

# A pilot-scale study of biohydrogen production from distillery effluent using defined bacterial co-culture

# T.M. Vatsala<sup>\*</sup>, S. Mohan Raj<sup>1</sup>, A. Manimaran<sup>2</sup>

Shri AMM Murugappa Chettiar Research Centre, Photosynthesis and Energy Division, Tharamani, Chennai, India, 600 113

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### ABSTRACT

We evaluated the feasibility of improving the scale of hydrogen (H<sub>2</sub>) production from sugar cane distillery effluent using co-cultures of Citrobacter freundii 01, Enterobacter aerogenes E10 and Rhodopseudomonas palustris P2 at 100 m<sup>3</sup> scale. The culture conditions at 100 ml and 2 L scales were optimized in minimal medium and we observed that the co-culture of the above three strains enhanced H<sub>2</sub> productivity significantly. Results at the 100 m<sup>3</sup> scale revealed a maximum of 21.38 kg of H<sub>2</sub>, corresponding to 10692.6 mol, which was obtained through batch method at 40 h from reducing sugar (3862.3 mol) as glucose. The average yield of H<sub>2</sub> was 2.76 mol mol<sup>-1</sup> glucose, and the rate of H<sub>2</sub> production was estimated as 0.53 kg/100 m<sup>3</sup>/h. Our results demonstrate the utility of distillery effluent as a source of clean alternative energy and provide insights into treatment for industrial exploitation. © 2008 International Association for Hydrogen Energy. Published by Elsevier Ltd. All rights

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# 1. Introduction

Hydrogen (H<sub>2</sub>) is a sustainable, non-polluting source of energy. Owing to its potential use, it has been considered as a prominent future fuel and energy vector. The United States and the European countries have already committed to establish an energy economy based on H<sub>2</sub> from fully sustainable or renewable sources. Based on the national H<sub>2</sub> program of the United States, the contribution of H<sub>2</sub> to total energy market will be ~8–10% by 2025 [1]. The market for H<sub>2</sub> and technologies for its production are potentially huge, and the level of investment into H<sub>2</sub> research has increased significantly in recent years, and  $H_2$  may, in the long-term, replace all other fuels. However, cost is currently the biggest impediment to  $H_2$  fuel production.  $H_2$  is produced from nonrenewable energy sources, such as oil, natural gas, and coal that require expensive energy input [2]. Thus, the development of cost-effective and/or economically viable processes is necessary for increasing the achievable dominance of  $H_2$  as an energy source. As a consequence, scientists are pursuing biological systems as a way to meet increasing energy demand. Biological  $H_2$  production stands out as an environmentally friendly process carried out under mild operating conditions that include fermentation of sugars, photolysis of water and water gas shift reactions [3–6].

<sup>\*</sup> Corresponding author. Present address: R&D, Hydrolina Biotech Private Ltd., 406, TICEL Biopark, Tharamani, Chennai, India, 600 113. Tel.: +91 44 2254 1199; fax: +91 44 2471 8155.

E-mail address: tmvatsala@rediffmail.com (T.M. Vatsala).

<sup>&</sup>lt;sup>1</sup> Present address: Department of Chemical and Biochemical Engineering, Pusan National University, Busan 609 735, Republic of Korea.

<sup>&</sup>lt;sup>2</sup> Present address: Department of Biotechnology and Food Technology, Durban University of Technology, Durban 4000, South Africa. 0360-3199/\$ – see front matter © 2008 International Association for Hydrogen Energy. Published by Elsevier Ltd. All rights reserved. doi:10.1016/j.ijhydene.2008.07.015

Biological H<sub>2</sub> production is the most challenging and exigent prefecture of renewable energy technology development, especially where it involves the treatment of organic waste. H<sub>2</sub> can also be produced by bacteria from renewable biomass. Several studies on biohydrogen production from various substrates, including wastes [7-10], have been reported at laboratory level. Vatsala and Seshadri [7] have studied the enzymatic hydrolysis of silk-cotton waste into H<sub>2</sub> and volatile fatty acids (VFAs) using Rhodospirillum rubrm ATCC 11170. Kapdan and Kargi [8] have published an excellent review on the use of various substrates, particularly simple sugars, corn starch wastes, sweet potato, food industrial wastes, wastewater, and waste sludge, for H<sub>2</sub> production under dark and phototrophic conditions. While, chemical wastewater was used successfully for biohydrogen production at lab-scale [9]. Yu et al. [10] have demonstrated the continuous production of H<sub>2</sub> from anaerobic acidogenesis of a highstrength rice winery wastewater by a mixed bacterial flora at 3 L upflow reactor. Various types of bioreactors with different capacities have also been used for H<sub>2</sub> production, including batch, fed-batch and continuous-flow, stirred tank reactors [11–14]. The concept of waste utilization through a microbial fermentation process represents an ideal alternative technology for coordinating pollutant elimination and energy production. In this sense, biohydrogen is considered an intriguing "future fuel". The energy required for aerobic wastewater treatment is huge, but can be reduced through economically viable biohydrogen production processes.

We have been studying H<sub>2</sub> production from various waste materials [15,16]. Among the waste materials examined, distillery effluent was the most suitable substrate for H<sub>2</sub> production [17]. H<sub>2</sub> production from distillery effluent was investigated at lab-scale and pilot-scale [17]. The seasonal variations on the molasses quality, particularly sugar content, sulfur content, and their influences on H<sub>2</sub> production using individual and/or co-culture of Citrobacter freundii and/or Rhodopseudomonas palustris P2 were investigated [17]. We also reported H<sub>2</sub> production from distillery effluent at 10 m<sup>3</sup> volume and the produced H<sub>2</sub> was subsequently tested for fuel cell application [18]. Here, the present investigation focuses on developing a cost-effective, viable process for producing clean  $H_2$  on a pilot-scale (100 m<sup>3</sup>) from distillery effluent. Apart from H<sub>2</sub> as an energy source, the treatment of distillery waste was also accomplished, with a drastic reduction in chemical oxygen demand (COD) and biological oxygen demand (BOD). To the best of our knowledge, this is the first report on H<sub>2</sub> production at 100 m<sup>3</sup> scale.

# 2. Materials and methods

## 2.1. Materials

The biochemical characterization kits for bacterial identification were purchased from Hi-Media (Mumbai, India). The genomic DNA isolation kit was procured from Promega (Madison, WI, USA). DNA gel purification kit was purchased from Qiagen (Mannheim, Germany). Primers were synthesized by Cosmotech Co. Ltd., Korea. All other chemicals, unless otherwise indicated, were obtained from E-Merck (Mumbai, India) and/or Sigma–Aldrich (St. Louis, MO, USA).

#### 2.2. Microorganisms and culture conditions

Heterotrophic bacterial strains (HTB) were isolated from the effluent treatment plant, EID Parry (I) Ltd., Nellikuppam, Tamil Nadu, India. Photoheterotrophic bacterial strains (PTB) were isolated from pond soil, Chengalpet, Tamil Nadu, India. Ormerod's minimal medium [19] was used for the regular maintenance of photoheterotropic isolates. To measure the H<sub>2</sub> production efficiency of the PTB isolates, experiments were conducted in serum bottles (125 ml total volume (tv); 100 ml working volume (wv)) at 37 °C, under photoheterotrophic conditions at 7000 lux intensity with an incandescent light source. Malate at 50 mM was used as a carbon source for PTB isolates. Heterotrophic isolates were cultured in H<sub>2</sub> production medium, which contained (per litre): 1.0 g yeast extract, 1.0 g NaCl, 1.0 g sodium thiosulphate and 10 g glucose, under dark anaerobic conditions for 24 h. This medium was fortified with a potassium phosphate buffer (50 mM; pH 7.0), and the experiments were conducted in serum bottles at 37 °C. Before inoculating the seed culture (HTB and/or PTB), the bottles were flushed with argon (Ar) gas (99.9%) for 10 min to ensure that the bottles were completely deprived of oxygen (O2). Bottles were then sealed with butyl rubber septa and aluminum caps. Isolates were tested for H<sub>2</sub> production efficacy with specific media. The best H<sub>2</sub> producing strains, HTB01, HTB10, and PTB2, were subjected to physiochemical characterization using the methodologies described previously [20].

To further confirm the identification of bacterial isolates, the 16S rRNA genes from the strains HTB01, HTB10, and PTB2 were amplified by using the primers: 27F (5'-AGA GTT TGA TCM TGG CTC AG-3') and 1492R (5'-TAC GGY TAC CTT GTT ACG ACT T-3') and sequenced by Cosmotech, Co. Ltd., Korea. Sequence alignment and analysis of similarity of the 16S rRNA genes were performed with CLUSTAL W [21] and DNA baser RC2 (v2.9.97) programs. A similarity search was done in the Ribosomal Database Project (RDP) [22].

To determine the efficiency of the selected microorganism for  $H_2$  production, several experiments were conducted at a lab-scale with 125 ml serum bottles and a glass bioreactor (Bioengineering, KLF, 2000, SW). Axenic strains were acclimatized on distillery effluent in the laboratory under dark anaerobic conditions to grow and produce  $H_2$  for several generations.

#### 2.3. Bioreactors designing, fabrication and operation

The feasibility of  $H_2$  production and effluent treatment was evaluated at the 100 m<sup>3</sup> scale. The lab-scale process of  $H_2$ production was then scaled-up from 1 L to 100,000 L. As shown in Fig. 1, a sequence of bioreactors was constructed for inoculum preparation for the 100 m<sup>3</sup> reactor. The total volumes of the reactors were 0.125, 1.25, 12.5 and 125 m<sup>3</sup>, and the working volumes were set from 0.1 to 100 m<sup>3</sup>. The heightto-diameter (H/D) ratio of each reactor was set to 1.28. All the reactors were fabricated using mild steel with epoxy coating to prevent chemical and/or biological corrosion. Temperature and pH were monitored using specialized probes for 10 and 100 m<sup>3</sup> reactors. However, temperature and pH were not controlled. A 10% inoculum was used for each reactor. The Download English Version:

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