



# Anodic and cathodic biofilms coupled with electricity generation in single-chamber microbial fuel cell using activated sludge

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## Abstract

Microbial fuel cell (MFC) is used to remove organic pollutants while generating electricity. Biocathode plays as an efficient electrocatalyst for accelerating the Oxidation Reduction Reaction (ORR) of oxygen in MFC. This study integrated biocathode into a single-chamber microbial fuel cell (BSCMFC) to produce electricity from an organic substrate using aerobic activated sludge to gain more insights into anodic and cathodic biofilms. The maximum power density, current density, chemical oxygen demand (COD) removal, and coulombic efficiency were  $0.593 \text{ W m}^{-3}$ ,  $2.6 \text{ A m}^{-3}$ ,  $83 \pm 8.4\%$ , and  $22 \pm 2.5\%$ , respectively. Extracellular polymeric substances (EPS) produced by biofilm from the biocathode were higher than the bioanode. Infrared spectroscopy and Scanning Electron Microscope (SEM) examined confirmed the presence of biofilm by the adhesion on electrodes. The dominant phyla in bioanode were Proteobacteria, Firmicutes, Bacteroidetes, and Actinobacteria, while the dominant phylum in the biocathode was Proteobacteria. Therefore, this study demonstrates the applicable use of BSCMFC for bioelectricity generation and pollution control.

**Keywords** Activated sludge · Biocathode · Biofilm viability · Power density

## Introduction

Resolving power shortages and pollution are the top challenges facing world developments [1]. The wastewater treatment plants generate large amounts of activated sludge which contains organics compounds at a high concentration (66%) [2]. Activated sludge treatment and disposal have become difficult and expensive problems [3]. Therefore, new approaches are highly required. Among these technologies are microbial fuel cells (MFCs) which are considered as energy transducers consists of anode and cathode converting the stored chemical energy in a variety of biodegradable materials, ranging from pure compounds to multifaceted substrates present in wastewater into electrical energy by tiny electrogens as a biocatalyst to support redox reactions. In the anodic section, the electrogens can oxidize organic

compounds anaerobically producing electrons and protons. Afterward, these electrons pass through an external circuit and combine with the transferred protons into the cathode by diffusion and a terminal electron acceptor usually oxygen producing water [4].

The efficiency of the electric current generation in MFCs is reduced due to overvoltage losses at the cathode electrode [5]. To eliminate this problem, MFC technology uses cathodes based on materials containing catalysts as platinum that actively remove excess electrons [6]. However, its use increases the technology cost. This leads to the search for modern, cheaper, and promising catalysts for cathodic processes [7]. One promising tool for cathodic MFC is the use of microorganisms as biocatalysts [8]. The aerobic biocathodes MFCs have the dominant range of bacterial species in a mixed population belonging to Alphaproteobacteria and Betaproteobacteria [9], Gammaproteobacteria [10], Bacteroidetes [11], and other less well-known groups.

Little studies illustrate the importance of extracellular polymeric substances (EPS) in the biocathode, which have many roles as cell attachment to solid surfaces, biofilm structure maturation, and protection from harsh environmental conditions [12]. EPS are typically made up of proteins, DNA, humic acids, polysaccharides, etc. that are secreted in

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pure and mixed cultures by microbes [12]. The redox properties of these EPS components could be used as an electron carrier in electrochemically active biofilms [13]. EPS are secreted by the bacteria when they form a biofilm on an electrode [14]. Because the EPS layer is not a solid barrier between cells and external electron donors/acceptors, these redox compounds can diffuse through the EPS and play a role in electron transfer between cells and electron donors/acceptors [15].

The wastewater treatment and energy production from artificial or real wastewater provide an overview of the MFC's efficiency with a range of wastewater [16]. Microorganisms naturally grow on many complex compounds found in wastewater. It is necessary to know the microbiome and its biochemical properties, and metabolic activity to obtain successful wastewater bioremediation and electricity generation [17]. To expand the BSCMFC technology scope in the future, keeping in mind the enormous secret bacterial diversity, usually, electricigens in nature, isolation of potentially active ones is necessary. 16SrRNA sequencing was used to study the electroactive involved in the power generation in MFC systems [18]. It explains how the microbial population dynamics is interrelated with the bioreactor that works as a treatment system for wastewater [19].

To our knowledge, no previous studies have been performed comparing the responses of biocathode-isolated electrogenic bacteria to in vitro biofilm formation. Another aspect, there were a few studies in biocathode MFCs. This study aimed to use BSCMFC with aerobic activated sludge as a substrate to increase the rate of ORR and electroactive microbial activities along with the elimination of organic matter through wastewater treatment and electricity generation. The BSCMFC performance was studied by monitoring the power generation, coulombic efficiency, and COD removal efficiency. The anode and cathode potentials were also evaluated. EPSs were extracted from the bioanode and biocathode and their protein and carbohydrate contents were analyzed. In addition, the chemical structure of each EPS fraction was characterized using FTIR. Moreover,

microbial communities were analyzed using SEM and 16SrRNA. In addition, under in vitro conditions, the biofilm forming abilities of each anodic and cathodic culturable bacterial isolate were separately determined and their biochemical characteristics were performed.

## Materials and method

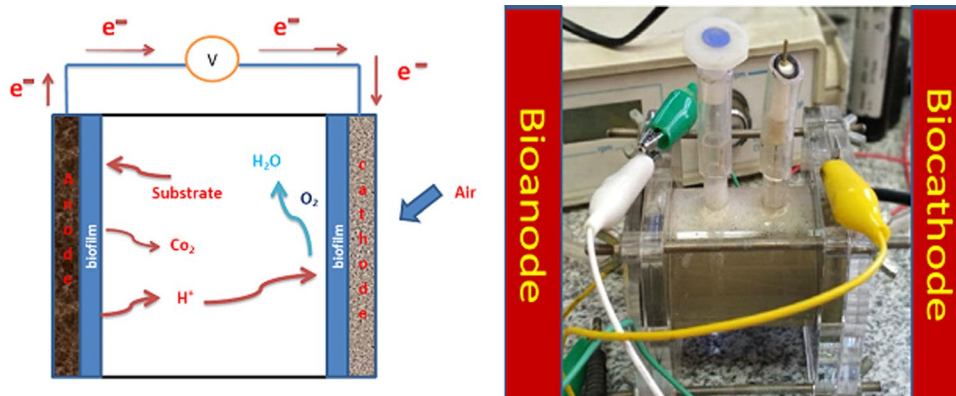
### Chemicals

All supplies and chemical materials were of analytical, and biochemical grade purity. All aqueous solutions were newly prepared throughout the experiments with de-ionized distilled water. Both carbon felt and carbon cloth were purchased from fuel cell store.

### Configuration of BSCMFC

Performing experiments for BSCMFC were achieved in fed-batch mode. MFCs consisting of a cylindrical plexi-glass tube containing the anode and the cathode electrodes, placed on two opposite sides of a liquid chamber reactor (6.0 cm long, 4.6 cm in diameter, 100 mL effective working liquid volume), as shown in Scheme 1 [20]. The anode consisted of an unmodified sheet of three-dimensional carbon felt (not waterproof, project surface area of 0.005288 m<sup>2</sup>) glued to the top of the externally connected anode port. The anode was positioned on the opposite side of the chamber (perpendicularly to the cathode at a distance of 5 cm) and free from the membrane. The gas diffusion carbon cloth electrode was used as the backbone cathode electrode (6 × 6 cm each, cathode coated project area of 0.001663 m<sup>2</sup>). For ORR, the cathode electrode was left open to the air. The current collector (titanium wire) was placed on both sides of the electrodes; this elongated side was used for multimeter connection.

**Scheme 1** Schematic diagram and photo of BCSCMFC



## Microbial populations enumeration, incubational and operational conditions

The biocathode was inoculated with the adapted aerobic activated sludge. The sludge was collected from Zenien wastewater treatment plant located in the Bolaq Aldakrur, Giza, Egypt. This sludge was directly used as an anodic and cathodic inocula. The sample was conveyed to the laboratory in dry plastic water bottles and kept at 4 °C. The sludge was characterized chemically upon arrival as follows (mg L<sup>-1</sup>): total nitrogen, 145.57; total phosphorus, 25.761; total chlorides, 24.85; total solids, 701.6; total dissolved solids, 652.8; total suspended solids, 48.8; biochemical oxygen demand (BOD), 160; COD, 524.16.

Microbiologically by the count plate method, the collected samples have comprised of the determination of the total number of culturable bacteria. The bacterial count was  $5.4 \times 10^7$  CFU mL<sup>-1</sup>. The sludge was initially operated under fed-batch mode of operation (over 1201 h) at  $30 \pm 2$  °C for a total of 50 days to allow a biofilm to form over the anode and the cathode surfaces [21]. The synthetic electrolyte contained CH<sub>3</sub>COONa, 2 g L<sup>-1</sup>; as the degradable organic substrate (electron donors) in a 50 mM phosphate buffer solution (PBS) amended with 12.5 mL micro metal solution and 12.5 mL vitamin solutions. The 50 mM PBS contained (g L<sup>-1</sup>): NaHCO<sub>3</sub>, 2.5; NH<sub>4</sub>Cl, 0.2; KH<sub>2</sub>PO<sub>4</sub>, 13.6; KCl, 0.33; NaCl, 0.3; K<sub>2</sub>HPO<sub>4</sub>, 17.4; CaCl<sub>2</sub>·2H<sub>2</sub>O, 0.15; MgCl<sub>2</sub>, 3.15; yeast extract, 1.0 resulting in an initial COD inlet of  $1472 \pm 17$  mg L<sup>-1</sup>. The batch was replenished with fresh media when the MFC voltage was under 0.1 V to initiate a new cycle MFCs performance in terms of the power generation and substrate degradation was evaluated using the previously reported methods.

## Analytics and computations of biocathode

During the MFCs experiments, the produced cell potentials data between the anode and the cathode were gathered via a multimeter (Lab jack U6-PRO) connected to a laptop by a personal data acquisition system. The multi-cycle technique was used to achieve maximum MFC power densities. The reactors were operated with an external circuit linked through 1000 Ω. The observed voltage was converted to current using the relationship  $I = V/R$ , where  $I$  is the current generated by an applied potential ( $V$ ) calculated at a definite external resistance,  $R$  is 1000 Ω loaded to the cell. Accordingly, the polarization and power curves were plotted using the single cycle technique by recording the pseudo-steady-state voltage peak by reducing the applied external resistance ( $R_{ext}$ ) over a range from 10 MΩ to 330 Ω in reducing order stepwise after the MFC was operated stably. Power density ( $P$ , mW m<sup>-3</sup>) was normalized by the total liquid volume of the MFC.

After stabilization of the open circuit potential (OCP), the polarization was plotted at zero current and the power curves were calculated using Joule's law. The internal resistance ( $R_{int}$ ) was determined using a linear regression related to the ohmic zone on the linear part of the polarization curve [22]. COD concentrations of the influent and effluent were calculated and analyzed according to the APHA procedure. The organic substance concentration is measured as the COD removal efficiency (COD %) using the following equation:  $COD\% = (COD_{initial} - COD_{final}) / COD_{initial} \times 100$  [23]. The Coulombic efficiency ( $C_E$ ) was determined by incorporating the calculated current with the theoretical current based on COD consuming,  $C_E (\%) = (C_p / C_T) \times 100$ . Where  $C_T$  is the theoretical amount of coulombs production.  $C_p$  is the actual current production collected by the anode during one batch cycle. Anode and cathode potentials were evaluated at room temperature ( $25 \pm 1$  °C) using the Voltmaster 6 potentiostat (PST006). Prior to measurements, MFCs was allowed to equilibrate until the open circuit voltage stabilized.

## Biofilm composition analysis

### Attachment assay

At the end of the MFC cycle, the anode and cathode biofilm electrodes were carefully removed from the BSCMFC configuration. To minimize the excess matters, the electrodes were gently washed with sterile distilled water, followed by washing with 0.1 M phosphate buffer saline (PBS, pH 7.0). Biocatalyst density on electrode surfaces was assessed by crystal violet assay. Electrodes were stained for a few seconds with a 1% (w/v) crystal violet stain and solubilized with 70% ethanol. This dye-ethanol solution was quantified at OD<sub>580</sub>. The generated OD value is directly proportional to biofilm thickness [24]. The fresh electrode was used as a cell attachment assay control.

### EPS analysis

For proteins and carbohydrates quantification, the biofilm formed by the microbes on the electrodes was analyzed. Using a sterilized scalpel, the biofilm electrodes were cut into dimensions of  $1 \times 1$  cm<sup>2</sup>. With 0.1 M PBS, the anode and cathode biofilms were washed twice and collected by vortexing then scraping in 1 mL 0.01 M KCl solution. Afterward, the biofilms were centrifuged for 20 min (3000 rpm, 4 °C) after sonication, and the supernatant was collected and filtered via a 0.22 μm membrane filter for EPS analysis. The phenol–sulfuric acid [25] and the Lowry methods [26] were used to quantify the amount of carbohydrate and protein, respectively.

## Biofilm characterization

### FTIR analysis

To confirm the formation of biofilms, samples were prepared by oven drying at 120 °C overnight to remove water on the anodic and cathodic electrodes, followed by grinding the scratched biofilm layer with dry KBr. The powders were analyzed in the Bruker Alpha 11 and recorded on an FTIR spectrophotometer in the region from 400 to 4000  $\text{cm}^{-1}$ .

### SEM analysis

The anodic and the cathodic biofilm surfaces were characterized by SEM (JEOL, JXA-840A) for determining the biofilm formation after 72 days of microbial culture incubation in the BSCMFC. The electrodes were fixed with 2.5% glutaraldehyde, then washed with water three times and dehydrated for 10 min by successively immersing them in ethanol solutions (from 30 to 100%). The specimens were dried, mounted onto specimen stubs using graphite paste, and then covered with gold.

## Analysis of microbial cultures

### Isolation of electroactive

The biofilms were removed from the BSCMFC at the end of operation cycles to enumerate the electroactive bacterial species. The biofilms were washed with PBS, placed in tubes containing PBS, and the bacterial cells removal was performed by sonication. The PBS with the detached bacterial cells was serially diluted and the appropriate dilutions were poured plated on Luria–Bertani (LB) agar plates followed by incubation at 37 °C for 48 h. The plates from the anode and cathode were incubated anaerobically and aerobically, respectively. After incubation, morphologically dissimilar colonies were randomly selected from all plates and purified by streaking.

### Assessment of biofilm formation

The isolated bacteria were tested for biofilm production according to the previously described method [27], in 96-well microtiter plates. The cell density per well was adjusted to be  $1 \times 10^6$  CFU  $\text{mL}^{-1}$  in 200  $\mu\text{L}$  LB medium. After incubation (37 °C, 48 h), the wells were emptied and filled with 200  $\mu\text{L}$  of crystal violet solution (2% v/v). After 5 min, the plates were emptied, washed three times with

distilled water and the dye was extracted by ethanol addition (300  $\mu\text{L}$ ). By a microplate reader, the  $\text{OD}_{600}$  was registered.

### 16SrRNA gene sequencing

Anodic and cathodic single colonies were grown in LB broth medium and incubated at 37 °C for 48 h. Afterward, the bacterial cultures were sent to the biotechnology Lab. (Fac. of Agriculture, Cairo University Research Park, FA-CURP, Egypt) for DNA extraction and PCR amplification. The DNA quantity and purity were measured spectrophotometrically (characterized by  $A_{260}/A_{280}$  nm absorbance ratio). The universal primers used for the PCR amplification were 27 forward primers (5'AGAGTTTGATCCTGG CTCAG3') and 1492 reverse primers (5'GGYTACCTTGTT ACGACTT3'). The amplified genes solution was subjected to agarose gel (1%) electrophoresis, purified and sequenced (MACROGEN sequencing company, Korea). PCR products were directly sequenced in one direction using the respective forward and reverse primers. The obtained nucleotide sequences were aligned using the Basic Local Alignment Search Tool (BLAST) and compared with the sequences in the NCBI database. A phylogenetic tree was constructed using the neighbor-joining model of the MEGA X program. The identified isolates were classified morphologically and biochemically using a standard procedures based on Bergey's manual.

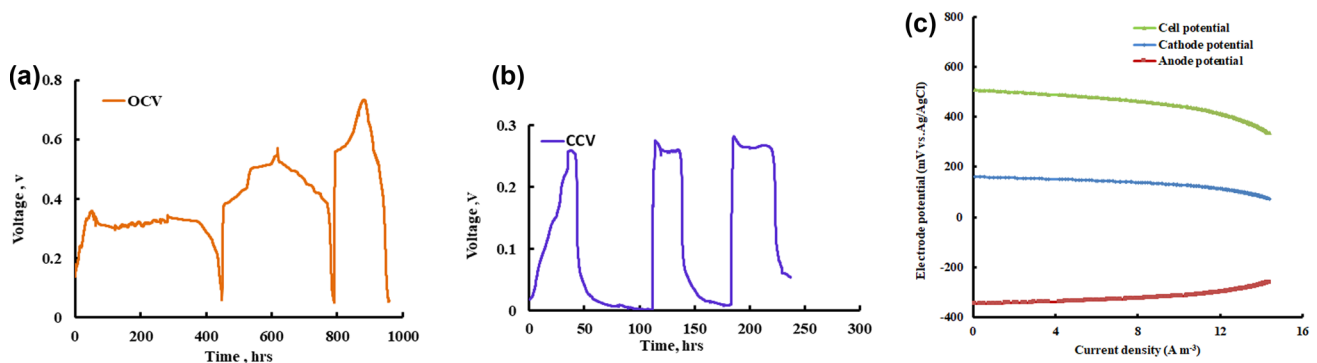
### Statistical analysis

Using the Microsoft Excel 2016 version, statistical analysis was carried out. All parameters, calculations, means, and standard deviations were involved in data analysis.

## Results and discussion

### MFC performance during startup, acclimation, and bioelectricity generation

The open circuit voltage (OCV) changed with time when inoculated with aerobic activated sludge after 3 months' long-term activity of continuous cultivation (Fig. 1a). At first, the maximum OCV was 363 mV (0–7.5 days), then followed by a rapid decrease in OCVs for 2 weeks to approximately 60 mV due to the microbial activity and an increase in substrate consumption. The slow speed of electricity generation can be associated with the lag time of bacterial cell formation on the anode surface at the initial stage of MFC activity [28]. Then, the MFC was inoculated with fresh media to regain the microbial activity for starting the 2nd cycle. These cycles took about (7.5–13 days) for increasing the voltage to 585 mV followed by decreasing the voltage



**Fig. 1** The bioelectrochemical performance of the BSCMFC; **a** time–voltage relationship in BSCMFC **b** the effect of external resistance (with an external resistor of 1k $\Omega$ ) on the voltage output and **c** electrode potentials

gradually according to decreasing the acetate concentration in the media leading to start the third refreshment cycle. During the 3rd cycle (13–50 days), the average voltage output was 755 mV. This result can be due to the development of an electrode biofilm that facilitated the transport of electrons and improved voltage production. The BSCMFC was run for three consecutive rounds and fresh media were introduced to reactivate the microorganisms and promote the growth of anodic biofilm [29]. Therefore, the OCV cycles were indicative of the successful biofilm formation.

Figure 1b shows the effect of 1 k $\Omega$  on the MFC outputs at an initial sodium acetate concentration of 2 gL<sup>-1</sup> under fed-batch mode after cultivating 3 months using the multi-cycle technique. The voltage was initially dropped from 755 mV in OCV to be 259 mV in closed circuit voltage (CCV) which dropped into 50 mV after 112 h. The voltage output of the BSCMFC exhibiting a relatively stable electricity production with the voltage output fluctuating around 282  $\pm$  6.6 mV in each batch after 300 h. These results could be accomplished that the progression of mature microbial communities on both anodic and cathodic electrodes [30]. In addition, the generation of electricity from BSCMFC also indicated the successful reduction process of oxygen that carried out by the formed bioelectrocatalysts on the cathode [31]. Also, the voltage values can be due to the biofilm development on the electrode surfaces during all three operating cycles in both OCV and CCV modes, as well as the bacterial content capacity to generate and transfer the electrons to the anode surface without chemical mediators.

In this stage, the individual electrode potentials of the anode and cathode were kept at a constant level and collected to check the BSCMFC's performance. At steady state, Fig. 1c recorded the biocathodic and bioanodic potentials, as well as the performance of the microbial fuel cell. In comparison to Ag/AgCl, the anode and cathode potentials were at 360 and 173 mV, respectively. As a result, the corresponding OCP vs. Ag/AgCl was around 533 mV, and the

performance of the microbial fuel cell was greatly improved. The variations in the current generation were found to be due to variances in both cathode and anode potential, according to an analysis of the electrode potential curves. Thus, it was essential to develop bioanode and biocathode for efficient BSCMFC.

The COD removal in a medium of initial COD (1472  $\pm$  17 mg L<sup>-1</sup>) increased in the 1st cycle to be 89% and reached to be 72% in the 3rd cycle. In the mature MFC during the long-term operation, the COD removal ratio was 83  $\pm$  8.4%. The COD removal ratio was 83  $\pm$  8.4% which was in line with previously recorded results [20]. They showed that at the end of the MFC operation period, the percentage of COD removal of acetate fed MFCS reached 96%. The difference in COD removal efficiency values could be explained by the various microbial sources (inoculum), the microbial population structures, electrode types, configuration of MFCs, and microbial activity.

Coulombic efficiency ( $C_E$ ), referred to the overall charge ratio transmitted to the anode over the maximum extractable charge after full oxidation of the electricity-converted substrate [32], was 22  $\pm$  2.5% and decreased slowly to 13  $\pm$  5% within 10d in the 4th cycle.

In the case of BSCMFC under fed-batch mode, the anodic biofilm reduced the organic substrate anaerobically into protons and electrons. The organic substrate was converted anaerobically to protons and electrons by the anodic biofilm. The anodic products are then transported into a biocathodic biofilm that may aerobically oxidize oxygen (electron acceptor) into water. After standard anaerobic anodic treatment (i.e. the excess COD from the anode is transferred into the biocathode, which led to significant growth of aerobic heterotrophs with the ability to remove slowly biodegradable COD in aerobic biocathodes), high COD removal efficiency occurred as a further aerobic cathodic step. [33]. According to Haung et al. [34], the optimum output of energy production is determined by the COD initial level concentration.

When the substrate COD concentration was increased, metabolic reaction rates quickened, which could aid to improve MFC efficiency [35]. It can be concluded that the sludge was an effective substrate for electroactive. Furthermore, the bacterial composition in wastewater treatment would be responsible for biodegradability [36].

To assess the performances of the BSCMFC, the power curve and the polarization tests should be achieved after 60 days of stable phase at the end of the operation to evaluate the contribution of the biocathode performance (Fig. 2). The recorded results showed that the power density raised with lowering of the external resistance from 10 M $\Omega$  to 330  $\Omega$ ; a maximum power density of 593 mW m<sup>-3</sup> with the external resistance of 1 k $\Omega$  at a maximum current density of 26,154 mA m<sup>-3</sup>, and the corresponding cell voltage was 244 mV. The internal resistance of the BSCMFC could be determined to be 195  $\Omega$  from the linear fitting of polarization curves. The polarization curve showed that the power was generated by the system considers the principal objective of BSCMFC. In addition, Cai et al. [37] showed that lower external resistance was beneficial for electrogenic bacterial growth and improved MFC efficiency by investigating the impact of external resistance on biofilms. This study reported that the enrichment of microorganisms in the cathode electrode worked as a source of an electroactive catalyst layer during the ORR, so that the aerobic biocathode can help in the improvement of overall BSCMFC electricity output [38]. The findings were compared and indexed in Table 1 with other preceding researches. Xu et al. [39] reported that the power density of SCMFC was 1056  $\pm$  110 mW m<sup>-2</sup> when using a carbon brush that has a high surface area. On contrary, Yang and his colleagues produced a power density of 0.104 mW m<sup>-3</sup> which was lower than our result [40]. The previous findings varied significantly according to

the volume and reactor form, the microorganism's source, and the materials used.

### EPS characterization of biofilms for anodic and cathodic electrodes

At the end of the BSCMFC operation (72 days), the absorbance of the bioanode and biocathode were evaluated to prove the significant growth of microbial biomass on the electrode surfaces. It could be observed that a higher biofilm thickness was observed on the cathode surface (1.422 OD) than that observed on the anode surface (0.952 OD). In addition, the content and composition of EPS are of great importance to the formation of these bacterial biofilms and related to their performance in BSCMFC, because it supports bacterial surface adhesion and cell aggregation. The polysaccharides and proteins contents in the EPS extracted from biocathodic and bioanodic biofilms were quantified. As shown in Fig. 3, The EPS protein content in the biocathode was 0.255 mg protein cm<sup>-2</sup> and with polysaccharides content of 0.347 mg cm<sup>-2</sup>, while in the case of anodic biofilm EPS protein content was measured to be 0.176 mg protein cm<sup>-2</sup> with polysaccharides content (0.229 mg cm<sup>-2</sup>). Thus, the EPS composition as determined from the polysaccharide/protein ratios to be 1.36 and 1.30 for cathodic and anodic biofilms, respectively. It was revealed that the EPS obtained from the aerobic enriched culture had a higher polysaccharide content than the protein level [49]. This implies the presence of an aerobic population on the cathode electrode's surface. This result is consistent with prior research using nitrification reactors. According to Jemaat et al. [50], polysaccharides were the most abundant EPS constituent in the aerobic ammonia-oxidizing bacteria-enriched granules, with polysaccharide/protein ratios frequently more than two [50]. Furthermore, in contrast to earlier research, the EPS of the anaerobic-enriched culture included significantly more polysaccharide in the bioanode, Jiang et al. [51] reported high protein/polysaccharide ratios of EPS from anaerobic ammonia-oxidizing bacteria-dominated sludge. The different methods utilized to extract EPS and the varying protocols used to quantify polysaccharides and proteins could be responsible for the inconsistent findings of several studies [52]. As a result, we can deduce that proteins are a prominent ingredient in anaerobic EPS and that proteins are likely to be responsible for the production of anodic aggregates; on the other hand, the polysaccharide fraction may be a crucial factor in aerobic cathode aggregates.

The biofilm includes polysaccharides, proteins, peptidoglycan, lipids, and nucleic acids. There was a link between the EPS concentration and the biofilm growth [53, 54]. The higher the secretion of EPS by the organism, the greater the biofilm density. More organism number is contained in a denser biofilm than in a lighter one. Therefore, the biofilm

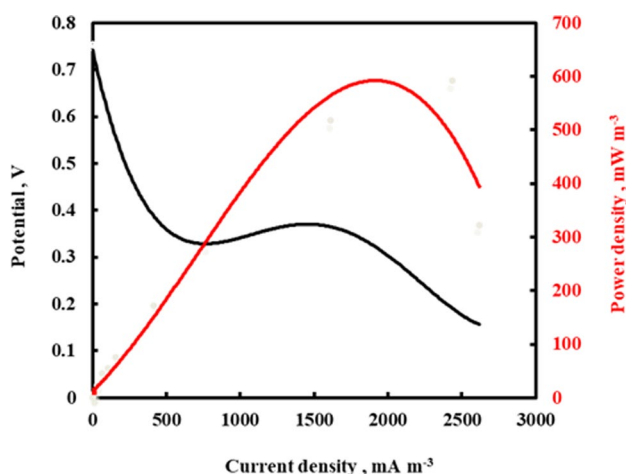


Fig. 2 Polarization and power curves of the BSCMFC reactor

**Table 1** The previous biocathode MFC studies

| MFC design         | Substrate       | Microorganisms                                   | Anode material                | Cathode material  | Max OCV (V) | Max CCV (V) | Max CD ( $\text{Am}^{-3}$ ) | Max PD ( $\text{Wm}^{-3}$ )      | Coulombic efficiency ( $C_E$ %) | Ref        |
|--------------------|-----------------|--|-------------------------------|---|-------------|-------------|-----------------------------|----------------------------------|---------------------------------|------------|
| Single chamber MFC | Sodium acetate  | Aerobic sludge                                   | A sheet of carbon felt        | Carbon cloth with gas diffusion layer   | 0.755       | 0.282       | 2.822                       | 0.796                            | $22 \pm 2.5$                    | This study |
|                    | Sodium acetate  | Clarifier effluent                               | Carbon brush                  | Composed of three layers a diffusion layer, a supporting layer, and a catalyst layer) | 0.55        |             | $\sim 2.5$                  | $1056 \pm 110 \text{ mW m}^{-2}$ | $19 \pm 5$                      | [41]       |
|                    | Sodium acetate  | Aerobic nitrifying sludge                        | Active-carbon felt            | Active-carbon felt  | 0.435       |             | 0.7                         | 0.104                            | $3.5 \pm 0.2$                   | [40]       |
|                    | Lactate         | Anaerobic sludge                                 | Carbon brush                  | Carbon Cloth containing Pt catalyst   | 0.7         | 0.27        |                             | $5.79 \text{ mW m}^{-3}$         |                                 | [42]       |
|                    | Dye waste-water | Mixed bacterial culture                          | Granular activated carbon bed | Granular activated carbon bed   | 0.39        |             |                             | 8                                |                                 | [43]       |
|                    | Sodium acetate  | Mixed liquor from the aerated nitrification tank | Non-wet proof carbon cloth    | Wet-proofed carbon cloth  | 0.561       |             |                             | 0.900                            | 26                              | [44]       |

Table 1 (continued)

| MFC design       | Substrate              | Microorganisms   | Anode material          | Cathode material                           | Max OCV (V) | Max CCV (V) | Max CD ( $\text{Am}^{-3}$ ) | Max PD ( $\text{Wm}^{-3}$ ) | Coulombic efficiency ( $C_E$ %) | Ref  |
|------------------|------------------------|------------------|-------------------------|--|-------------|-------------|-----------------------------|-----------------------------|---------------------------------|------|
| Dual chamber MFC | Diary manure           | Anaerobic sludge | Graphite fiber brush    | Graphite fiber brush                       | 1.04        | 0.502       |                             | $8.15 \pm 0.20$             | $39.8 \pm 1.6$                  | [45] |
|                  | Acetate and propionate | Sewage sludge    | Graphite fiber brush    | Graphite fiber brush                       | 0.94        |             |                             | $13.2 \pm 1.7$              | $40.8 \pm 9.0$                  | [46] |
|                  | Domestic wastewater    | Anaerobic sludge | Plain granular graphite | Plain granular graphite                    | 0.437       | 0.223       | 15.36 mA                    | 19.53                       |                                 | [47] |
|                  | Glucose                | Activated sludge | Graphite granules       | Graphite fiber brush and graphite granules | 0.829       |             | $240.62 \pm 14.60$          | $99.83 \pm 3.72$            | $75 \pm 3$                      | [48] |

density increases relating to the cells age [55]. In conclusion, EPS has an important effect on the bioelectrochemical properties of the electrode surfaces and on biofilm adhesion the BSCMFC.

It is widely known that the protein content of microbial aggregates has a considerably stronger relationship with surface properties such as hydrophobicity and surface charge than the polysaccharide content [56]. While polysaccharide content might supply bacterial cells with a multitude of binding sites, it can also have a significant impact on cell aggregation. In the presence of multivalent ions, the impact of polysaccharides in cell aggregation can be considerably enhanced due to the availability of functional groups such as carboxylates [57].

### FTIR analysis

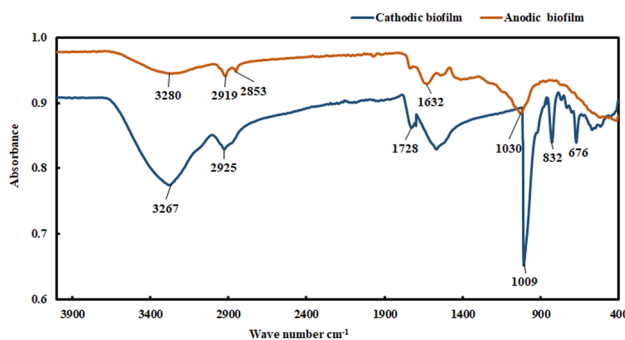
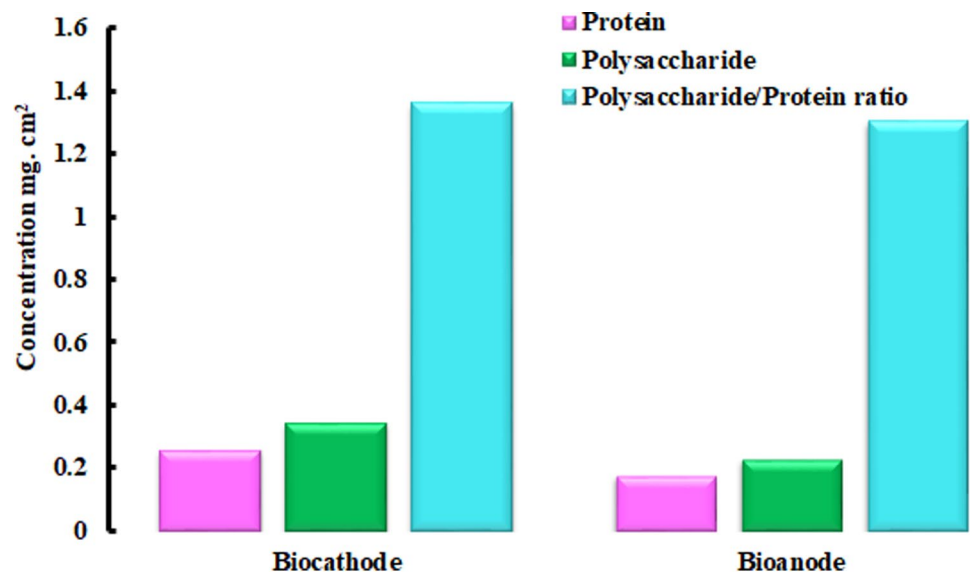
The functional groups derived from proteins, polysaccharides, and amino acids that are involved in EPS were conducted presented using FTIR analysis spectra as indicated in Fig. 4. The  $3900\text{--}400\text{ cm}^{-1}$  region provides the main information regarding the compositions and functionalities of the EPS constituents. The broad band's observed at  $3280$  and  $3267\text{ cm}^{-1}$  are due to the functional groups of O–H and N–H, which can form hydrogen bonds with the carbonyl group of protein peptide bonds [58]. The peaks at  $2925$ ,  $2919$ , and  $2853\text{ cm}^{-1}$  can be attributed to the functional groups of membrane fatty acids and also to certain side chain amino acid vibrations, as the characteristic C–H stretching vibrations of the functional groups –CH<sub>3</sub> and =CH<sub>2</sub> dominate [59]. The observed band at  $1632\text{ cm}^{-1}$  corresponds to the primary amide band ( $1630\text{--}1655\text{ cm}^{-1}$ ) that is associated with the C=O stretching vibrations of peptide groups in proteins Furthermore, the band observed in  $1030$  and  $1009\text{ cm}^{-1}$  is attributable to the symmetric stretching vibration of PO<sub>2</sub> groups in nucleic acid and to C–O–C and C–O–P stretching [60], likely this originating from the damage of bacterial cells during the extraction of EPS. Naumann [60] reported that the presence of carbohydrates and polysaccharides ( $1200$  and  $900\text{ cm}^{-1}$ ) in the cell wall influenced the nucleic acids. The peaks in  $676\text{ cm}^{-1}$  indicated the glycosidic linkage exist between the glycosyl groups [61]. These findings showed the presence of EPS matrix on the electrode surfaces.

### SEM analysis

Bioanode and biocathode surfaces were visualized to determine the relationship between the above-mentioned electrochemical analysis and biofilm coverage. A naked eye revealed that there is a dense rough layer of electroactive on carbon felt and carbon cloth surfaces. Figure 5a, b indicate the fresh structure of the electrodes at



**Fig. 3** Chemical characteristics of EPS from aerobic biocathode and anaerobic bioanode enriched mixed cultures



**Fig. 4** FTIR spectra (3900–400 cm<sup>-1</sup>) of the anodic and cathodic EPS

beginning of the experiment, while Fig. 4c, d depicts the SEM images of the bioanode and biocathode, respectively. It could be observed that the anodic biofilm had a partial and heterogeneous structure with two morphological distinct rod-shaped cells, which was similar to [62]. The morphology variation is assumed to be mainly caused due to the substrate feeding and electrogens, which can either acclimate and colonize with the anodic electrode by secreting matrix or connected together with thick layers of exopolysaccharide using pilus like structures [63]. While SEM image for the biocathode showed that the carbon cloth based biofilm loaded with the dense microbial consortium (Fig. 5d). It can be concluded that the biofilm was intended to consist of aerobes, which acted as a living membrane that prevented or minimized oxygen penetration into the anode. Hence, it can undergo the reduction process of O<sub>2</sub> on the surface of cathode electrode with enhancement of BSCMFC [64]. Moreover, the carbon cloth had a fully accessible surface for microorganisms with a particular surface area and these colonized

microorganisms effectively increased the COD reduction rate and the power production. Therefore, the microbial community in both electrodes supported cooperation between the anaerobic acetate oxidation process on the bioanode and aerobic oxygen reduction on the biocathode for electricity production [65].

### Analysis of microbial cultures

The electroactive bacteria were isolated from the biocathode and bioanode in BSCMFC. The morphologically different isolates were obtained from the biofilm formed on electrodes. Twenty-one isolates were purified.

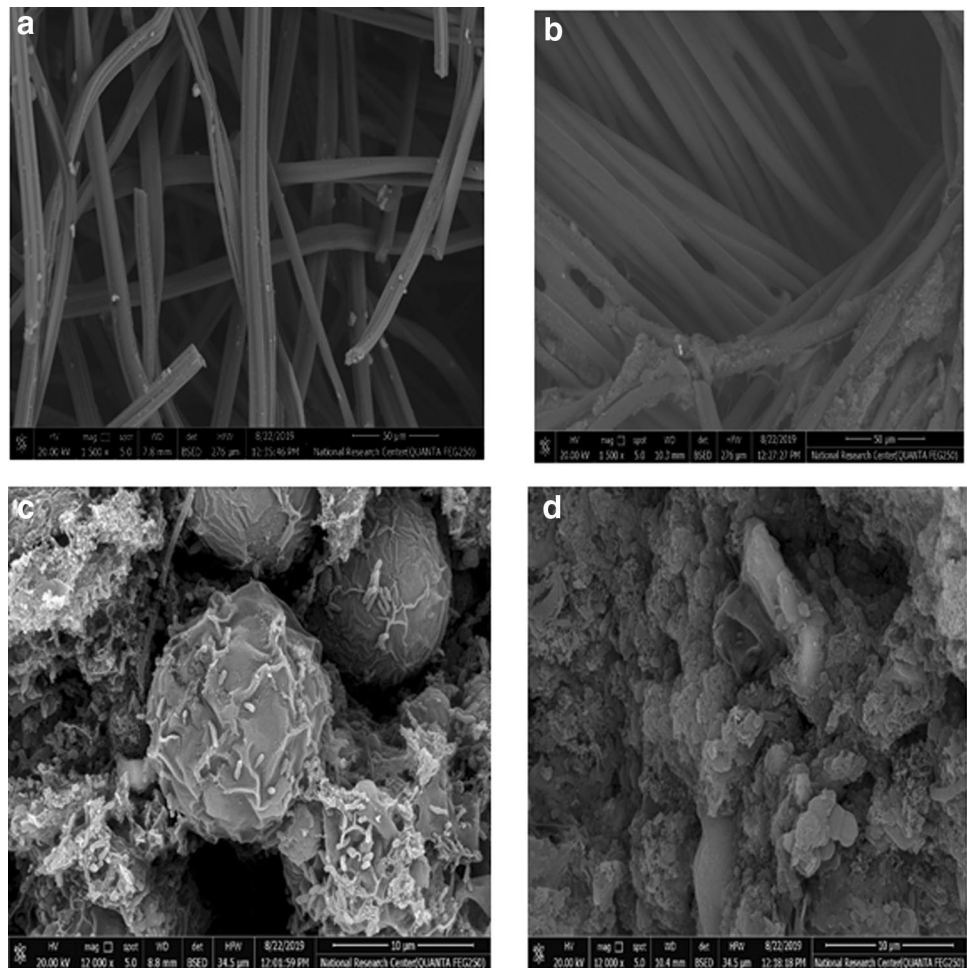
### Biofilm ability

There was limited knowledge of the biofilm formation of the isolates and their significance in bioelectricity production. Table 2 shows that 90.48% and 9.52% of the isolates were strong and moderate biofilm formers, respectively. This result was contradictory with Zafar et al. [66], who reported that 5% of the anodic isolates were strong and 15% were moderate biofilm formers. With the EPS help, biofilm forming bacteria can highly organize themselves on the electrode surfaces and contribute to greater electricity generation in MFCs [67]. The isolated strains would offer the ability to degrade organic substances effectively and to generate electricity in the MFC system.

### Molecular analysis

The 16S rRNA gene sequences for the 13 isolates that had high biofilm formers were performed to classify the compositional difference between the anodic and cathodic

**Fig. 5** SEM images of the anode and cathode electrodes before inoculation with aerobic sludge (**a, b**), SEM image of bioanode (**c**) and biocathode (**d**) after the inoculation of BSCMFC



bacterial populations. The absorbance ratio of the extracted DNA samples was equal or above 1.8 (Table 2). The agarose gel showed the reproducible PCR bands were considered as PCR positive, while isolates of negative PCR findings were regarded as PCR negative. Furthermore, eight isolates with very faint bands were overlooked (Fig. 6a), but the others with reproducible PCR products were amplified and sequenced.

The bioanodic diversity was dominated by Proteobacteria phylum (42.86%), accompanied by phyla Firmicutes (28.57%), Bacteroidetes (14.29%), and Actinobacteria (14.29%) (Fig. 6b) with seven strains: *Kocuria polaris*, *Comamonas nitratorans*, *Alcaligenes faecalis*, *Flavobacterium lindanitolerans*, *Bacillus wiedmannii*, *Bacillus safensis*, and *Pseudomonas pseudoalcaligenes* (Table 3). The majority of these species were facultative anaerobes; this may be consistent with the fact that biofilm strains are limited to species capable of transporting electrons [68]. While cathodic diversity was dominant only by one phylum known as Proteobacteria with six strains: *Pseudomonas putida*, 2 *Pseudomonas monteilii*, *Pseudomonas plecoglossida*, *Achromobacter aegrifaciens*, and *Castellaniella denitrificans*. Table 3

demonstrates the electroactive identification and their accession numbers (MN538880 to MN538892). Some identified percentages of bacteria were found to be 96.74, 96.41, 96.25, 96.14, 95.56, 93.89, 92.99, and 90.82%, suggesting that the electroactives were novel in our MFC. A phylogenetic dendrogram was constructed to clarify the genetic relationship between the electroactives based on 16SrRNA (Fig. 7).

The microbial composition demonstrated the synergistic activity of various functional groups in the BSCMFC to convert acetate into bioelectricity. Bio-electroactives play a role in the electrons transfer by outer membrane cytochromes, pili conductivity, and extracellular excretion between the biofilms and the anolyte electrode [69]. In previous studies, some of our species were known as electroactive. *Bacillus* sp. known to transmit the electrons into the anode through soluble redox-active mediators [70] could be used for bioremediation of sugar industry wastewater and electricity production [71]. Clearly, no MFC had been recorded for *Kocuria*, but *Comamonas* was previously reported [72]. The microbial electrochemical systems had been previously associated with members of the *Flavobacterium* genus, but

**Table 2** Biofilm strength and nucleic acid quantification of the extracted DNA samples of the electroactives

| Isolates no                  | Biofilm formation<br>OD at 570 nm | Nucleic acid quantification |                       |
|------------------------------|-----------------------------------|-----------------------------|-----------------------|
|                              |                                   | Nucleic acid(ng/μL)         | OD <sub>260/280</sub> |
| <b>Bioanodic community</b>   |                                   |                             |                       |
| 1                            | 1.33 ± 0.22 (S)                   | 16.2                        | 1.96                  |
| 2                            | 0.28 ± 0.00 (M)                   | 73.1                        | 1.87                  |
| 3                            | 0.84 ± 0.08 (S)                   | 1091.2                      | 1.93                  |
| 4                            | 0.37 ± 0.02 (S)                   | 175.3                       | 2.00                  |
| 5                            | 0.33 ± 0.03 (S)                   | 1.6                         | 2.35                  |
| 6                            | 0.29 ± 0.01 (S)                   | 338.9                       | 2.10                  |
| 7                            | 0.61 ± 0.17 (S)                   | 604.6                       | 1.88                  |
| 8                            | 0.26 ± 0.07 (M)                   | 16,515.5                    | 1.86                  |
| 9                            | 0.74 ± 0.03 (S)                   | 216.8                       | 1.92                  |
| 10                           | 0.74 ± 0.03 (S)                   | 1089.0                      | 2.10                  |
| 11                           | 0.83 ± 0.04 (S)                   | 55.3                        | 1.63                  |
| 12                           | 1.18 ± 0.24 (S)                   | 421.0                       | 1.97                  |
| <b>Biocathodic community</b> |                                   |                             |                       |
| 13                           | 0.62 ± 0.10 (S)                   | 935.5                       | 1.95                  |
| 14                           | 0.34 ± 0.02 (S)                   | 1089.0                      | 2.10                  |
| 15                           | 1.01 ± 0.18 (S)                   | 1302.3                      | 1.98                  |
| 16                           | 1.42 ± 0.15 (S)                   | 494.6                       | 1.78                  |
| 17                           | 0.32 ± 0.03 (S)                   | 625.4                       | 2.06                  |
| 18                           | 0.88 ± 0.08 (S)                   | 537.3                       | 1.93                  |
| 19                           | 0.78 ± 0.07 (S)                   | 656.1                       | 2.08                  |
| 20                           | 0.35 ± 0.04 (S)                   | 7.8                         | 1.20                  |
| 21                           | 1.09 ± 0.03 (S)                   | 600.3                       | 1.74                  |
| Control                      | 0.07 ± 0.06 (-)                   | –                           | –                     |

Data are mean ± standard deviations

The cut-off (ODC) is defined as the mean OD value of the negative control

Based on the OD, strains are classified as non-biofilm producers (OD ≤ ODC), weak (ODC < OD ≤ 2 × ODC), moderate (2 × ODC < OD ≤ 4 × ODC) or strong biofilm producers (4 × ODC < OD) (Borges et al., 2012)

M moderate; S strong

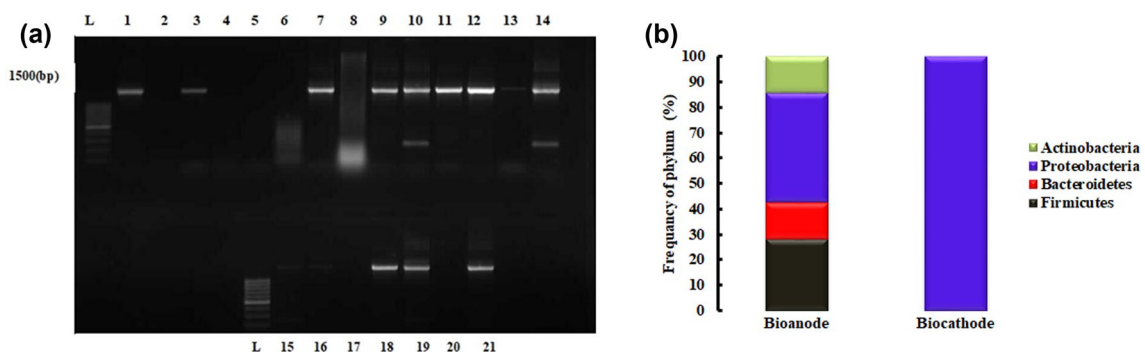
their role remains unknown for its large metabolic ability in current production [73].

*Alcaligenes* genus isolated from the mixed consortium in an MFC [74] participates in the heavy metal intracellular decrease in the MFC cathode chamber [75]. These strains have the unique ability to metabolize ammonium and have nanowires for the possible delivery of electrons to electrode surfaces externally [76]. *Pseudomonas* is another identified bioanode, which can also produce redox mediators which not only enhance its capacity for electrical generation but also provide electron transfer mediators for other microorganisms [72]. In the cathode, *Pseudomonas* can electrochemically reduce oxygen [77]. *P. putida* could be treated the oil refinery wastewater and generate an electric current in an air–cathode chamber [78]. Various species, such as *P. aeruginosa*, *P. fluorescens*, *B. subtilis*, *Shewanella putrefaciens*, and other various species had been documented for cathodic oxygen reduction to water [79]. Further study opens the way for understanding the role of *Castellaniella* and *Achromobacter* in current electrosynthesis production.

Over the numerous mixed-species biocathode, several electrotroths with different metabolic and respiratory capacities remained to be identified [80]. Thus, the exoelectrogens abundance and certain synergetic bacteria could enhance the anodic biofilm oxidation and the oxygen reduction by biocathode without the aid of artificial redox mediators or other catalysts coated on the cathode surface. Based on the 16SrRNA sequence, some genera were novel that requiring further identification and application in MFC. Biofilms can help us unveiling new electrogenic microorganisms and improve our understanding of their role in the functioning of BSCMFC.

**Biochemical characterization**

Table 4 shows the microscopic characteristics, biochemical and sugar fermentation tests for the electroactive. Among



**Fig. 6** a PCR products of extracted DNA molecules on Agarose gel (1%) and b phylum distribution of bacterial community of the bioanode and biocathode

**Table 3** 16SrRNA gene sequences recovered from the NCBI library for electroactives

| Isolate no            | Accession no | Name of bacterium                                   | Per identity (%) | Query cover (%) |
|-----------------------|--------------|---|------------------|-----------------|
| Bioanodic community   |              |   |                  |                 |
| 1                     | MN538880     | <i>Kocuria polaris</i> strain SAKR 1                | 98.20            | 95              |
| 3                     | MN538881     | <i>Comamonas nitratorans</i> strain SAKR 3          | 93.89            | 97              |
| 7                     | MN538882     | <i>Alcaligenes faecalis</i> strain SAKR 7           | 97.56            | 97              |
| 9                     | MN538883     | <i>Flavobacterium lindanitolerans</i> strain SAKR 9 | 92.99            | 98              |
| 10                    | MN538884     | <i>Bacillus wiedmannii</i> strain SAKR 10           | 96.74            | 98              |
| 11                    | MN538885     | <i>Bacillus safensis</i> strain SAKR 11             | 97.26            | 99              |
| 12                    | MN538886     | <i>Pseudomonas pseudoalcaligenes</i> strain SAKR 12 | 97.57            | 99              |
| Biocathodic community |              |   |                  |                 |
| 13                    | MN538887     | <i>Pseudomonas putida</i> strain SAKR 13            | 90.82            | 98              |
| 15                    | MN538888     | <i>Pseudomonas monteilii</i> strain SAKR 15         | 96.25            | 98              |
| 16                    | MN538889     | <i>Pseudomonas monteilii</i> strain SAKR 16         | 96.14            | 99              |
| 18                    | MN538890     | <i>Pseudomonas plecoglossicida</i> strain SAKR 18   | 97.73            | 98              |
| 19                    | MN538891     | <i>Achromobacter aegrifaciens</i> strain SAKR 19    | 96.41            | 99              |
| 21                    | MN538892     | <i>Castellaniella denitrificans</i> strain SAKR 21  | 95.56            | 100             |

these, ten strains were gram negative rods while, the other three strains were gram positive rods and cocci. All electroactive were non-spore former except *B. wiedmannii* and *B. safensis*. The biochemical features revealed significant changes in metabolic capacities. All strains tested for catalase were positive meaning that the electroactives were strict aerobes or facultative anaerobes and most of them were motile. All strains except *P. pseudoalcaligenes* were indole negative. Using the methyl red test, *B. wiedmannii* demonstrated positive acid fermentation. *A. faecalis*, *B. wiedmannii*, and *B. safensis* were Voges Proskauer positive. Eight strains were positive for simmons citrate, while the remaining five strains showed no change.

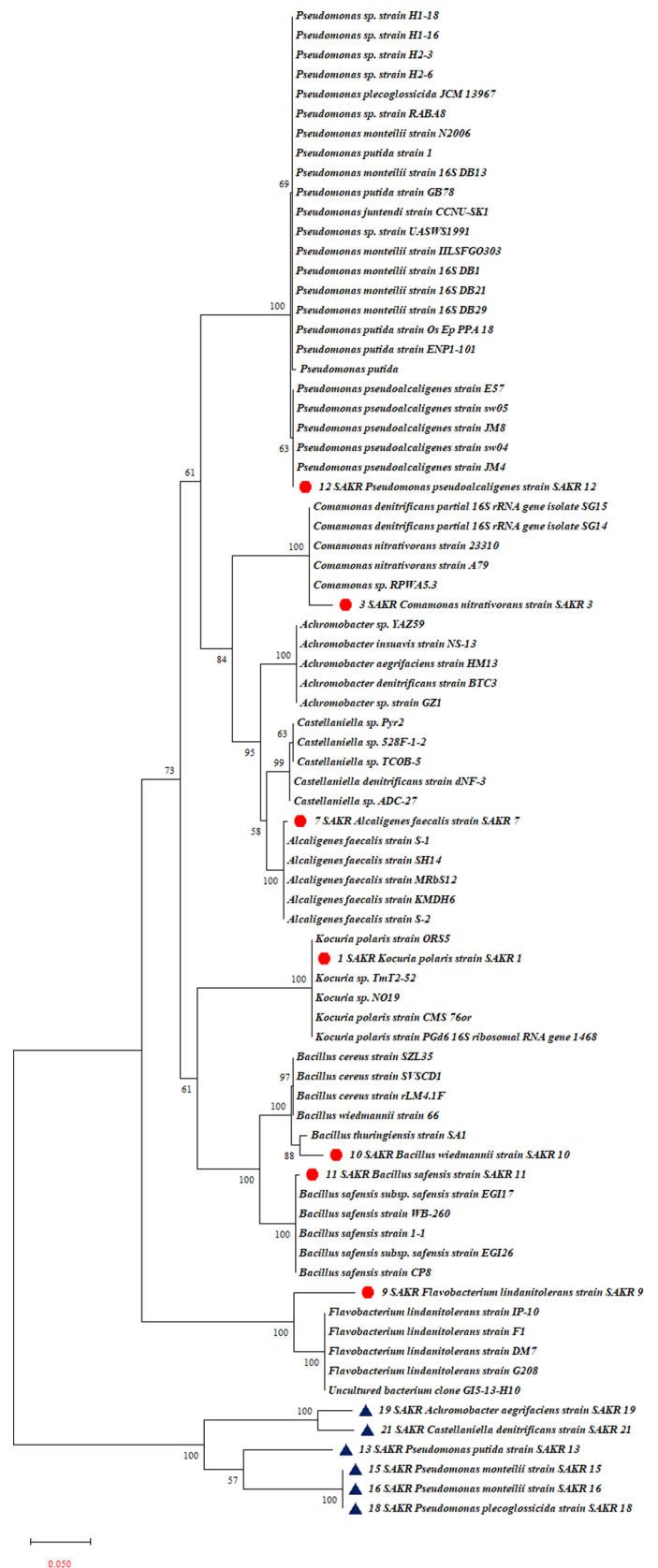
Furthermore, *P. putida* and *A. aegrifaciens* were H<sub>2</sub>S positive and demonstrated that they could minimize their sulfur compound to sulfides, whereas the others were H<sub>2</sub>S negative. In addition, six strains were nitrate positive indicated that these strains could remove nitrogen compounds from wastewater and the others could not reduce NO<sub>3</sub>. *A. aegrifaciens* was urease positive that could degrade the urea and cleaned up waste products into the ammonia. This strain could, therefore, also be used for wastewater treatment with high ammonia content. Additionally, only *B. wiedmannii* and *B. safensis* was able to use the extracellular development of gelatinase enzymes in polypeptides and amino acids. Based on the lipid hydrolysis, two strains had shown positive lipase enzyme, which could be used in bioremediation as an indication of lipolytic activity.

*K. polaris*, *A. faecalis*, and *B. wiedmannii* were positive for starch hydrolysis. Table 4 reveals that four strains had sugar fermentation capacities. It implied that these electroactive did not specifically participate in the pathway of electrode reduction and they were rather breaking down complex sources of carbon that supplied simpler substrates with non-fermenting bacteria. Our results were consistent with Vejarano et al. [81] and Kim et al. [82]; they showed that electrochemically active bacteria had been restricted to simple organic acids as electron donors and based on fermentation products.

## Conclusion

This study demonstrated that the located biofilm on the surface of the cathode enhanced the current response (2615 mA m<sup>-3</sup>) of the BSCMFC and power generation (593 mW m<sup>-3</sup>) with the aid of aerobic exoelectrogens. These exoelectrogens can reduce oxygen without the aid of exogenous mediators. Also, most the electrogenic bacteria were strongly biofilm former in vitro. The 16SrRNA sequencing technology is successfully applied to analyze the microbial community for both bioanode and biocathode. Biofilms can help us unveiling new electrogenic microorganisms and improve our understanding of their role in the functioning of BSCMFC.

**Fig. 7** Phylogenetic relationships between electroactive isolates and 16S rRNA gene sequences retrieved from the GenBank database. The red circle represents the bioanodic dominant population while the blue triangle represents cathodic dominant population (colour figure online)



**Table 4** Biochemical characteristics of electroactives

| S/N                   | Electroactive strains         | Gram | Endospore | Catalase | Motility | Indole production | Methyl-roskauer | Vegetative | Acid production from different carbohydrates |                             |                   |                   |                    |                   |          |            |             |               |          |             |         |         |         |         |        |          |   |   |   |   |   |   |   |   |   |   |
|-----------------------|-------------------------------|------|-----------|----------|----------|-------------------|-----------------|------------|--|-----------------------------|-------------------|-------------------|--------------------|-------------------|----------|------------|-------------|---------------|----------|-------------|---------|---------|---------|---------|--------|----------|---|---|---|---|---|---|---|---|---|---|
|                       |                               |      |           |          |          |                   |                 |            | Citrate utilization                          | H <sub>2</sub> S production | Nitrate reduction | Urease hydrolysis | Gelatin hydrolysis | Starch hydrolysis | D-xylose | L-rhamnose | D-arabinose | Glucofuranose | Fructose | D-Galactose | Mannose | Maltose | Lactose | Sucrose | Starch | Mannitol |   |   |   |   |   |   |   |   |   |   |
| Bioanodic community   |                               |      |           |          |          |                   |                 |            |  |                             |                   |                   |                    |                   |          |            |             |               |          |             |         |         |         |         |        |          |   |   |   |   |   |   |   |   |   |   |
| 1                     | <i>Kocuria polaris</i>        | +    | Cocci     | -        | -        | -                 | -               | -          | +  | -                           | -                 | +                 | +                  | +                 | +        | +          | +           | +             | +        | +           | +       | +       | +       | +       | +      | +        | + | + | + | + | + | - | + |   |   |   |
| 2                     | <i>Com. nitrativorans</i>     | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - |   |   |
| 3                     | <i>Alcal. Fae-calis</i>       | -    | Rod       | -        | +        | -                 | -               | +          | -  | -                           | -                 | +                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | + |   |
| 4                     | <i>Flavo. lindanitolerans</i> | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | - |   |
| 5                     | <i>B. wiedmannii</i>          | +    | Rod       | +        | +        | -                 | +               | +          | -  | +                           | -                 | -                 | +                  | +                 | +        | +          | +           | +             | +        | -           | -       | -       | -       | +       | +      | +        | + | + | + | + | + | + | + | - | - |   |
| 6                     | <i>B. safensis</i>            | +    | Rod       | +        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | +                  | -                 | +        | +          | +           | +             | +        | -           | -       | -       | -       | -       | +      | +        | + | + | + | + | + | + | + | + | + | + |
| 7                     | <i>Ps. pseudocaligenes</i>    | -    | Rod       | -        | +        | +                 | +               | -          | -  | -                           | +                 | +                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | - |   |
| Biocathodic community |                               |      |           |          |          |                   |                 |            |  |                             |                   |                   |                    |                   |          |            |             |               |          |             |         |         |         |         |        |          |   |   |   |   |   |   |   |   |   |   |
| 8                     | <i>Ps. putida</i>             | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | +       | +       | +      | +        | + | + | + | + | + | + | + | + | + | + |
| 9                     | <i>Ps. monotelitii</i>        | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | - | - |
| 10                    | <i>Ps. monotelitii</i>        | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | - | - |
| 11                    | <i>Ps. plecostictida</i>      | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | +       | +       | +      | +        | + | + | + | + | + | + | + | + | + | + |
| 12                    | <i>Achrom. aegrifa-ciens</i>  | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | - | - |
| 13                    | <i>Cast. denitrificans</i>    | -    | Rod       | -        | +        | -                 | -               | -          | -  | -                           | -                 | -                 | -                  | -                 | -        | -          | -           | -             | -        | -           | -       | -       | -       | -       | -      | -        | - | - | - | - | - | - | - | - | - | - |

(-) Negative results  
(+) Positive results

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**Data availability** All data generated or analyzed during this study are included in this published article.

## Declarations

**Conflict of interest** The authors declare no conflict of interest.

**Ethical approval** Not applicable.

**Consent to participate** Yes.

**Consent for publication** Not applicable.

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