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## Isolation and Identification of Bacteria's from Cattle Dung used in Microbial Fuel Cells to Generate Bioelectricity

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**Abstract:** In recent times, recycling of biomass based organic waste has emerged as very vital aspect of wastes management. Cattle dung is extensively studied for its use as bio agriculture fertilizers and widely explored as prominent, potential and alternative fuel. The unique characteristics to carry diversified microbes in dung are exploited to use it as substrate in recently developed novel technology of Microbial Fuel Cells (MFC). In this study an inexpensive lab scale H-shaped double chamber MFC consisting of two chambers separated by salt bridge was fabricated. Role of cattle dung for biological utilization based bacteria to make biofilm and generate energy in microbial fuel cell was observed. The research studies are able to enumerate Total Viable Count in the range of 1.9 X10<sup>6</sup> to 2.8 X10<sup>6</sup> cfu/gram of dung sample and subsequently primarily nine isolates Bacillus subtilis, Escherichia coli, Streptococcus spp, Pseudomonas aeruginosa, Clostridium Spp, Peptostreptococcus Spp, Bacillus Cereus, Klebsiella Spp, Bacteroides Species. While examining bio film majority of bacteria were of anaerobic nature which contribute substantially in bioelectricity generation. Undoubtedly, the exploitation of cattle dung bacterial community may contribute substantially in sustainability of energy generation in

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MFC, but to identify electrifying bacteria and thoroughly understanding the bio mechanism with the help of recent advances in research in engineering and biotechnology is a key of success.

**Keywords:** bacteria, biomass, biofilm, bioelectricity cattle dung, MFC

**1. Introduction:** Energy consumption across the globe has increased exponentially in recent past and to meet the ever increasing energy demand, there is a dire need to identify more and all feasible sources of energy (Gagandeep: 2017). Rigorous use of fossil fuels to meet demand has posed a threat to life with secondary effects of global warming and environmental pollution (Lovely:2006, Rahimnejad: 2015). In recent times, recycling of biomass based organic waste has emerged as very vital aspect of wastes management. Many of such wastes like municipal solid waste, food wastes, night soils and livestock manures such as cattle dung, sheep, goat dung, poultry drooping and plant wastes that can be very easily managed through biological methods. Cattle dung is the organic biomass mixture of semi-digested and undigested residues of digested matter excreted through the cow's digestive system from bovine's animal species. Cattle dung is extensively studied for its use as bio agriculture fertilizers and widely explored as prominent, potential and alternative fuel in the form of biogas with high methane values. Cow dung carries a wide diversity of microorganism containing different species of bacteria like Bacillus spp., Corynebacterium spp. and Lactobacillus spp., Citrobacter, Enterobacter, Escherichia coli, Klebsiella spp, Pasteurella spp along with protozoa and yeast (Nene: 1999, Kartikey: 2016). Versatility of microorganisms in dung has encouraged the researchers to seek alternative sources for energy by utilizing traditional biomass sources by employing modern tools of technologies developed recently. As a consequence of these efforts, one of the recently proposed alternatives is energy derived from fuel cells utilising presently existing biomasses. Microbial Fuel Cells (MFC) are novel devices that use bacterial community as the catalyst for the oxidation of organic or inorganic matter and lead to generate current (Logan: 2006). Therefore a bioelectricity is generated in MFC between bacterial metabolic due to

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development of biopotential (Heilmann: 2006). A bio-potential developed between the bacterial metabolic and these conditions leads to generate a bioelectricity in MFCs. Anaerobic conditions are necessary in anode chamber as oxygen will hinder the production of electricity (Davis: 1962, Rahimnejad: 2015). In present lab scale studies, an inexpensive and widely used configuration of lab scale H-shaped double chamber MFC consisting of two chambers separated by salt bridge have been assembled as shown in Figure 1. These H-

shaped are widely acceptable for examining power production with different materials,

microbes and mediators.

Research on role of cattle dung as a source of renewable energy subjected to optimum biological utilization based bacteria to make biofilm and generate energy in microbial fuel cell is on full swing. Our objective for this study is to isolate, identify and characterized the bacteria from cattle dung substrate with different morphological and biochemical basis and to study their usefulness in the MFC to make bio-film and subsequently generate electricity.

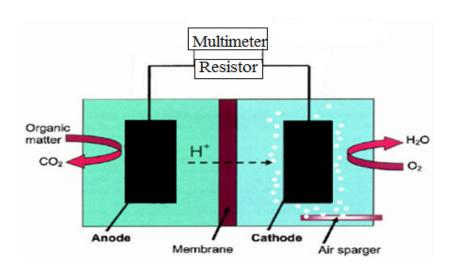


Figure 1: Schematic of Basic Components of Double Chamber MFC

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**2. Material and Methods:** Cattle dung is naturally occurring rich organic matter having different bacterial colonies. In the present study, the random samples of recently excreted fresh cattle dung samples were collected from three different locations. All animals were routinely fed with locally available green fodder along with the concentrate ration.

Cattle dung is the main substrate used in the MFC along with the sterile distilled water or other medium. In MFC anodic chamber under anaerobic environment with some of the specific group of bacteria may increase in number and make a bio film on anode electrode and enhance power generating capabilities of cells by many folds. Therefore, enumeration of the bacteria in cattle dung substrate before being utilized in MFC, and then after removing bio-film from anode electrode.

**2.1 Enumeration of Bacterial Population in Samples:** Enumeration of bacterial population was determined by the serial dilution methods which is developed by (Frazier:1995, **Talaro: 2009).** For the microbiological analysis of dung, fresh cattle dung samples are homogenized thoroughly. For the preparation of samples, one gram of fresh dung, out of the collected dung is taken and mixed with 100 ml distilled water in beaker. The homogenized dung sample is marked as 1A and serial diluted in seven tubes with distilled water up to 10<sup>-7</sup>. Further step, 0.1 ml of solution is taken from tube 1 and dropped on Petri plate already having the Nutrient Agar for growth of bacteria. Later the solution is made to spread on Petri plate and plate is ready to place in aerobic/ anaerobic conditions and so on for other. After incubation, the plates been analysed under microscope and colonies were counted with multiplier 10<sup>1</sup>. Similarly plate 2, had been analyzed and counted colonies were reported with multiplier  $10^2$  & so on. After that, colony forming units (cfu) are counted as total viable count TVC. Standard plate count method is used for culturing as well as calculating total viable count (TVC). The same serial dilution process is repeated for sample 1 B. The average of two TVC is taken as result. Again fresh cattle dung is collected from another two regions and marked it as sample 2 and sample 3 and same process of TVC determination is applied.

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2.2 Isolation of Bacterial Colonies from the Cow Dung and Electrode Biofilm: During

incubation, variety of colonies observed on the cultures plates are selected and purified with

the help of selective media like brain heart infusion, eosin methylene blue agar, brilliant

green agar and Maconkey lactose agar (MLA) . The different bacterial species isolated from

fresh dung are determined by standard bacteriological identification methods according to

Bergey's manual (7th edition). Morphological features like gram staining, shape and motility

are examined. By using broth culture, various biochemical test are conducted which mainly

includes Catalyze, Oxidase, Triple Sugar Iron, Sugar fermentation, IMViC series which

consist of four definite tests i.e Indole production, methyl red, Voges Proskauer and Citrate

utilization (Talaro: 2009).

3. Results and Discussion: Cattle dung is a special type of organic matter which is quite

rich with bacteria. Under specific anaerobic conditions these bacteria grow in millions and

help to act as active catalyst to increase the pace of chemical reactions. These reactions

improve the power generation capability of cells. The conditions for enough microbial

growth are maintenance of anaerobic environment, sufficient substrate and required pH

value. The microbiological analysis of dung samples mainly includes total bacterial count

which is necessary in MFC operation for transfer of electrons to electrode. Bacteria attach to

the surface of the anode act as biocatalysts to pull electrons from substrate. The results of

total viable count of bacteria, isolation and identification of bacteria are discussed below.

**3.1 Total Viable Count of Bacteria**: Total viable count (TVC) is an important parameter to

evaluate the quality of dung to be used as manure as well as bio-energy source. Enumeration

of microbial loads has been calculated as colony forming units (cfu/gram) of all the 3 dung

samples. Each sample has been serially diluted in duplicate to calculate TVC and results are

shown in table 1. Result shows that, average values of TVC are 2.8 X10<sup>6</sup> cfu/gram, 1.9 X10<sup>6</sup>

cfu/gram and 2.5 X10<sup>7</sup> cfu/gram in sample one, two and three respectively. From these

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plated 15 distinct colonies were further streaked onto the general as well as selective culture medium to get the pure and isolated colony (Teo: 2011).

**Table 1: Total Bacterial Count** 

Biomass	TVC A	TVC B	Average TVC		
	(cfu/gm)	(cfu/gm)	(cfu/gm)		
Cattle Dung Sample 1	$3.4 \times 10^{6}$	$2.2 \times 10^{6}$	$2.8 \times 10^{6}$		
Cattle Dung Sample 2	1.5 ×10 <sup>6</sup>	$2.3 \times 10^{6}$	1.9 ×10 <sup>6</sup>		
Cattle Dung Sample 3	$2.9 \times 10^7$	$2.1\times10^{7}$	$2.5 \times 10^7$		

**3.2 Isolation and Identification of Isolates:** Bacterial isolates (from the cattle dung as well as from the biofilm of anodic chamber) are classified on the basis of their features in different forms of species. To enhance the power generating capabilities of MFC, identification of power developing bacteria is significant. Identification has helped in addition of particular type of bacteria as mediator for augment of power density.

**3.2.1 Morphological Features of Bacterial Isolates:** In the present study, 15 bacterial species have been isolated from the dung samples. Only 9 (B1-B9) isolates have been selected for study and remaining six isolates are ignored being similar in morphological features. The detailed morphological characteristics of nine isolates mentioned in table 2 shows the distinct morphological features of bacterial isolates. The features include bacterial colony characteristics of colony shape, colour, size, margin, appearance, colour pigmentations, gram staining reactions, bacterial morphology, spore staining and motility. Some isolated bacteria colonies and their microscopic views of gram staining are shown in

figure 2. The results of bacterial colony characteristics of this study are comparable to (**Dhadse: 2012,** and **Yuti: 2013).** 

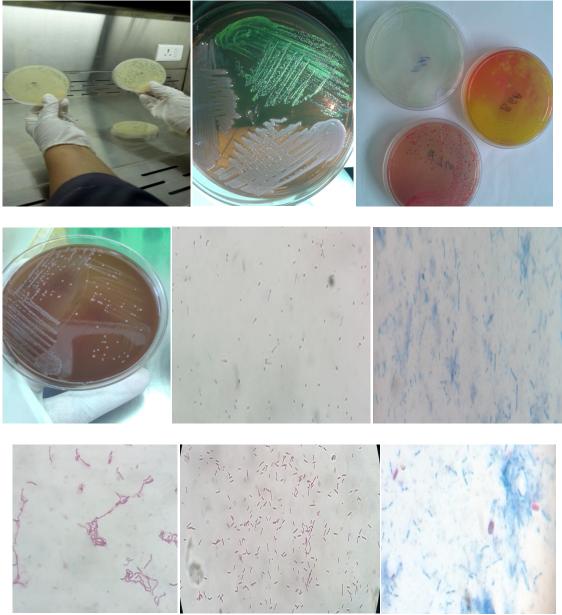


Figure 2: Isolated Bacterial Colonies and their Microscopic view with Gram Staining

- 3.2.2 Biochemical Analysis of Isolates: Nine bacterial species have been isolated from the cattle dung, characterised and identified by standard bacteriological identification procedure. Various kind of biochemical tests have been conducted by inoculating broth culture of the isolates into the variety of media and chemicals. These tests include variety of biochemical, enzymatic and Sugar Fermentation Test (Glucose, sucrose, maltose lactose etc.). Table 3 presents the biochemical and enzymatic features of bacterial isolates which includes biochemical tests, O-F test, methyl red test and citrate utilization test. Table 4 presents the sugar fermentation results of bacterial isolates which includes identifying the presence of mainly glucose, fructose, sucrose and lactose. Biochemical characteristics of isolated bacteria revealed extreme variations in their metabolic capacities. These unique features have been used to analyse and identify the isolates. The isolates are B1 (Bacillus subtilis), B2 (Escherichia coli), B3 (Streptococcus spp), B4 (Pseudomonas aeruginosa), B5 (Clostridium Spp), B6 (Peptostreptococcus Species), B7 (Bacillus Cereus), B8 (Klebsiella Spp), B9 (Bacteroides Species). On the anodic chamber biofilm contains almost anaerobic bacteria which could be the helpful in the generating electricity in the MFC. The major anaerobic bacteria B1 (Bacillus subtilis), B5 (Clostridium Spp), B6 (Peptostreptococcus Species), B7 (Bacillus Cereus), B9 (Bacteroides Species). The isolate findings of this study are comparable to Adegunloye: 2007, Gopinath: 2014, Shivkumar: 2012, Nene:1999, Sawant: 2007, and Kartikey: 2016.
- **4. Conclusion:** The study suggests cattle dung is house of numerous varieties of microbes with unique and variable properties. Exploitation of cattle dung bacterial community may contribute substantially in sustainability of energy generation with agriculture waste. Undoubtedly, more detailed studies of micro flora of dung as a substrate and elaborated research on bio film components in MFC will prove to be a milestone. These will be positive steps to combat energy crisis and depleted fossil energy resources along with environmental problems that eventually controls the global warming. The optimum use of cattle dung micro flora with considerable potential can result in the eco-friendly and better sustainability. To identify electrifying anaerobic bacteria and thoroughly understanding the bio mechanism with

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the help of effective utilization of recent advances in engineering and biotechnology is the need of an hour. There is tremendous scope for research and development in dung based MFC to take into the industrial field along with commercialization. On positive node it can precisely say that dung based biomass may be presumed to be an easily available bioresource material that carries a huge potential to meet sustainable energy development in the near future.

**Table 2: Distinct Morphological Features of Bacterial Isolates** 

Colony		Isolated bacterial colony								
Characteristics	B1	B2	В3	B4	В5	В6	В7	В8	В9	
Colony shape	Rough/ Wrinkled	Round	Round	Round	Round	Round	Round	Round	Round	
Elevation / Colour	Flat	Flat	Raised	Flat	Flat	Flat	Convex	flat	convex	
Optical	Opaque	Opaque	Translucent	Opaque	Dusky	Translucent	Cloudy	Opaque	opaque	
property /	Small	Large	Large	Large	Large	Large	Small	Small	Small	
Size of colony										
Margin	Entire	Entire	Curled	Scattered edges	Undulate	Entire	Entire	Entire	Entire	
Appearance	Dull & Mucoid	smooth	Mucoid	Shiny	Dull	Shiny	Shiny	mucoid	Moist	

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Colony	brownish	Greyish	White	Diffuse	White	Blackish	Greenish	Off	Light Grey
colour	Whitsh	White		greenish		Yellow	White	White	
Gram staining	Positive	Negative	Positive	Negative	Positive	Positive	Positive	Negative	Negative
Reaction									
Bacterial	Bacilli	Bacilli	Cocci in	Bacilli	Bacilli	Cocci in	Bacillus	Bacilli	Short Rods
Morphology		Rods	chain	Medium	Large Rod	Chain	Rods		
				Rod					
Spore staining	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Negative	Negative
Motility	Motile	Motile	Non Motile	Motile	Motile	Non Motile	Motile	Non	Non
								Motile	Motile

Table 3: Biochemical and Enzymatic Features of Bacterial Isolates

Biochemical	B1	B2	В3	<b>B</b> 4	B5	В6	<b>B</b> 7	B8	В9
Tests									
Catalsae	Positive	Positive	Negative	Positive	Negative	Negative	Positive	Positive	Negative
Oxidase	Variable	Negative	Negative	Positive	Negative	Negative	Negative	Negative	Negative
Coagulase	NA	Negative	Negative	NA	NA	Negative	NA	NA	Negative
production									
O-F Test	NA	F	F	O	F	NA	NA	F	F
Habitat*	A/FAn	FAn	FAn	OAn	An	An	A/FAn	FAn	An
Hemolysis	Positive	Negative	Positive	Positive	Negative	Variable	Positive	Negative	Negative
H <sub>2</sub> S	Negative	Negative	Negative	Negative	Positive	Positive	Negative	Negative	Variable
production									
Urease	Negative	Negative	Negative	Negative	Negative	Negative	NA	Positive	Negative
activity									

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Nitrate	Negative	Positive	NA	Positive	Negative	Positive	Variable	Positive	Negative
Reduction									
Indole	Negative	Positive	NA	Negative	Negative	Negative	Negative	Negative	Variable
Production									
Methy red	Negative	Positive	NA	Negative	Negative	NA	Negative	Negative	Negative
Test									
Voges	Negative	Negative	Negative	Negative	Negative	Positive	Positive	Positive	Negative
Proskaur test									
Citrate	Positive	Negative	NA	Positive	Positive	Negative	Positive	Positive	Negative
<b>Utilization test</b>									

<sup>\*</sup>A/Fan-aerobic/facultative anaerobic An-anaerobic FAn-facultative anaerobic NA-not applicable OAn-obligate anaerobic F-fermentative O-oxidative

**Table 4: Sugar Fermentation Results of Bacterial Isolates** 

Sugar	B1	B2	В3	B4	B5	B6	B7	B8	В9
Fermentation									
Glucose	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Positive
Fructose	Positive	NA	Positive	NA*	Positive	Negative	Positive	NA	Positive
Arabinose	Positive	Positive	Negative	NA	Negative	Negative	Negative	Positive	Negative
Galcatose	Positive	NA	Positive	NA	Negative	Variable	Negative	NA	Positive
Arabitol	Negative	NA	Negative	NA	Negative	Negative	Negative	Positive	NA
Dulcitol	Negative	NA	Negative	NA	Variable	Negative	NA	NA	Negative
Maltose	Positive	NA	Positive	NA	Negative	Negative	Positive	Positive	Positive
Sucrose	Positive	Positive	Positive	Negative	Negative	Negative	Variable	Positive	Positive
Xylose	Positive	NA	Negative	NA	Negative	Negative	Negative	Positive	Negative
Lactose	Negative	Positive	Positive	Negative	Negative	Negative	Positive	Positive	Positive

<sup>\*</sup>NA-not applicable

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