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Sustainable Electric Power Generation from Live Anaerobic Digestion of Sugar Industry Effluents Using Microbial Fuel Cells

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Supplementry Information

Experimental Section

 Table S1 Input parameters of each operating condition, where BL - biomass loading, IL

 inoculum loading and EC - electrical conductivity of anolyte

| No | Input parameters | Unit | Output parameters |
|----|------------------|------|---|
| 1 | nН | _ | OCV (mV)/current density (mA m ⁻²)/ |
| Ŧ | pri | | power density (mW m ⁻²) |
| 2 | DI | 0/ | OCV (mV)/current density (mA m ⁻²)/ |
| 2 | Z BL % | | power density (mW m ⁻²) |
| 2 | | 0/ | OCV (mV)/current density (mA m ⁻²)/ |
| 5 | IL. | /0 | power density (mW m ⁻²) |
| Л | | | OCV (mV)/current density (mA m ⁻²)/ |
| 4 | | | power density (mW m ⁻²) |



Fig. S1. The schematic representation of situation explaining the neural networks are adjusted, or trained, so that a particular input leads to a specific target output taken from MATLAB manual



Fig. S2. A schematic layout of the ANN network containing inputs as, **IL** - percentage of inoculum loading, **BL** - percentage of biomass loading, and **EC** - electrical conductivity

16S V3 - V4 metagenome sequencing and analysis

Sequencing methodology

25 ng of DNA was used to amplify 16S rRNA hyper variable region V3 - V4. The reaction includes KAPA HiFi HotStart Ready Mix and 100 nm final concentrations of modified 341F and 785R primers.^a The PCR involved an initial denaturation of 95 °C for 5 min followed by 25

cycles of 95 °C for 30 s, 55 °C for 45 s and 72 °C for 30 s and a final extension at 72 °C for 7 min. The amplicons were purified using Ampure beads to remove unused primers. Additional 8 cycles of PCR were performed using Illumina barcoded adapters to prepare the sequencing libraries. V3 - V4 primers used for the sequencing are shown in **Table S2**.

| Sl. no. | Primer name | Primer sequence 5' - 3' |
|---------|-------------|-------------------------|
| 1 | V3V4F | CCTACGGGNGGCWGCAG |
| 2 | V3V4R | GACTACHVGGGTATCTAATCC |

Table S2 V3 - V4 primers used for sequencing

Sequence data QC

The sequence data was generated using Illumina MiSeq. Data quality was checked using FastQC [http://www.bioinformatics.babraham.ac.uk/projects/fastqc/] and MultiQC software. The data was checked for base call quality distribution, % bases above Q20, Q30, %GC, and sequencing adapter contamination (**Fig. S3**). All the samples have passed QC threshold (Q20 > 95%).



Fig. S3. Histogram of reads with average sequence quality scores

Data analysis

The reads were trimmed (20 bp) from 5' end to remove the degenerate primers. The trimmed reads were processed to remove adapter sequences and low quality bases using Trimgalore. The QC passed reads were imported into mothur and the pairs were aligned with each other to form contigs.^b The contigs were screened for errors and only those between 300 bp and 532 bp were retained. Any contig with ambiguous base calls is rejected. The high quality contigs were checked for identical sequences and duplicates were merged. After this process, the gaps and the overhang at the ends from the contigs were removed and processed for chimera removal which may have formed due to PCR errors. UCHIME algorithm was used to flag contigs with chimeric regions.^c A known reference of all the chimeric sequences was used to identify and remove possible chimeric sequences. The filtered contigs were processed and classified into taxonomical outlines based on the GREENGENES v.13.8 - 99 database.^d The contigs were then clustered into OTUs (Operational Taxonomic Units). After the classification, OTU abundance was estimated. PICRUSt was used to predict gene family abundance.^e PICRUSt program was designed to estimate the gene families contributing to a metagenome by bacteria or archaea identified using 16S rRNA sequencing. The 16S rRNA copy numbers were normalized by PICRUSt's precalculated files. The metagenomes were predicted using predict_metagenomes.py script. The predicted pathways were collapsed into higher categories. OTU contributions for the particular functions were estimated by metagenome contributions.py script. The 16S rRNA metagenome work flow is shown in Fig. **S4**.

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Fig. S4. The workflow of 16S rRNA metagenome

<u>Results</u>

| Parameters | Sugarcane Effluent (SE) |
|------------------------|----------------------------|
| рН | 4.6 |
| Temperature | 32.8 °C |
| Colour | Dark greenish brown |
| COD | 4538 mg L ⁻¹ |
| TSS | 348 mg L ⁻¹ |
| TDS | 882 mg L ⁻¹ |
| TVS | 829 mg L ⁻¹ |
| Total carbohydrates | 18 μg mL ⁻¹ |
| Total proteins | 80 μg mL ⁻¹ |
| C:N | 16:3 |
| Conductivity | 10 mS cm ⁻¹ |

 Table S3 Physico-chemical characterization of sugarcane effluent before pretreatment

Table S4 Fitted Nyquist parameters and total internal resistance of the fabricated MFCs with

 different ionic conductivity

| Systems | R _s (Ω cm²) | R _c (Ω cm²) | R _a (Ω cm²) | IR (Ω cm²) |
|-------------------------------|---------------------------|---------------------------|---------------------------|------------------------|
| SE (10 mS cm ⁻¹) | 7.47 × 10 ² | 1.83 × 10 ³ | 6.29 × 10 ² | 3.20 × 10 ³ |
| SE (20 mS cm ⁻¹) | 6.30×10^{1} | 1.03 × 10 ³ | 4.86 × 10 ² | 1.58 × 10 ³ |
| SE (30 mS cm ⁻¹) | 1.80×10^{1} | 9.76 × 10 ² | 3.00×10^{1} | 1.02 × 10 ³ |
| SE (40 mS cm ⁻¹) | 7.30×10^{1} | 2.52 × 10 ³ | 3.70×10^{1} | 2.63 × 10 ³ |
| SE (50 mS cm ⁻¹) | 3.01×10^{2} | 4.37 × 10 ³ | 1.52 × 10 ² | 4.82 × 10 ³ |



Fig. S5. (**A**) Cell voltage, (**B**) current density and (**C**) power density curve of MFCs fabricated with varying biomass loading with the course of time. Error bars represent standard deviation of four independent experiments.



Fig. S6. (**A**) Cell voltage, (**B**) current density and (**C**) power density curve of MFCs fabricated with varying pH with the course of time. Error bars represent standard deviation of four independent experiments.



Fig. S7. (**A**) Cell voltage, (**B**) current density and (**C**) power density curve of MFCs fabricated with varying ionic conductivity with the course of time. Error bars represent standard deviation of four independent experiments.



Fig. S8. (A) Fitted results of Nyquist plots of MFCs with varying ionic conductivity (-0-)10 mS cm⁻¹, (- Δ -) 20 mS cm⁻¹ (- \Box -) 30 mS cm⁻¹ (- \bullet -) 40 mS cm⁻¹ (- Δ -) 50 mS cm⁻¹ and Equivalent circuit used for fitting the nyquist plots of MFCs with varying ionic conductivity

ANN modelling and its validation:

The effect of interaction among the four different variables on the enhancement of MFC performance is studied using ANN. The Design of Experiments (DOE; Design expert, version 7.0.0, USA) is used for the statistical modelling of MFC.^f The experimental design consists of 54 runs with two replicas each and is shown in **Table S5.**

| | | Input | variables | Output variables | | | |
|-----|---------------------------|----------------------------|-----------|--|-------------------------|---|--|
| | Variable | Variable | Variable | Variable / | Response | Response | Response |
| Run | 1 | 2 | 3 | Valiable 4 | 1 | 2 | 3 |
| | Biomass Ioading (%) | Inoculum Ioading (%) | рН | Conductivity (mS cm ⁻¹) | Cell voltage (mV) | Current density (mA cm ⁻ ²) | Power density (mW m ⁻ 2) |
| 1 | 30 | 30 | 7 | 20 | 743 | 6.45 | 4792 |
| 2 | 30 | 30 | 8 | 20 | 728 | 6.30 | 4586 |
| 3 | 30 | 30 | 9 | 20 | 741 | 6.51 | 4824 |
| 4 | 30 | 30 | 7 | 25 | 759 | 6.53 | 4956 |
| 5 | 30 | 30 | 8 | 25 | 765 | 6.62 | 5064 |
| 6 | 30 | 30 | 9 | 25 | 745 | 6.44 | 4798 |
| 7 | 30 | 30 | 7 | 30 | 781 | 6.80 | 5312 |
| 8 | 30 | 30 | 8 | 30 | 772 | 6.58 | 5079 |
| 9 | 30 | 30 | 9 | 30 | 764 | 6.86 | 5241 |
| 10 | 30 | 35 | 7 | 20 | 707 | 6.22 | 4397 |
| 11 | 30 | 35 | 8 | 20 | 704 | 6.25 | 4400 |
| 12 | 30 | 35 | 9 | 20 | 698 | 6.23 | 4348 |
| 13 | 30 | 35 | 7 | 25 | 709 | 5.99 | 4247 |
| 14 | 30 | 35 | 8 | 25 | 707 | 6.31 | 4461 |
| 15 | 30 | 35 | 9 | 25 | 719 | 6.44 | 4630 |
| 16 | 30 | 35 | 7 | 30 | 718 | 6.32 | 4537 |
| 17 | 30 | 35 | 8 | 30 | 729 | 6.23 | 4541 |
| 18 | 30 | 35 | 9 | 30 | 747 | 6.52 | 4870 |

Table S5 DOE for ANN modeling using four input variable and its outputs

| 19 | 30 | 40 | 7 | 20 | 528 | 4.16 | 2196 |
|----|----|----|---|----|-----|------|------|
| 20 | 30 | 40 | 8 | 20 | 555 | 4.78 | 2652 |
| 21 | 30 | 40 | 9 | 20 | 516 | 4.61 | 2378 |
| 22 | 30 | 40 | 7 | 25 | 606 | 4.64 | 2811 |
| 23 | 30 | 40 | 8 | 25 | 603 | 5.19 | 3120 |
| 24 | 30 | 40 | 9 | 25 | 647 | 5.77 | 3733 |
| 25 | 30 | 40 | 7 | 30 | 621 | 5.87 | 3645 |
| 26 | 30 | 40 | 8 | 30 | 660 | 6.10 | 4026 |
| 27 | 30 | 40 | 9 | 30 | 654 | 6.48 | 4237 |
| 28 | 50 | 30 | 7 | 20 | 666 | 4.21 | 2803 |
| 29 | 50 | 30 | 8 | 20 | 699 | 6.11 | 4270 |
| 30 | 50 | 30 | 9 | 20 | 729 | 6.41 | 4673 |
| 31 | 50 | 30 | 7 | 25 | 725 | 4.21 | 3052 |
| 32 | 50 | 30 | 8 | 25 | 717 | 6.30 | 4517 |
| 33 | 50 | 30 | 9 | 25 | 738 | 5.98 | 4413 |
| 34 | 50 | 30 | 7 | 30 | 732 | 4.29 | 3140 |
| 35 | 50 | 30 | 8 | 30 | 733 | 6.75 | 4947 |
| 36 | 50 | 30 | 9 | 30 | 925 | 7.83 | 7242 |
| 37 | 50 | 35 | 7 | 20 | 747 | 4.98 | 3720 |
| 38 | 50 | 35 | 8 | 20 | 727 | 6.82 | 4958 |
| 39 | 50 | 35 | 9 | 20 | 751 | 7.16 | 5377 |
| 40 | 50 | 35 | 7 | 25 | 731 | 5.67 | 4144 |
| 41 | 50 | 35 | 8 | 25 | 735 | 5.87 | 4314 |
| 42 | 50 | 35 | 9 | 25 | 759 | 6.88 | 5222 |
| 43 | 50 | 35 | 7 | 30 | 734 | 6.08 | 4462 |
| 44 | 50 | 35 | 8 | 30 | 745 | 5.16 | 3844 |
| 45 | 50 | 35 | 9 | 30 | 776 | 7.84 | 6084 |
| 46 | 50 | 40 | 7 | 20 | 502 | 4.53 | 2274 |
| 47 | 50 | 40 | 8 | 20 | 463 | 5.60 | 2592 |
| 48 | 50 | 40 | 9 | 20 | 451 | 6.93 | 3125 |
| 49 | 50 | 40 | 7 | 25 | 496 | 4.89 | 2425 |

| 50 | 50 | 40 | 8 | 25 | 510 | 5.86 | 2988 |
|----|----|----|---|----|-----|------|------|
| 51 | 50 | 40 | 9 | 25 | 536 | 6.17 | 3308 |
| 52 | 50 | 40 | 7 | 30 | 522 | 4.98 | 2599 |
| 53 | 50 | 40 | 8 | 30 | 534 | 6.20 | 3310 |
| 54 | 50 | 40 | 9 | 30 | 547 | 6.00 | 3282 |

According to DOE matrix, a combination of 50% of biomass loading, 30% of inoculum loading, ionic conductivity of 30 mS cm⁻¹ and pH of 9 is optimum for maximal MFC performance. Further, the concurrent effect of the variables on biofilm formation is determined using ANN and to correlate the power generation efficiency in the sugarcane effluent based MFC using anaerobic inoculum. Four input variables viz, Biomass loading, % of Inoculum added, pH, and ionic conductivity are used to feed forward neural network. Three output variables: cell voltage (mV), current density (mA cm⁻²) and power density (mW m⁻²) are used for validation. The MLP layer consists of an input layer of N neurons, a hidden layer of M neurons and an output layer of K neurons, the relationship between different layers can be described by equation (1),

$$g_k = F\left[\sum_{j=1}^M W_{kj} f\left(\sum_{i=1}^N w_{ji} x_i + \theta_j\right) + b_k\right]_{\dots,\dots,(1)}$$

where, j = 1, 2...M; i = 1,2...M and k = 1,2...K. In equation 1, w_{ji} is the weight connecting i^{th} neuron in the input layer to the j^{th} neuron in the hidden layer, θ_j is the bias of the j^{th} neuron in the hidden layer. Similarly, W_{kj} is the weight connecting the j^{th} neuron in the hidden layer to the k^{th} neuron in the output layer and b_k is the bias of the k^{th} neuron in the output layer.

Optimization of the topology of the ANN is the first crucial step. Of the various architectures investigated, one hidden layer with ten neurons is considered as the optimum configuration, which is based upon the regression values obtained. The dataset contains 216 input/output patterns for fitting. Out of these, 152 samples (70%) are taken for training and 32 samples (15%) each is used for validation and testing. The training data are the biggest set of data used by neural network to learn the pattern presented in the data by updating the network weights. Diagram of neural network used for the present study is shown in **Fig. S9**.



Fig. S9. Diagram of neural network used for the present study

The testing data is used for the evaluation of the quality of the network. The final performance and generalization ability of the trained network are carried out using validation data. To determine the optimum number of hidden nodes, various topologies are used, where the number of nodes in the hidden layer is varied from 2 to 12. The error function used in the present study is Root Mean Square Error (RMSE). Based on the minimum value of RMSE, we used 10 neurons in the hidden layer for the present study (**Fig. S10**). The best regression curve obtained using 10 neurons in the hidden layer are shown in **Fig. S11**.



Fig. S10. The graph showing the relationship between number of neurons and root mean square error (RMSE)



Fig. S11. Training, validation, and testing mean square error for the Levenberg – Marquardt algorithm with the best validation performance of 1170.8423 at epoch 8

The correlation coefficients for training, validation and testing for the data used in the present ANN model are 0.96063, 0.97251, and 0.96001, respectively are shown in **Fig. S12**. The training is stopped after 14 epochs as shown in **Fig. S11**, where the best performance is obtained for epoch 8 with a value of 1170.84. For the best fit, the data should fall along a 45 degree line, where the network outputs are equal to the experimental outputs. It evident from the figure that, almost all the data points fall on this line, except a few data points. **Fig. S13** shows the error histogram of the present work. The blue bars in the figure represent training data, the green bars represent validation data and the red bars represent the testing data. It is observed that, most of the errors fall between -21 and 21. But some of the training data points are outside this range. These further confirm that, neural network model can effectively reproduce the experimental results.



Fig. S12. Training, validation, and testing regression for the Levenberg–Marquardt algorithm



Fig. S13. The error histogram with 20 bins showing that most of the errors falls between - 21 and 21



Fig. S14. Trend of open cell voltage with time of (-0-) MFC - 1, (- Δ -) MFC - 2 and (- \Box -) MFC-3. Error bars represent standard deviation of four independent experiments.



Fig. S15. CLSM images of anode in different anolyte of (A1 and A2) - MFC - 1, (B1 and B2) - MFC - 2 and (C1 and C2) - MFC - 3



Fig. S16. Graphical representation of the COD removal efficiency and Columbic efficiency of fabricated MFCs



Fig. S17. (**A**) Histogram representing contig length distribution and (**B**) Rarefaction curve shows the measure of diversity that has been captured by a given number of reads in a sample



Fig. S18. Alpha diversity indices showing the richness in bacterial diversity with Chao 1 (318), ACE (329), Shannon (3.49), Simpson (0.92), InvSimpson (13.03) and Fisher (58.08)



Fig. S19. Frequency of top 10 genus that present in the bacterial community of sugarcane effluent based Microbial Fuel Cell



Fig. S20. Frequency of top 10 species that present in the bacterial community of sugarcane effluent based Microbial Fuel Cell

| Poagonto | Quantity | Annual |
|----------------------------------|----------------------|------------|
| Reagents | needed | cost (USD) |
| Fe ₂ TiO ₅ | 2 g L ⁻¹ | 0.01 |
| NaOH | 5% | 0.04 |
| HCI | 3% | 0.04 |
| SnCl ₂ | 10 g L ⁻¹ | 0.09 |
| PdCl ₂ | 10 g L ⁻¹ | 1.56 |
| NiSO ₄ | 30 g L ⁻¹ | 0.18 |
| $C_4H_6O_4$ | 25 g L ⁻¹ | 0.14 |
| $NaPO_2H_2$ | 25 g L ⁻¹ | 0.03 |
| Total | | 2.01 |

Table S6 Reagent cost for the development of anode materials

| Items | Remark | Unit cost (USD) | Annual cost (USD) | |
|--------------------|-------------------|--------------------|----------------------|--|
| Costs of routine | Once in a month | 7.02 | 84.28 | |
| maintenance | (12 times a year) | 7.02 | | |
| Costs of water | Once in a month | 7.02 | 01 70 | |
| quality monitoring | (12 times a year) | 7.02 | 04.20 | |
| Total | | | 168.56 | |

Table S7 Cost of operating and maintenance other than reagents

Table S8 Reagent cost for catholyte

| Reagents | Quantity needed | Annual Cost (USD) |
|----------------------------------|------------------------|-------------------|
| NaH ₂ PO ₄ | 4.4 g L ⁻¹ | 0.04 |
| Na ₂ HPO ₄ | 3.4 g L ⁻¹ | 0.04 |
| $K_3[Fe(CN)_6]$ | 16.6 g L ⁻¹ | 0.95 |
| Total | | 1.04 |

Table S9 Cost of other accessories

| Items | Cost (USD) |
|---|------------|
| Double - chambered plexiglass (40.5 mL) | 11.24 |
| Mild steel (Rs. 660/m ²) | 0.42 |
| Nafion 117 (10 × 10 cm) | 15.11 |
| Gaskets (Rs. 1000/m ²) | 0.70 |
| Stainless steel | 0.79 |
| Total | 28.26 |

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