HPCD) and *Ec-P* (expressing PACD alone) were constructed, and the protein expressions of these strains were assayed. Fig 3 shows the expression profile of the desired recombinant proteins. The heterologous genes of PCT, PACD and HPCD could be expressed in a soluble form in the recombinant *E. coli*. PACD (with a molecular weight of 49 kDa) was found to be the most abundant protein in the cell lysate of *Ec-P*. When

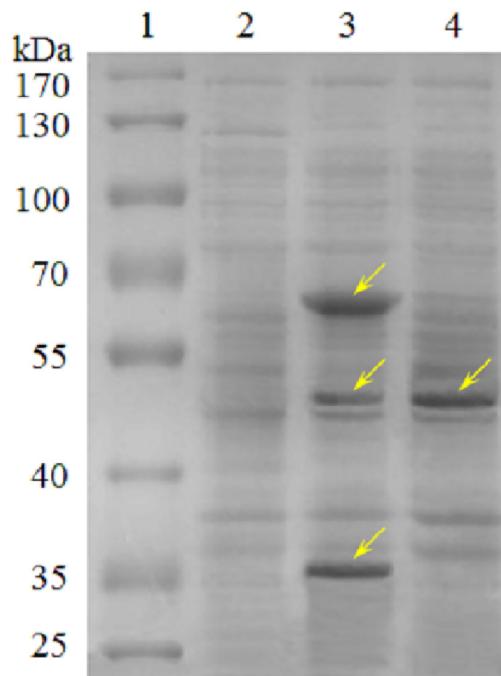


Fig 3. SDS-PAGE analysis of cell extract of the host *E. coli*, *Ec-PPH* and *Ec-P*. (Lane 1) protein standard, (Lane 2) host *E. coli*, (Lane 3) *Ec-PPH* expressing PACD, PCT and HPCD, (Lane 4) *Ec-P* with PACD expression alone. The arrow marks indicate the corresponding target proteins.

doi:10.1371/journal.pone.0156286.g003

PACD was co-expressed with PCT and HPCD in *Ec*-PPH, its expression level decreased to some extent (Fig 3).

Cultivation of the recombinant *E. coli* strains for 3-HP production

To evaluate the potential of the recombinant strains for the production of 3-HP through the propionyl-CoA pathway, the cultivation of *EC-P* and *Ec*-PPH was conducted in MI medium with 0.5% propionic acid at 30°C (Fig 4A). When PACD is expressed alone (*Ec*-P), the final 3-HP titer at 25 h was 1.33 mM. In comparison, when heterologous PCT and HPCD were co-expressed with PACD (*Ec*-PPH), the yield of 3-HP was increased 6-fold to 8.11 mM.

As mentioned, propionyl-CoA utilizing metabolic pathways exist in *E. coli* [24, 26, 37], which lead to a reduced amount of propionyl-CoA being converted to acryloyl-CoA. Furthermore, in contrast to the low K_m value of some *E. coli* enzymes to propionyl-CoA (for example, the K_m value of PrpC to propionyl-CoA is 17 μM) [24], the K_m value of PACD to propionyl-

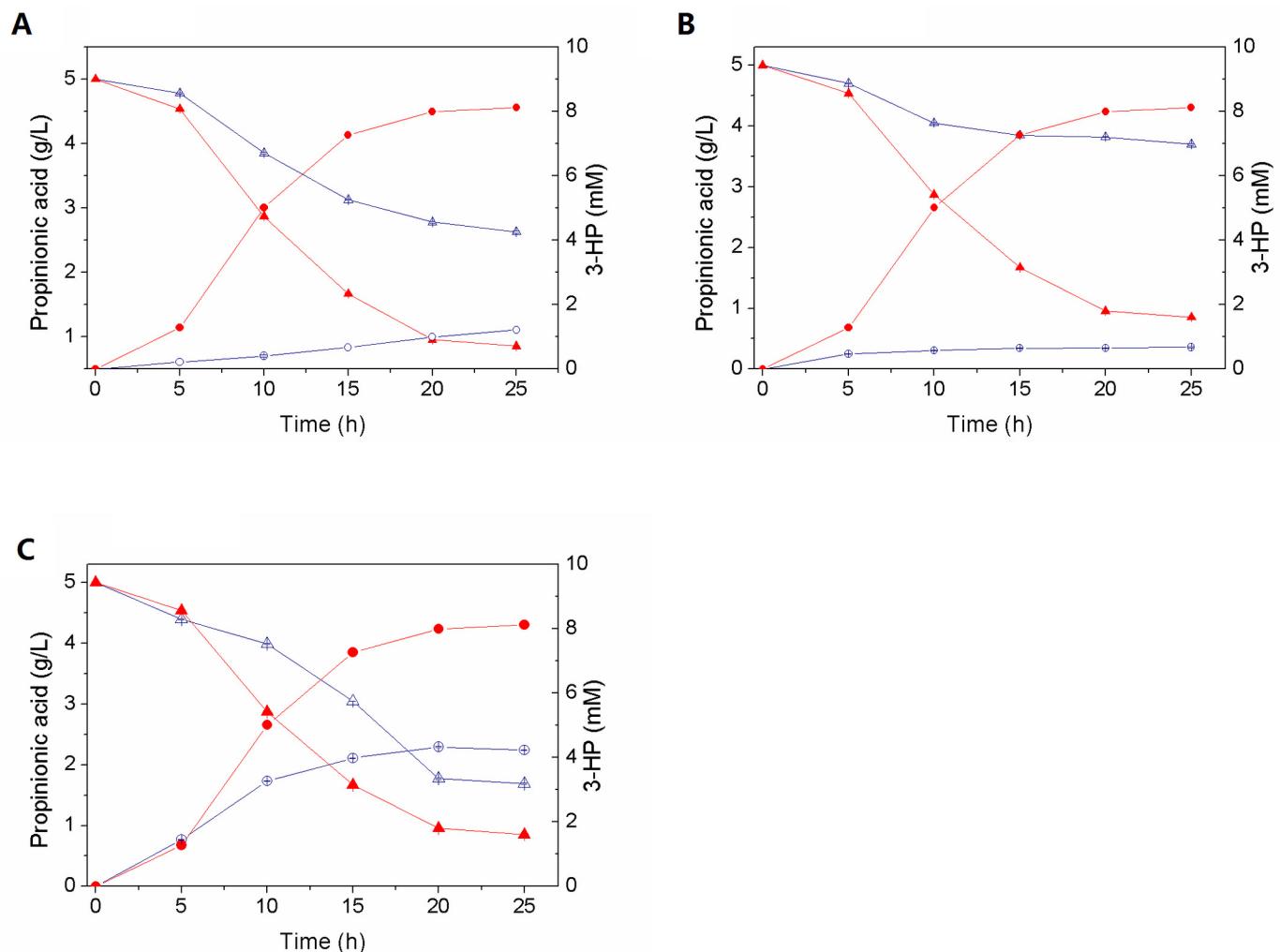


Fig 4. Time-course profile of propionic acid consumption and production of 3-HP for two strains, *Ec*-PPH and *Ec*-P. (A) *Ec*-PPH (expressing PACD PCT and HPCD, closed symbols) and *Ec*-P (expressing PACD, open symbols) were cultured in MI medium at 30°C. (B) *Ec*-PPH was cultivated in medium with glucose (1%, closed symbols) and medium without glucose (open symbols) at 30°C. (C) *Ec*-PPH cultivation in MI medium at 30°C (closed symbols) and 37°C (open symbols). Symbols: 3-HP (circle), propionic acid (triangle). The experiment was done in triplicate. The error bars show standard errors and are not shown if the error did not exceed the size of the symbol.

doi:10.1371/journal.pone.0156286.g004

CoA was reported to be as high as 40.86 μM [21]. Therefore, improvement of the PACD activity is necessary for efficient functioning of the modified β-oxidation pathway. On the other hand, inactivation of the *prpC* gene of the host should be helpful to increase the pool of intracellular propionyl-CoA, and the result of this experiment will be stated later in this work.

The production of 3-HP by *Ec-P* overexpressing PACD alone indicates the presence of the propionyl-CoA synthetase and 3-hydroxypropionyl-CoA dehydratase activities. In the past decades, distinctive types of CoA transferase have been reported in many bacterial strains. Some of these enzymes have been purified from *Escherichia coli* [38], *Peptostreptococcus elsdenii* [18], *Ralstonia eutropha* [39], *Clostridium propionicum* [40] and *Clostridium sp.* strain SB4 [41] and some of them show catalytic activity for the conversion of propionate to propionyl-CoA. In addition, the 3-hydroxypropionyl-CoA dehydratase activities of various hydroxyacyl-CoA dehydratases have been demonstrated to be present in *Escherichia coli* [42] and several other species such as *Metallosphaera sedula* [43] *Sulfolobus tokodaii* [44] and *Nitrosopumilus maritimus* [45].

Compared to *Ec-P*, *Ec-PPH* showed a much higher 3-HP yield, which might be due to the high catalytic efficiencies of the co-expressed enzymes, PCT and HPCD. The overexpression of PCT in *Ec-PPH* was expected to increase the intracellular level of propionyl-CoA, which is an immediate substrate for PACD. The synthesis of propionyl-CoA by PCT does not need ATP, while the production of propionyl-CoA from propionate in *Ec-P* requires the function of its endogenous PrpE, which requires additional ATP [35]. The overexpression of HPCD in *Ec-PPH* could accelerate the consumption of the PACD product, namely acryloyl-CoA, leading to the increased 3-HP yield.

The co-expression of heterologous PCT and HPCD helps to strengthen the desired pathway and reduce the generation of by-products. The results in Fig 4A suggest that to promote 3-HP production via the propionyl-CoA pathway, the PACD activity as well as the propionyl-CoA concentration and acryloyl-CoA conversion in the recombinant cells should be increased.

Glucose, a low-cost and highly efficient substrate, was considered in this work as an additional carbon source that may accelerate the 3-HP synthesis by *Ec-PPH* (Fig 4B).

In the absence of glucose, the production of 3-HP was only 0.67 mM, which is presumably owing to the insufficient ATP supply. As reported in Ref. 46, the synthesis of plasmid-encoded proteins is energetically demanding and leads to a redirection of metabolic fluxes in the cell, e.g. to an increased flow through the TCA cycle [46]. Without the glucose supplementation, propionic acid tends to be used for energy production, thus decreasing the flow through the propionyl-CoA pathway. However, propionic acid is not a preferable energy source as glucose is for *E. coli*, and ATP generated from propionate is far from being enough for the synthesis of growth-related and heterologous proteins, further weakening the 3-HP production.

The addition of glucose fulfilled the rising energy demands and resulted in a 12-fold increase in the 3-HP yield. As demonstrated here, the presence of glucose or another carbon source is required for the highly efficient production of 3-HP.

A previous study [21] reported that the PACD activity was the highest at 30°C. Nevertheless, it decreased to 30% of its original activity when incubated for 2 h at 37°C. At contrast, two other enzymes of the pathway (PCT, HPCD) had higher activity at 37°C, and this temperature is also optimal for *E. coli* [18]. Therefore, We compared the 3-HP production by *Ec-PPH* at 30 and 37°C.

As shown in Fig 4C, *Ec-PPH* produced only 4.23 mM 3-HP at 37°C (with the 3-HP yield from propionate of 8.2%), whereas 8.11mM 3-HP synthesized at 30°C (with the 14.5% yield). It suggests a crucial role of heterologous PACD in the process, thus revealing propionyl-CoA conversion to acryloyl-CoA as a rate-limiting step of the constructed pathway. In contrast, an increase of the temperature to 37°C (which is optimal for *E. coli* metabolism) results in increased by-products formation, weakening desirable 3-HP accumulation.

Effect of *ygfH* and *prpC* deletion on 3-HP production

E. coli is known to be capable to metabolize propionyl-CoA [24, 26, 37]. For example, in the presence of succinate, propionyl-CoA can be converted by CoA-transferase into propionate and succinyl-CoA, and the succinyl-CoA is then utilized in the TCA cycle [26]. Propionyl-CoA could also be converted via the MC cycle to pyruvate, a key intermediate in cellular metabolism [24]. Thus, native *E. coli* metabolism of propionyl-CoA would result in a reduced amount of propionyl-CoA being converted to acryloyl-CoA.

As propionyl-CoA is a key precursor for 3-HP formation, the *E. coli* strain was further engineered to supply more propionyl-CoA into the 3-HP biosynthesis pathway by disrupting the competing metabolic pathways. 3-HP production from propionic acid was examined on a flask scale using four engineered *E. coli* host strains (wild type *E. coli*, *Ec-ΔygfH*, *Ec-ΔprpC*, and *Ec-ΔygfH ΔprpC*) containing *pacd*, *pct* and *hpcd* (Fig 5); they were tested three times independently.

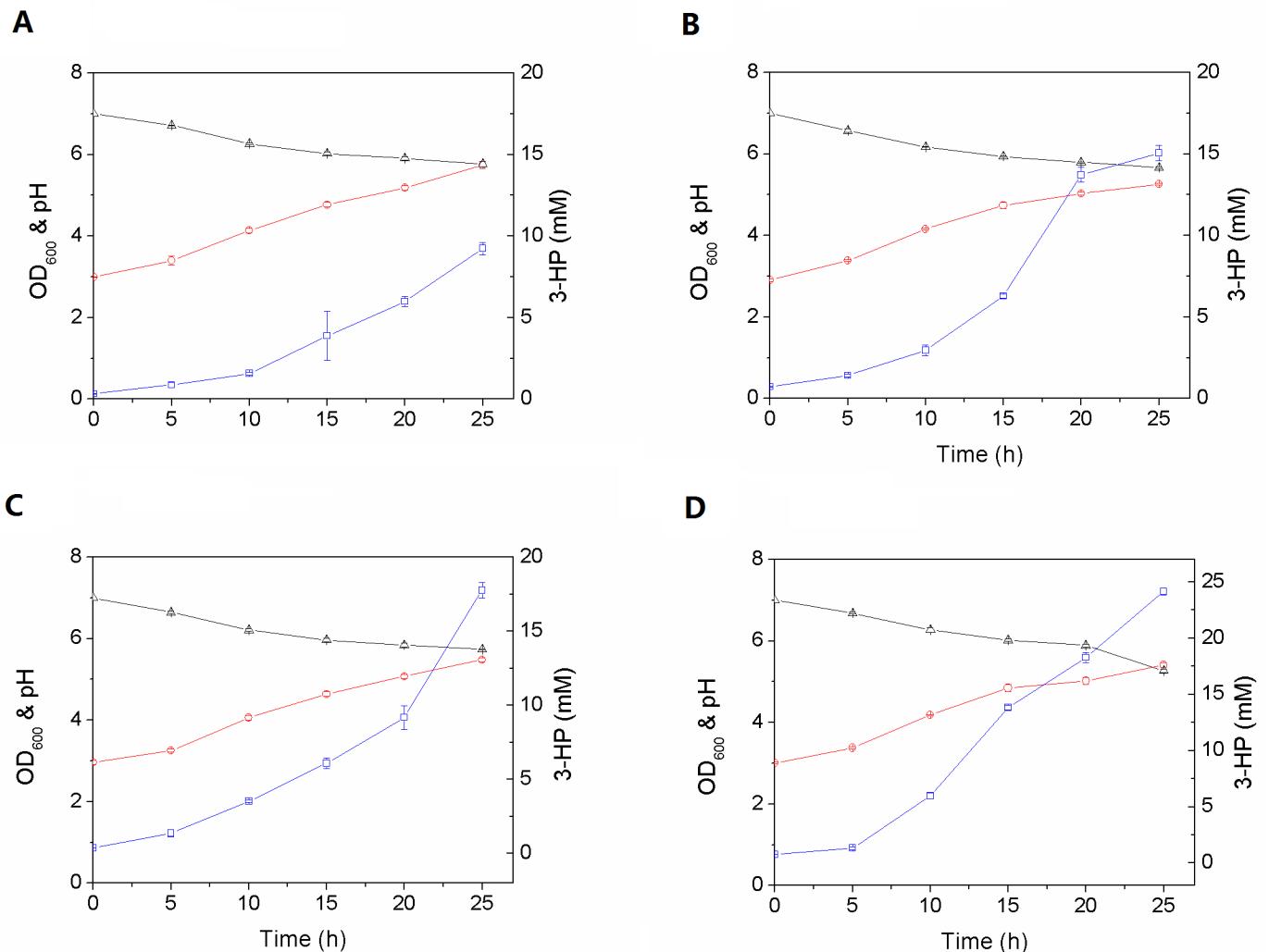


Fig 5. Production of 3-HP by recombinant *E. coli* deletion mutants expressing *pacd*, *pct* and *hpcd*. (A) WT *Ec*, (B) *Ec-ΔygfH*, (C) *Ec-ΔprpC*, and (D) *Ec-ΔygfH ΔprpC*. Symbols: 3-HP (square), cell mass biomass (circle), pH (triangle). The error bars denote standard errors of the mean from triplicate flasks and are not shown where the error did not exceed the size of the symbol.

doi:10.1371/journal.pone.0156286.g005

As shown in Fig 5, the growth of all these strains was very similar, whereas the deletion of the *ygfH* and/or *prpC* genes significantly increased the 3-HP accumulation in the engineered strains (*Ec-ΔY-PPH*, *Ec-ΔP-PPH* and *Ec-ΔY-ΔP-PPH*). Indeed, *Ec-ΔY-PPH* and *Ec-ΔP-PPH* strains produced 15.04 and 17.76 mM of 3-HP, respectively, whereas deletion of both *ygfH* and *prpC* genes in the *Ec-ΔY-ΔP-PPH* strain increased the 3-HP production from 9.23 to 24.14 mM, i.e. 2.6-fold. This underscores the importance of redirecting the metabolic fluxes through the designed pathway by disrupting processes that compete for propionyl-CoA.

Discussion

In this work, recombinant *E. coli* strains were engineered to express the key enzymes (PACD, PCT and HPCD) of the propionyl-CoA pathway for 3-HP production. Recombinant *E. coli* could produce 3-HP when cultured in a propionate-containing medium. Considering that propionyl-CoA is the key intermediate in this pathway and that propionate could be produced by some types of microorganisms [47], including the popular engineering host *E. coli* [16], to produce 3-HP, the propionyl-CoA pathway is a promising alternative route to the commonly reported pathways that use glucose or glycerol as the carbon source [6, 12].

If no propionate was fed in the medium, 3-HP could not be detected in the broth of the recombinant strains during the whole period of cultivation (0–25 h). This result suggests that the propionyl-CoA pool of the host strain was not sufficient to produce a detectable amount of 3-HP. In contrast to glucose or glycerol, propionate is not a usual and economical carbon source for bio-based product synthesis. As propionate itself is a biological product and its biosynthesis pathways in *Propionibacterium* [47] and *E. coli* [16] have been well established, further efforts are required to improve the intracellular propionyl-CoA pool by introducing a propionate/propionyl-CoA synthesis pathway into the recombinant strains mentioned above.

The intracellular propionyl-CoA pool might also be enhanced by inhibiting the endogenous propionyl-CoA catabolism, so we disrupted the metabolic pathways diverted to the MC cycle and/or the TCA cycle (Fig 1). Based on previous reports that the conversion of propionyl-CoA to target products could be improved by the deletion of the *ygfH* gene encoding propionyl-CoA:succinate CoA-transferase [26, 48] or of the *prpC* gene encoding 2-methylcitrate synthase [29, 48], the effect of deleting these genes on the 3-HP production was examined (Fig 5).

The results indicate that the effect of *ygfH* gene deletion on the 3-HP titer was similar to that of *prpC* gene deletion, and the combination of the *ygfH* and *prpC* gene deletions increased the 3-HP titer, exhibiting an efficiency higher than that of either single gene deletion. However, in a previously reported work on the accumulation of a propionyl-CoA derived product, poly (3-hydroxybutyrate-co-3-hydroxyvalerate), in *E. coli* [48], deletion of the *prpC* gene did not significantly increase the 3-HV (3-hydroxyvalerate) fraction, while deletion of the *ygfH* gene significantly increased the 3-HV fraction in the copolymer. By comparing our work to that of Chen et al [48], there were differences in the effects of *ygfH* and *prpC* gene deletion on the product accumulations, suggesting that the metabolic flux from propionyl-CoA to the MC or TCA cycle in *E. coli* was product-dependent.

By culturing these recombinant strains in propionate-containing medium, the conversion yields of propionate to 3-HP were typically in the range of 3% to 15%, implying that the native metabolism of propionate detracts propionyl-CoA from the desired 3-HP biosynthesis. By disrupting the genes for key enzymes of the competing metabolic pathways, *ygfH* and *prpC*, the 3-HP yield on propionate for *Ec-ΔY-ΔP-PPH* increased to 35.4%, not a very high value yet, suggesting that there are other unidentified propionyl-CoA metabolic pathway(s) in the *E. coli* host. Propionyl-CoA carboxylase (PCC), an enzyme that can effectively convert propionyl-CoA to (2S)-methylmalonyl-CoA, has been reported in many species [43, 49], although not in

E. coli. Considering that the carboxylations of acetyl-CoA and propionyl-CoA are often shown to be catalyzed by a single ATP-dependent enzyme [43, 50], the function of propionyl-CoA carboxylation in *E. coli* might be applied by acetyl-CoA carboxylase (ACC). However, ACC is essential for the central metabolism of the host cell and can not be deleted. In addition, the acetyl-CoA:short-chain fatty acid CoA-transferase in *E. coli* might act as the isoenzyme of YgfH because of the broad substrate specificity of bacterial CoA transferases for short-chain acyl-CoA thioesters [18, 38], and the intracellular propionyl-CoA concentration would be reduced by the reaction of CoA transferase even in the ΔygfH strain. Thus, further down-regulating the propionyl-CoA catabolic flux and increasing the intracellular propionyl-CoA pool by, e.g., introducing genes for the synthesis of propionyl-CoA in the host might contribute to the future improvement of the 3-HP yield.

In conclusion, a PACD gene from *Candida rugosa* was heterologously expressed in *E. coli* and its effect on 3-HP production was demonstrated. By overexpressing PCT and HPCD along with PACD, 3-HP production improved six-fold due to the strengthened metabolic flux to 3-HP. The presence of glucose in the medium and the use of an appropriate temperature ensured that there was a sufficient energy source and optimal enzyme thermostability, which contributed to higher 3-HP titers. Furthermore, disruption of the competing propionyl-CoA metabolic pathways by deleting the *ygfH* and *prpC* genes resulted in a 2.6-fold improvement in the 3-HP titer. It is expected that the 3-HP yield of the recombinant strains could be increased by further minimizing the carbon flux to the other metabolites and balancing the activities among the targeted enzymes. To the best of our knowledge, this is the first report on the bio-synthesis of 3-HP in a recombinant strain via the propionyl-CoA pathway.

Supporting Information

S1 Table. Primers used in the study. The primer sequences and the PCR conditions are shown. The restriction enzyme sites in the primer for subsequent cloning are underlined. (DOCX)

Acknowledgments

The authors thank Prof. Huimin Yu at Tsinghua University (Beijing, China) for the generous gift of plasmids pKD13, pKD46 and pCP20.

Author Contributions

Conceived and designed the experiments: HL YC. Performed the experiments: XL DZ ZN DLQ. Analyzed the data: HL XL DZ DLQ YC. Contributed reagents/materials/analysis tools: HL YC. Wrote the paper: HL XL DZ DLQ.

References

1. Della Pina C, Falletta E, Rossi M. A green approach to chemical building blocks. The case of 3-hydroxypropanoic acid. *Green Chem.* 2011; 13(7):1624.
2. Gokarn R, Selifonova O, Jessen H, Steven J, Selmer T, Buckel W. 3-hydroxypropionic acid and other organic compounds. *PCT/US2001/043607*, 2001.
3. Nikolau BJ, Perera M, Brachova L, Shanks B. Platform biochemicals for a biorenewable chemical industry. *Plant J.* 2008; 54(4):536–45. doi: [10.1111/j.1365-313X.2008.03484.x](https://doi.org/10.1111/j.1365-313X.2008.03484.x) PMID: [18476861](#)
4. Lipscomb TE, Lynch MD, Gill RT. Methods, systems and compositions for increased microorganism tolerance to and production of 3-hydroxypropionic acid (3-hp). *WO 2010/011874 A2*, 2010.
5. Henry CS, Broadbelt LJ, Hatzimanikatis V. Discovery and analysis of novel metabolic pathways for the biosynthesis of industrial chemicals: 3-hydroxypropanoate. *Biotechnol Bioeng.* 2010; 106(3):462–73. doi: [10.1002/bit.22673](https://doi.org/10.1002/bit.22673) PMID: [20091733](#)

6. Rathnasingh C, Raj SM, Jo JE, Park S. Development and evaluation of efficient recombinant *Escherichia coli* strains for the production of 3-hydroxypropionic acid from glycerol. Biotechnol Bioeng. 2009; 104(4):729–39. doi: [10.1002/bit.22429](https://doi.org/10.1002/bit.22429) PMID: [19575416](#)
7. Luo LH, Kim CH, Heo S-Y, Oh B-R, Hong W-K, Kim S, et al. Production of 3-hydroxypropionic acid through propionaldehyde dehydrogenase PduP mediated biosynthetic pathway in *Klebsiella pneumoniae*. Bioproc Technol. 2012; 103(1):1–6. doi: [10.1016/j.biotech.2011.09.099](https://doi.org/10.1016/j.biotech.2011.09.099) PMID: [22018749](#)
8. Takamizawa K, Horitsu H, Ichikawa T, Kawai K, Suzuki T. β-hydroxypropionic acid production by *Byssochlamys* sp. grown on acrylic acid. Appl Microbiol Biotechnol. 1993; 40(2–3):196–200.
9. Lynch MD, Gill RT, Lipscomb TE. Methods for producing 3-hydroxypropionic acid and other products. WO 2011/038364, 2011.
10. Saini R, Kapoor R, Kumar R, Siddiqi T, Kumar A. CO₂ utilizing microbes—A comprehensive review. Biotechnol Adv. 2011; 29(6):949–60. doi: [10.1016/j.biotechadv.2011.08.009](https://doi.org/10.1016/j.biotechadv.2011.08.009) PMID: [21856405](#)
11. Jessen HJ, Liao HH, Gort SJ, Selifonova OV. Beta-alanine/alpha-ketoglutarate aminotransferase for 3-hydroxypropionic acid production. US 2011/8030045, 2011.
12. Rathnasingh C, Raj SM, Lee Y, Catherine C, Ashok S, Park S. Production of 3-hydroxypropionic acid via malonyl-CoA pathway using recombinant *Escherichia coli* strains. J Biotechnol. 2012; 157(4):633–40. doi: [10.1016/j.biote.2011.06.008](https://doi.org/10.1016/j.biote.2011.06.008) PMID: [21723339](#)
13. Bar-Even A, Flamholz A, Noor E, Milo R. Rethinking glycolysis: on the biochemical logic of metabolic pathways. Nature chemical biology. 2012; 8(6):509–17. doi: [10.1038/nchembio.971](https://doi.org/10.1038/nchembio.971) PMID: [22596202](#)
14. Finley SD, Broadbelt LJ, Hatzimanikatis V. Thermodynamic analysis of biodegradation pathways. Biotechnol Bioeng. 2009; 103(3):532–41. doi: [10.1002/bit.22285](https://doi.org/10.1002/bit.22285) PMID: [19288443](#)
15. Hasegawa J, Ogura M, Kanema H, Kawaharada H, Watanabe K. Production of β-Hydroxypropionic acid from propionic acid by a *Candida rugosa* mutant unable to assimilate propionic acid: Studies on β-hydroxycarboxylic acids (IV). Journal of fermentation technology. 1982; 60(6):591–4.
16. Haller T, Buckel T, Rétey J, Gerlt JA. Discovering new enzymes and metabolic pathways: conversion of succinate to propionate by *Escherichia coli*. Biochemistry. 2000; 39(16):4622–9. PMID: [10769117](#)
17. Berry A. Methods for producing 3-hydroxypropionic acid and compounds thereof. WO 2009/089457, 2009.
18. Tung K, Wood W. Purification, new assay, and properties of coenzyme A transferase from *Peptostreptococcus elsdenii*. J Bacteriol. 1975; 124(3):1462–74. PMID: [369](#)
19. Wiesenborn DP, Rudolph F, Papoutsakis E. Coenzyme A transferase from *Clostridium acetobutylicum* ATCC 824 and its role in the uptake of acids. Appl Environ Microbiol. 1989; 55(2):323–9. PMID: [2719476](#)
20. Green D, Mii S, Mahler H, Bock RM. Studies on the fatty acid oxidizing system of animal tissues III. Butyryl coenzyme A dehydrogenase. J Biol Chem. 1954; 206(1):1–12. PMID: [13130521](#)
21. Zhou FL, Zhang YG, Zhang RB, Liu W, Xian M. Expression and characterization of a novel propionyl-CoA dehydrogenase gene from *Candida rugosa* in *Pichia pastoris*. Appl Biochem Biotechnol. 2011; 165(7–8):1770–8. doi: [10.1007/s12010-011-9393-x](https://doi.org/10.1007/s12010-011-9393-x) PMID: [21960276](#)
22. Gokarn RR, Selifonova OV, Jessen HJ, Gort SJ, Selmer T, Buckel W. 3-hydroxypropionic acid and other organic compounds. US7186541, 2007.
23. Teufel R, Kung JW, Kockelkorn D, Alber BE, Fuchs G. 3-hydroxypropionyl-coenzyme A dehydratase and acryloyl-coenzyme A reductase, enzymes of the autotrophic 3-hydroxypropionate/4-hydroxybutyrate cycle in the *Sulfolobales*. J Bacteriol. 2009; 191(14):4572–81. doi: [10.1128/JB.00068-09](https://doi.org/10.1128/JB.00068-09) PMID: [19429610](#)
24. Textor S, Wendisch VF, De Graaf AA, Müller U, Linder MI, Linder D, et al. Propionate oxidation in *Escherichia coli*: evidence for operation of a methylcitrate cycle in bacteria. Arch Microbiol. 1997; 168(5):428–36. PMID: [9325432](#)
25. Allen S, Kellermeier R, Stjernholm R, Jacobson B, Wood HG. The isolation, purification, and properties of methylmalonyl racemase. J Biol Chem. 1963; 238(5):1637–42.
26. Zhang H, Boghigian BA, Pfeifer BA. Investigating the role of native propionyl-CoA and methylmalonyl-CoA metabolism on heterologous polyketide production in *Escherichia coli*. Biotechnol Bioeng. 2010; 105(3):567–73. doi: [10.1002/bit.22560](https://doi.org/10.1002/bit.22560) PMID: [19806677](#)
27. Uchiyama H, Tabuchi T. Properties of methylcitrate synthase from *Candida lipolytica*. Agric Biol Chem. 1976; 40(7):1411–8.
28. Yang JE, Choi YJ, Lee SJ, Kang K-H, Lee H, Oh YH, et al. Metabolic engineering of *Escherichia coli* for biosynthesis of poly (3-hydroxybutyrate-co-3-hydroxyvalerate) from glucose. Appl Microbiol Biotechnol. 2014; 98(1):95–104. doi: [10.1007/s00253-013-5285-z](https://doi.org/10.1007/s00253-013-5285-z) PMID: [24113828](#)

29. Park SJ, Kang KH, Lee H, Park AR, Yang JE, Oh YH, et al. Propionyl-CoA dependent biosynthesis of 2-hydroxybutyrate containing polyhydroxyalkanoates in metabolically engineered *Escherichia coli*. *J Biotechnol*. 2013; 165(2):93–8. doi: [10.1016/j.biote.2013.03.005](https://doi.org/10.1016/j.biote.2013.03.005) PMID: [23524059](https://pubmed.ncbi.nlm.nih.gov/23524059/)
30. Sambrook J RD. Molecular cloning: a laboratory manual 3rd ed. 2001.
31. Datsenko KA, Wanner BL. One-step inactivation of chromosomal genes in *Escherichia coli* K-12 using PCR products. *Proc Natl Acad Sci U S A*. 2000; 97(12):6640–5. PMID: [10829079](https://pubmed.ncbi.nlm.nih.gov/10829079/)
32. Pfeifer BA, Admiraal SJ, Gramajo H, Cane DE, Khosla C. Biosynthesis of complex polyketides in a metabolically engineered strain of *E. coli*. *Science*. 2001; 291(5509):1790–2. PMID: [11230695](https://pubmed.ncbi.nlm.nih.gov/11230695/)
33. Ash R, Baird G. Activation of volatile fatty acids in bovine liver and rumen epithelium. Evidence for control by autoregulation. *Biochem J*. 1973; 136:311–9. PMID: [4359516](https://pubmed.ncbi.nlm.nih.gov/4359516/)
34. Wakil SJ. Studies on the fatty acid oxidizing system of animal tissues IX. Stereospecificity of unsaturated acyl CoA hydrase. *Biochim Biophys Acta*. 1956; 19:497–504. PMID: [13315310](https://pubmed.ncbi.nlm.nih.gov/13315310/)
35. Jiang X, Meng X, Xian M. Biosynthetic pathways for 3-hydroxypropionic acid production. *Appl Microbiol Biotechnol*. 2009; 82(6):995–1003. doi: [10.1007/s00253-009-1898-7](https://doi.org/10.1007/s00253-009-1898-7) PMID: [19221732](https://pubmed.ncbi.nlm.nih.gov/19221732/)
36. Bahnsen BJ, Anderson VE, Petsko GA. Structural mechanism of enoyl-CoA hydratase: Three atoms from a single water are added in either an E1cb stepwise or concerted fashion. *Biochemistry*. 2002; 41(8):2621–9. PMID: [11851409](https://pubmed.ncbi.nlm.nih.gov/11851409/)
37. Yuzawa S, Chiba N, Katz L, Keasling JD. Construction of a part of a 3-hydroxypropionate cycle for heterologous polyketide biosynthesis in *Escherichia coli*. *Biochemistry*. 2012; 51(49):9779–81. doi: [10.1021/bi301414q](https://doi.org/10.1021/bi301414q) PMID: [23181847](https://pubmed.ncbi.nlm.nih.gov/23181847/)
38. Sramek SJ, Frerman FE. Purification and properties of *Escherichia coli* coenzyme A-transferase. *Arch Biochem Biophys*. 1975; 171(1):14–26. PMID: [1103739](https://pubmed.ncbi.nlm.nih.gov/1103739/)
39. Lindenkamp N, Schürmann M, Steinbüchel A. A propionate CoA-transferase of *Ralstonia eutropha* H16 with broad substrate specificity catalyzing the CoA thioester formation of various carboxylic acids. *Appl Microbiol Biotechnol*. 2013; 97(17):7699–709. doi: [10.1007/s00253-012-4624-9](https://doi.org/10.1007/s00253-012-4624-9) PMID: [23250223](https://pubmed.ncbi.nlm.nih.gov/23250223/)
40. Selmer T, Willanzheimer A, Hetzel M. Propionate CoA-transferase from *Clostridium propionicum*. *Eur J Biochem*. 2002; 269(1):372–80. PMID: [11784332](https://pubmed.ncbi.nlm.nih.gov/11784332/)
41. Barker H, Jeng I-M, Neff N, Robertson J, Tam F, Hosaka S. Butyryl-CoA: acetoacetate CoA-transferase from a lysine-fermenting *Clostridium*. *J Biol Chem*. 1978; 253(4):1219–25. PMID: [624727](https://pubmed.ncbi.nlm.nih.gov/624727/)
42. Yang SY, Elzinga M. Association of both enoyl coenzyme A hydratase and 3-hydroxyacyl coenzyme A epimerase with an active site in the amino-terminal domain of the multifunctional fatty acid oxidation protein from *Escherichia coli*. *J Biol Chem*. 1993; 268(9):6588–92. PMID: [8454629](https://pubmed.ncbi.nlm.nih.gov/8454629/)
43. Hügler M, Krieger RS, Jahn M, Fuchs G. Characterization of acetyl-CoA/propionyl-CoA carboxylase in *Metallosphaera sedula*. *Eur J Biochem*. 2003; 270(4):736–44. PMID: [12581213](https://pubmed.ncbi.nlm.nih.gov/12581213/)
44. Teufel R, Kung JW, Kockelkorn D, Alber BE, Fuchs G. 3-hydroxypropionyl-coenzyme A dehydratase and acryloyl-coenzyme A reductase, enzymes of the autotrophic 3-hydroxypropionate/4-hydroxybutyrate cycle in the *Sulfolobales*. *J Bacteriol*. 2009; 191(14):4572–81. doi: [10.1128/JB.00068-09](https://doi.org/10.1128/JB.00068-09) PMID: [19429610](https://pubmed.ncbi.nlm.nih.gov/19429610/)
45. Könneke M, Schubert DM, Brown PC, Hügler M, Standfest S, Schwander T, Schada von Borzyskowski L, Erb TJ, Stahl DA, Berg IA. Ammonia-oxidizing archaea use the most energy-efficient aerobic pathway for CO₂ fixation. *Proc Natl Acad Sci U S A*. 2014; 111(22):8239–44. doi: [10.1073/pnas.1402028111](https://doi.org/10.1073/pnas.1402028111) PMID: [24843170](https://pubmed.ncbi.nlm.nih.gov/24843170/)
46. Hoffmann F, Weber J, Rinas U. Metabolic adaptation of *Escherichia coli* during temperature-induced recombinant protein production: 1. Readjustment of metabolic enzyme synthesis. *Biotechnol Bioeng*. 2002; 80(3):313–9. PMID: [12226864](https://pubmed.ncbi.nlm.nih.gov/12226864/)
47. Wang Z, Ammar EM, Zhang A, Wang L, Lin M, Yang ST. Engineering *Propionibacterium freudenreichii* subsp. *shermanii* for enhanced propionic acid fermentation: effects of overexpressing propionyl-CoA: Succinate CoA transferase. *Metab Eng*. 2015; 27:46–56. doi: [10.1016/j.ymben.2014.10.005](https://doi.org/10.1016/j.ymben.2014.10.005) PMID: [25447642](https://pubmed.ncbi.nlm.nih.gov/25447642/)
48. Chen Q, Wang Q, Wei G, Liang Q, Qi Q. Production in *Escherichia coli* of poly(3-hydroxybutyrate-co-3-hydroxyvalerate) with differing monomer compositions from unrelated carbon sources. *Appl Environ Microbiol*. 2011; 77(14):4886–93. doi: [10.1128/AEM.00091-11](https://doi.org/10.1128/AEM.00091-11) PMID: [21652742](https://pubmed.ncbi.nlm.nih.gov/21652742/)
49. Menendez C, Bauer Z, Huber H, Gad'on N, Stetter K-O, Fuchs G. Presence of acetyl coenzyme A (CoA) carboxylase and propionyl-CoA carboxylase in autotrophic *Crenarchaeota* and indication for operation of a 3-hydroxypropionate cycle in autotrophic carbon fixation. *J Bacteriol*. 1999; 181(4):1088–98. PMID: [9973333](https://pubmed.ncbi.nlm.nih.gov/9973333/)
50. Cronan JE, Waldrop GL. Multi-subunit acetyl-CoA carboxylases. *Prog Lipid Res*. 2002; 41(5):407–35. PMID: [12121720](https://pubmed.ncbi.nlm.nih.gov/12121720/)