

# Bacteriological Quality of Water Contaminated with Faecal Wastes from Livestock

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**Abstract:** Water is very important to life but when contaminated, it can be deleterious to life. Livestock from various farms produce large amount of faecal wastes. Most cattle farmers move their animals from place to place around Imo State, Nigeria, in search of pasture, these animals defecate indiscriminately. Again, many farmers indulge in indiscriminate disposal of animal wastes. These faecal wastes can be carried by runoff water into surrounding water sources causing them to become contaminated. This study was therefore carried out to determine the prevalence of possible pathogenic bacteria in water contaminated with poultry litters, cow and pig dungs. Faecal samples from poultry, cows and pigs were collected from School of Agriculture and Agricultural Technology (SAAT) farm in the Federal University of Technology Owerri (FUTO), Imo State, Nigeria. The contaminated water samples were examined for the prevalence of bacteria. Isolation and characterization of bacteria were done using standard microbiological protocols. Results revealed the presence of *Escherichia coli*, *Enterobacter aerogenes*, *Serratia marcescens*, *Erwinia amylovora*, *Vibrio cholerae*, *Pseudomonas aeruginosa*, *Citrobacter freundii*, *Proteus vulgaris*, *Staphylococcus aureus*, *Bacillus subtilis*, *Streptococcus faecalis* and *Salmonella typhi*. *Pseudomonas aeruginosa* was the most prevalent isolate followed by *Escherichia coli* and *Vibrio cholerae*. The presence of these organisms of public health importance in water contaminated by livestock faeces calls for continuous surveillance of various water sources prone to faecal contamination. Appropriate treatment and disposal methods of these faecal samples as well as appropriate treatment of water source prone to such contaminants should be adopted so as to avert the possible health hazards resulting from use of water from such contaminated sources.

**Keywords:** Bacteria, Contaminated Water, Cow Dung, Pig Dung, Poultry Litters

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## 1. Introduction

Water covers 71% of the Earth's surface [1]. It is vital for all known forms of life. Only 2.5% of the Earth's water is 'freshwater' and 98.8% of that water is in ice and groundwater. Less than 0.3% of all freshwater is in rivers, lakes, and the atmosphere, and an even smaller amount of the Earth's freshwater (0.003%) is contained within biological bodies and manufactured products [1]. Water is our lifeline that bathes us and feeds us. In ancient cultures, water represented the very essence of life. It provides the Earth with the capacity of supporting life [2].

Water pollution is any contamination of water with either chemicals or other foreign substances that are detrimental to

human, plant or animal health [3]. These pollutants may include agricultural wastes, fertilizers and pesticides from agricultural runoff, sewage and food processing waste, lead mercury and other heavy metals, chemical contamination from hazardous waste sites as well as domestic wastes. Worldwide nearly two billion people drink water from sources that could be harmful to health [3]. Water pollution affects plants and organisms living in the bodies of water; in most cases the effects get to natural biological communities [4]. Consumption of contaminated water has far reaching public health import causing water-borne diseases which include: diarrhoea, typhoid fever, nausea, cholera as well as

viral infections [4, 5].

In Imo State, herdsmen and animal farmers engage in indiscriminate grazing and rearing of their animals. These animals defecate randomly in different places and their faeces are deposited via run off to the nearest water bodies. Furthermore, when some farmers clean their pens and farms, they as well, deposit the wastes most times without proper treatment into water bodies. Some of these unassuming farmers may not understand the effect of such indiscriminate dumping of animal wastes products.

When faeces from animals such as cow, pig and poultry come in contact with a water source, they contaminate such water source with their inherent microorganisms (most of which are pathogenic). Such contaminated water if ingested by an individual, can cause high health risk to such individual. Some of these bacteria for example include some strains of *Escherichia coli* which causes diarrhoea and stomach cramp [6]. *Pseudomonas aeruginosa* [7], *Vibrio cholerae* which causes cholera and *Salmonella typhi* which causes typhoid fever[8]. Animal wastes just like other substances such as sewage and agricultural wastes are usually high in organic matter and nutrient. Hence they could cause increase in the microbial flora of the water bodies, thereby resulting in high heterotrophic bacteria counts [9].

**Aim of study:** This study was carried out in order to identify and characterize possible pathogenic bacteria present in water contaminated with poultry litters, cow and pig dung with a view to creating awareness of the existence of such pathogenic organisms in livestock faeces and their public health importance.

## 2. Methodology

### 2.1. Sample Collection and Preparation

Faecal samples from cow, pig and poultry were collected from SAAT farm, located in Federal University of Technology, Owerri West Local Government Area of Imo State. These were carefully put inside appropriately labeled sterile whirl packs and carried to the laboratory for further use.

Twenty eight and half grams of each sample were put into well labeled sterile containers and Six hundred milliliter (600 ml) of sterile water was poured into the containers habouring the faecal samples, giving a value of 47.5 mg/l.

The labeling of the containers was done according to the respective faecal samples used as follows:

|        |                         |
|--------|-------------------------|
| Label: | Faecal sample           |
| A:     | Cow dung                |
| B:     | Poultry litters         |
| C:     | Pig dung                |
| D:     | Control (sterile water) |

The contaminated water samples were then left for 7days to allow/encourage bacterial growth, before the bacteriological analysis commenced.

### 2.2. Isolation and Characterization of Microorganisms

All glasswares, media and diluents used for culture were sterilized by autoclaving at 121°C for 15 minutes at 15 psi, while wire loops were sterilized by flaming as recommended by [10].

The respective samples (contaminated water) were thoroughly homogenized, and 1ml of each was introduced into a sterile test tube containing 9 ml sterile water using a sterile glass pipette. A ten- fold serial dilution was then carried on each sample

Aliquots (0.1 ml) of three different dilutions ( $10^{-3}$ ,  $10^{-5}$  and  $10^{-6}$ ) of each sample were inoculated on the respective agar plates by the pour plate method as described by [10] these were then incubated at 37°C for 24 hours.

After incubation, colonies observed on different plates, were noted. The morphological characteristics of the isolates were also recorded.

### 2.3. Isolation of Pure Culture

Sub-culturing of observed colonies was done onto sterile nutrient agar plates. The streaking technique was employed to obtain pure cultures prior to characterization/ identification of the isolates.

### 2.4. Identification of Bacterial Isolates

Bacteria isolated from the samples were characterized based on the colonial, morphological, standard microbiological and biochemical reactions of the pure cultures as described by [10].

The following biochemical tests were carried out: motility, indole, methyl red, catalase, voges proskauer, coagulase, oxidase and sugar fermentation tests. These tests were carried out in order to confirm the identities of the bacteria isolated. The results obtained from biochemical tests were cross matched with the Bergey's manual of determinative bacteriology [5].

## 3. Results

Table 1 shows the characteristics of different microbial isolates from the respective faecally contaminated water samples labeled A – C and a control labeled D.

## 4. Discussion and Conclusion

### 4.1. Discussion

Results from this study revealed a total of 19 isolates including both Gram negative and Gram positive bacteria. Characterization of the bacterial isolates revealed twelve species (Table 1). Singleton [11] has reported similar species in water contaminated with livestock waste. The presence of these organisms in water is of public health concern because some of these organisms are considered pathogenic. The public health importance of these organisms are highlighted below.

**Table 1.** Characterization/Identification of bacterial isolates from the three different faecal contaminated water samples and control.

| Sampl<br>les | Isolates | Morphological<br>characteristics on<br>nutrient agar | Gram<br>reaction          | Motility | Indole | Methyl red | Catalase | Glucose | Lactose | Oxidase | Coagulase | Voges<br>Proskauer | Probable<br>organism          |
|--------------|----------|--|---------------------------|----------|--------|------------|----------|---------|---------|---------|-----------|--------------------|-------------------------------|
| A            | A1       | Round, cream colonies                                | -ve rods                  | +        | -      | -          | +        | +       | +       | -       | -         | +                  | <i>Enterobacter aerogenes</i> |
|              | A2       | Clustered creamy colonies                            | +ve cocci in clusters     | -        | -      | +          | +        | +       | +       | -       | +         | -                  | <i>Staphylococcus aureus</i>  |
|              | A3       | Smooth, raised, cream colonies                       | -ve rods                  | +        | +      | +          | +        | +       | +       | -       | -         | -                  | <i>Escherichia coli</i>       |
|              | A4       | Circular, reddish colonies                           | -ve rods                  | +        | -      | -          | +        | +       | -       | -       | -         | +                  | <i>Serratia marcescens</i>    |
|              | A5       | Circular grey colonies                               | -ve rods                  | +        | +      | +          | +        | +       | -       | -       | -         | -                  | <i>Erwinia amylovora</i>      |
|              | A6       | Round creamy colonies                                | -ve rods                  | +        | -      | -          | +        | +       | +       | -       | -         | +                  | <i>Enterobacter aerogenes</i> |
| B            | B1       | Circular, yellowish colonies                         | -ve curved rods           | +        | +      | +          | +        | +       | -       | +       | -         | -                  | <i>Vibrio cholerae</i>        |
|              | B2       | Circular, mucoid smooth, greenish                    | -ve rods                  | +        | -      | +          | +        | -       | -       | +       | -         | -                  | <i>Pseudomonas aeruginosa</i> |
|              | B3       | Circular, mucoid smooth, greenish                    | -ve rods                  | +        | -      | +          | +        | -       | -       | +       | -         | -                  | <i>Pseudomonas aeruginosa</i> |
|              | B4       | Circular, mucoid, smooth, greenish                   | -ve rods                  | +        | -      | +          | +        | -       | -       | +       | -         | -                  | <i>Pseudomonas aeruginosa</i> |
|              | B5       | Creamy, flat colonies                                | +ve rods                  | +        | -      | +          | +        | +       | +       | -       | +         | -                  | <i>Bacillus subtilis</i>      |
|              | B6       | Round creamy colonies                                | -ve rods                  | +        | -      | +          | +        | +       | +       | -       | -         | -                  | <i>Citrobacter freundii</i>   |
| C            | C1       | Circular, mucoid, smooth, greenish                   | -ve rods                  | +        | -      | +          | +        | -       | -       | +       | -         | -                  | <i>Pseudomonas aeruginosa</i> |
|              | C2       | Circular, mucoid smooth, greenish                    | -ve rods                  | +        | -      | +          | +        | -       | -       | +       | -         | -                  | <i>Pseudomonas aeruginosa</i> |
|              | C3       | Mucoid, yellow coloured, small                       | +ve cocci in short chains | -        | -      | -          | -        | +       | +       | -       | +         | -                  | <i>Enterococcus faecalis</i>  |
|              | C4       | Round, swarming, cream, flat                         | -ve rods                  | +        | +      | +          | -        | +       | -       | -       | -         | -                  | <i>Proteus vulgaris</i>       |
|              | C5       | Circular, yellowish                                  | -ve rods                  | +        | +      | +          | +        | +       | -       | +       | -         | -                  | <i>Vibrio cholerae</i>        |
|              | C6       | Grey, raised, smooth                                 | -ve rods                  | +        | +      | +          | +        | +       | +       | -       | -         | -                  | <i>Escherichia coli</i>       |
|              | C7       | Cream, round, raised mucoid colonies in chains       | -ve rods                  | +        | -      | +          | +        | +       | -       | -       | -         | -                  | <i>Salmonella typhi</i>       |
| D            | NG       | No observable growth                                 |                           |          |        |            |          |         |         |         |           |                    | NG                            |

Key: -ve = Negative, +ve = Positive, NG = No observable growth.

*Escherichia coli* are a large group of bacteria that can infect someone via ingestion of contaminated water. Most strains of *E. coli* are harmless, however some strains such as *E. coli* 0157:H7 produce a toxin that can cause diseases like diarrhea (often with blood) and stomach cramps. Serious complication of *E. coli* 0157:H7 infection is hemolytic uremic syndrome (kidney failure) [6, 8].

*Enterobacter aerogenes* are called opportunistic pathogens, they can cause numerous infections to humans such as cerebral abscesses, pneumonia, meningitis and septicaemia. This bacterium can infect someone who drinks water

contaminated by it [8, 12].

*Serratia marcescens* are pathogenic bacteria responsible for a variety of infection which includes bacteraemia and intravenous catheter associated infections (10, 12).

*Erwinia amylovora* are plant pathogenic bacteria that infect plant (such as pear and apple) and cause the disease "fire blight" to them. They can be infected when they come in contact with faecally contaminated water either through groundwater uptake or through runoff water.

*Vibrio cholerae* is a human pathogenic bacterium that causes cholera. Cholera is a major public health problem

confronting developing countries. It is associated with poverty and poor sanitation. This disease is characterized by devastating watery diarrhoea and leads to dehydration and sometimes death [8, 9, 10]. It is associated with epidemics and pandemics [10].

*Pseudomonas aeruginosa* is an opportunistic pathogen, it is called opportunistic because it affects immunocompromised hosts (e.g HIV/AIDS, cystic fibrosis and diabetes mellitus). The common site of infection is the lower respiratory tract [10].

*Citrobacter freundii* is an opportunistic nosocomial pathogen that can cause urinary tract infections, blood stream infection and neonatal meningitis that can lead to brain abscesses [11, 13].

*Proteus vulgaris* has been associated with urinary tract infection such as urolithiasis (stone formation in the kidney or bladder) [12].

*Salmonella typhi* is a bacterium that causes typhoid fever. This disease is characterized by the sudden onset of a sustained and systematic fever, severe headache, nausea, loss of appetite, diarrhoea and possible development of meningitis. Untreated typhoid fever can result to death [14].

*Staphylococcus aureus* is a common member of the human micro flora, it can however, produce diseases of adverse health effect [5, 10, 15] such as skin sepsis, post operative wound infections, enteric infections and many more [5, 10]. It is relatively spread in the environment, but found mainly on the skin and mucus membranes of animals. It has also been detected in sewage and in drinking water supplies [15].

*Enterococcus faecalis* formerly known as *Streptococcus faecalis* are facultatively anaerobic, catalase-negative, Gram positive cocci. They are normal inhabitants of the intestinal tract of animals and their oral cavity [5, 10]. They are consistently the second or third most common agent in urinary tract infections, wound infections, and bacterial infections [10]. The hosts for this bacterium include human, pets and livestock [16].

*Bacillus subtilis* is an endospore forming bacterium. The endospore allows it to withstand extreme temperatures. Therefore water contaminated by livestock waste will require a proper treatment to make the water fit for drinking and other purposes. However, this bacterium is not considered strictly pathogenic [17].

*Pseudomonas aeruginosa* was the most prevalent organism followed by *Escherichia coli* and *Vibrio cholerae* each being found in at least two of the faecally contaminated sample. The contamination of water by these bacteria is worrisome: The presence of *Escherichia coli* in water not only pose problem of being pathogenic but also indicates the presence of other pathogenic organisms in such water body because it is a known "indicator organism"[18]. *Vibrio cholerae* on its own is a very versatile epidemic causing organism especially among children [8, 9, 10]. *Pseudomonas aeruginosa* is a known recalcitrant organism [8].

#### 4.2. Conclusion

Based on the findings of this study, it is evident that,

contamination of water with livestock waste can lead to introduction of various types of bacteria of public health importance. These can easily be transferred to man and even his livestock via consumption, wadding and bathing activities.

There is therefore need for deliberate policies and laws by the government to restrict illegal dumping of faecal waste from livestock farms into water sources. Public awareness in form of workshops should be organized for livestock farmers and indeed the entire community around such environment to educate them on the importance of water sanitation and good faecal waste disposal methods. Livestock farmers should be assisted via incentives which can help manage livestock waste and this management process should be supervised by waste management experts.

Closed grazing system where the livestock waste products are utilized within the system and does not portend a burden to the environment and water sources around the environment should be adopted by livestock farmers. Furthermore livestock farms should have improved hygiene standard within and around their environment. These will in turn help to alleviate pollution of the environment especially water resources around such areas thereby conserving the water that the populace that uses it for sundry activities.

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