



Contents lists available at ScienceDirect

Bioresource Technology

journal homepage: www.elsevier.com/locate/biortech



Production of validamycin A from hemicellulose hydrolysate by *Streptomyces hygroscopicus* 5008



Tan-Che Zhou^a, Jian-Jiang Zhong^{a,b,*}

^a State Key Laboratory of Microbial Metabolism, and Laboratory of Molecular Biochemical Engineering, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University, 800 Dong-Chuan Road, Shanghai 200240, China

^b Shanghai Collaborative Innovation Center for Biomanufacturing Technology (SCICBT), East China University of Science and Technology, 130 Meilong Road, Shanghai 200237, China

HIGHLIGHTS

- Corn cob hydrolysate was used as a low-cost feedstock for antibiotic fermentation.
- Xylose in corn cob hydrolysate had a major contribution to VAL-A biosynthesis.
- Higher VAL-A production from corn cob hydrolysate was achieved by an engineered strain.

ARTICLE INFO

Article history:

Received 12 August 2014

Received in revised form 9 October 2014

Accepted 10 October 2014

Available online 20 October 2014

Keywords:

Hemicellulose hydrolysate

D-Xylose

Streptomyces hygroscopicus

Antibiotics fermentation

Validamycin A

ABSTRACT

Validamycin A (VAL-A) is an important agricultural antibiotic produced by *Streptomyces hygroscopicus* 5008, which uses starch as carbon source occupying about 20% of total production cost. To reduce the medium cost, corn cob hydrolysate – a hemicellulose hydrolysate was applied as a low-cost substrate to VAL-A fermentation. It was found that three major sugars in corn cob hydrolysate including D-glucose, D-xylose and L-arabinose could all be utilized by *S. hygroscopicus* 5008 to produce VAL-A while D-xylose was the main contributor. A higher VAL-A production titer from D-xylose was achieved by using a genetically engineered strain TC03 derived from *S. hygroscopicus* 5008, which resulted in 1.27-fold improvement of VAL-A production from the medium containing 13% (v/v) corn cob hydrolysate compared to that by its original strain. A medium cost analysis was done and compared with previous reports. This work indicates a great potential of the hemicellulose hydrolysate as substrate for antibiotic fermentation.

© 2014 Elsevier Ltd. All rights reserved.

1. Introduction

Validamycin A (VAL-A) is a well-known and important C₇N-aminocyclitol antibiotic, which is produced by *Streptomyces hygroscopicus* var. *jinggangensis* 5008 (hereafter *S. hygroscopicus* 5008). It has been used to efficiently prevent and treat sheath blight disease of crops including rice, and its current market in China is around 30–50 thousand ton per year with annual revenue of billions of RMB (Chinese Yuan). Recently this antibiotic is also recognized as an important raw material for production of an antidiabetic drug, voglibose (Yu et al., 2005). Because of its commercial significance, enhancing the VAL-A production has been the focus of many investigations during past decades (Wei et al., 2012; Zhou et al., 2014,

2011). In another aspect, it is evident that reducing the VAL-A production cost is critical to its large-scale production due to its rather low price. The medium in industrial-scale VAL-A production contains more than 100 g L^{−1} of starch (like rice and corn powder) as carbon source, which takes about 70% of the total medium cost (Wei et al., 2012). Therefore, developing a cheap culture medium by using biomass feedstock as substrates could save the industrial production cost, reduce the competition of the starch based food consumption, and also contribute to the sustainable growth of society and economy.

Hemicellulosic biomass is the second most available renewable resource in nature which has great potential as a substrate for microbial fermentation (Wang et al., 2010). D-Xylose is the most abundant pentose sugar in hemicellulosic biomass and can be utilized as an inexpensive substrate by many microorganisms (Wong et al., 1991; Wu et al., 2014; Ye et al., 2013). In bacteria, D-xylose is usually first isomerized by D-xylose isomerase (encoded by *xylA*) to form D-xylulose, then phosphorylated to generate D-xylulose 5-phosphate by D-xylulose kinase (encoded by *xylB*), and

* Corresponding author at: State Key Laboratory of Microbial Metabolism, and Laboratory of Molecular Biochemical Engineering, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University, 800 Dong-Chuan Road, Shanghai 200240, China. Tel.: +86 21 34206968; fax: +86 21 34204831.

E-mail address: jjzhong@sjtu.edu.cn (J.-J. Zhong).

subsequently metabolized through the pentose phosphate (PP) pathway (Wong et al., 1991). According to the genome sequencing result of *S. hygroscopicus* 5008 (GenBank accession: CP003275) from our State Key Laboratory of Microbial Metabolism at Shanghai Jiao Tong University (Wu et al., 2012), two putative D-xylose utilizing genes – *xylA* and *xylB* were identified in the genome. These two genes in *S. hygroscopicus* 5008 are clustered and show 91% amino acid identity to the reported genes in *S. rubiginosus* (Wong et al., 1991), implying the former could also utilize D-xylose as carbon source. On the other hand, sedoheptulose-7-phosphate (S-7-P), the key intermediate metabolite of the PP pathway, is the direct precursor for VAL-A biosynthesis. It has been identified that biosynthesizing 1 mol VAL-A requires 2 mol S-7-P (Bai et al., 2006). Consequently, it seems straightforward to use D-xylose as carbon source for the VAL-A biosynthesis since it could be a direct precursor supply.

Corn cob hydrolysate (or called corn cob molasses, xylose mother liquid) is a typical kind of lignocellulosic biomasses and widely available in China. It is an acid hydrolysate waste which is generated during the industrial-scale production of D-xylose from corn cob (Wang et al., 2010). Corn cob hydrolysate contains a high concentration of mixed sugars (including D-xylose, L-arabinose and D-glucose) and D-xylose occupies their two-third amount (Cheng et al., 2011). Recently, this low cost and high carbohydrate content feedstock has attracted increasing attention for the production of industrially important chemicals such as acetone, butanol and ethanol (Xiao et al., 2012), propionic acid (Liu et al., 2012), L-lactic acid (Wang et al., 2010), and xylitol (Cheng et al., 2011). However, there is still lack of reports on application of such cheap feedstocks to produce commercially interesting antibiotics like VAL-A.

In this study, the possibility of VAL-A production from corn cob hydrolysate was evaluated, and the contribution of each sugar in the corn cob hydrolysate to VAL-A biosynthesis by *S. hygroscopicus* 5008 was investigated. As an engineered strain of *S. hygroscopicus* 5008, which contained 3–5 copies of *val* gene clusters by tandem amplification via the *zouA* system, was recently constructed by our group to enhance the VAL-A production (Zhou et al., 2014), this engineered VAL-A producer was further applied to fermentation with corn cob hydrolysate by fed-batch cultivation to achieve a higher production titer. This work would provide valuable information for researches on antibiotic fermentation using biomass hydrolysis as feedstock.

2. Methods

2.1. Bacterial strains

The *S. hygroscopicus* 5008 wild-type strain (Yu et al., 2005) and its engineered strain TC03, which contained 3–5 copies of *val* gene clusters by the *zouA* tandem amplification (Zhou et al., 2014), were used in this study. *S. hygroscopicus* 5008 was grown on solid Soya-Flour Mannitol (SFM) agar medium (20 g L⁻¹ agar, 20 g L⁻¹ mannitol and 20 g L⁻¹ soybean powder, pH 7.2) at 30 °C for 5 days. Then the spores were collected and stored in 25% glycerol solution at –80 °C. The strain TC03 was grown in liquid Yeast Extract-Malt Extract (YEME) medium, and the mycelia were collected and suspended in 25% glycerol, then stored at –80 °C (Zhou et al., 2014).

2.2. Fermentation conditions

Corn cob hydrolysate containing approximately 500 g L⁻¹ D-xylose, 190 g L⁻¹ L-arabinose and 150 g L⁻¹ D-glucose was supplied by Jiahe Sugar Co., Ltd., Shandong, China (Cheng et al., 2011). The seed medium containing 2% (v/v) corn cob hydrolysate,

2.2 g L⁻¹ soybean meal, 10 g L⁻¹ yeast extract, 2 g L⁻¹ NaCl and 0.8 g L⁻¹ KH₂PO₄ was used for corn cob hydrolysate fermentation. The modified Super Optimal Broth (SOB) medium containing 20 g L⁻¹ tryptone, 5 g L⁻¹ yeast extract, 0.5 g L⁻¹ NaCl and 2.5 mM KCl was applied for both single and mixed sugar fermentation. Fermentation medium contained 23 g L⁻¹ soybean flour, 5 g L⁻¹ yeast extract, 1 g L⁻¹ NaCl and 1.5 g L⁻¹ KH₂PO₄ supplemented with corn cob hydrolysate (at a concentration of 7%, 10%, 13% or 15%, (v/v) or sugars.

For seed culture, 100 µl of spores or mycelia was inoculated into 500 ml shake flasks containing 100 ml seed medium at 30 °C and 220 rpm for 24 h. Then 5 ml of seed cultures was transferred to a 250 ml flask containing 50 ml fermentation medium and cultured at 37 °C and 220 rpm for 5 days. Four independent experiments were performed for each condition.

For the fed-batch cultivation with corn cob hydrolysate as carbon source, two feeding modes were performed: (a) the initial corn cob hydrolysate concentration was 6.5% (v/v), and then about 1.63% (v/v) corn cob hydrolysate was fed at the 36th h, 48th h, 60th h and 72nd h, respectively; and (b) the initial corn cob hydrolysate concentration was 13% (v/v), and then corn cob hydrolysate was fed from 72nd h to 84th h with addition of 0.1% (v/v) per hour. For each condition, four independent experiments were set up.

2.3. Sampling and analytical methods

For each fermentation experiment, a total of 2 ml fermentation broth was collected from quadruplicate flasks at the designated time. Cells and supernatants were separated by centrifugation at 9000×g for 10 min and stored at –80 °C for further analyses. For dry cell weight, due to the insoluble medium component, total intracellular protein was used to represent cell growth as reported earlier (Wei et al., 2012). Standard Bradford method was used to determine the intracellular protein concentration released from the sonication-treated cells (Zhou et al., 2014).

All the metabolites were analyzed by high performance liquid chromatography (1200 Series HPLC, Agilent, Waldbronn, Germany) with different detectors. The supernatants were diluted 5-fold using deionized water and filtered through 0.22-µm-pore-size filters before HPLC analysis. The concentrations of D-glucose, D-xylose and L-arabinose were detected by a refractive index detector (RID) with an Aminex HPX-87H column (Bio-Rad, Hercules, CA). The mobile phase (5 mM H₂SO₄) was applied with a flow rate of 0.6 ml min⁻¹ at 65 °C (Fan et al., 2013). VAL-A and validoxylamine A production were determined by HPLC as previously reported (Yu et al., 2005).

2.4. Statistical analysis

The data consisted of the average of the measurements of four independent sample measurements with the error bars indicating the standard deviation. Statistical significance was analyzed using the Student's *t*-test, where *P* < 0.05 in a two-tailed analysis indicates significance.

3. Results and discussion

3.1. Effect of initial corn cob hydrolysate concentration on VAL-A production

In general, a relatively high sugar concentration may lead to a higher VAL-A titer in its fermentation (Wei et al., 2012). To investigate the feasibility of using corn cob hydrolysate as a low-cost raw material by *S. hygroscopicus* 5008, various initial concentrations of corn cob hydrolysate were attempted for VAL-A fermentation. As

Download English Version:

<https://daneshyari.com/en/article/680313>

Download Persian Version:

<https://daneshyari.com/article/680313>

[Daneshyari.com](https://daneshyari.com)