Bioresource Technology 129 (2013) 599-605

Contents lists available at SciVerse ScienceDirect

Bioresource Technology

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

Spatial uniformity of microbial diversity in a continuous bioelectrochemical system



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HIGHLIGHTS

- ► A bioelectrochemical system with exchangeable electrodes was constructed.
- ▶ Microbial diversity did not differ between electrodes within a time point.
- ▶ Temporal dynamics can be related to community development rather than spatial variation.

ARTICLE INFO

Article history: Received 30 June 2012 Received in revised form 20 November 2012 Accepted 20 November 2012 Available online 4 December 2012

Keywords: Bioelectrochemical system Microbial fuel cell Microbial communities Electroactive biofilms Spatial variation

ABSTRACT

Bioelectrochemical systems (BESs) are emerging as a technology with diverse future applications. Anodeassociated microbial diversity and activity are known to change over time, but the consequences of these dynamics on BES functioning are poorly understood. A novel BES with exchangeable anodic electrodes that facilitates characterisation of microbial communities over time was constructed. The BES, received a mixture of volatile fatty acids and produced 0.13 mA cm⁻² of anodic electrode surface, leading to the removal of 14 g chemical oxygen demand per m² electrode per day at a coulombic efficiency of 76%. Pyrosequencing of 16S rRNA genes revealed no differences in the diversity of microbial communities associated with different electrodes within a single time point. This finding validates the design for temporal studies as changes in microbial diversity observed over time can be related to community development rather than spatial variation within the reactor.

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1. Introduction

Bioelectrochemical Systems (BESs) exploit microorganisms to catalyse oxidation and/or reduction reactions at anodic and cathodic electrodes, respectively. Key applications for BESs include electrical power generation, *i.e.* microbial fuel cells (Davis and Yarbrough, 1962), bioremediation (Gregory and Lovley, 2005) and production of (bio)chemicals (Hongo and Iwahara, 1979). A wide range of electron donors, including organic compounds associated with wastewater, can be oxidised at the anode and facilitate cathodic production of compounds of interest *e.g.* caustic soda (Rabaey et al., 2010), methane (Clauwaert et al., 2008; Cheng et al., 2009) or acetate (Nevin et al., 2010). While modifications to the architecture of BESs and the materials used in their construction have led to significant advances in BES technologies (Logan et al., 2006; Du et al., 2007), a better understanding of

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BES microbiology is critical to achieve stable and optimal performance.

A wide range of parameters, including pH, starting inoculum and electrode potential, are known to influence the composition of anode-associated microbial communities (Torres et al., 2009; White et al., 2009; Patil et al., 2011). Nonetheless, little is known about how electrode-associated communities develop over time and influence anodic and/or cathodic processes. Previous studies have demonstrated pH stratification and differential gene expression within Geobacter sulfurreducens biofilms (Franks et al., 2010), behavioural changes of Shewanella oneidensis MR-1 cells in relation to electron transfer over time (Harris et al., 2010), and temporal structure in the spatial arrangement of different species in coculture (Read et al., 2010). These studies indicate that electrodeassociated microbial communities exhibit differences over time; however, the influence of these changes on the functional stability of BESs is poorly understood. Due to a lack of appropriate BES designs, there are currently no studies that describe in detail the temporal dynamics of electrode-associated microbial communities in detail.



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