

https://doi.org/10.47430/ujmr.2493.010

Received: 26<sup>th</sup> February 2024

Accepted: 15th May 2024



# Bacteriological Quality of Borehole Water in Gusau Metropolis

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#### Abstract

Microbes are ubiquitous and are known to contaminate materials, including food and water. We examined the bacteriological quality of borehole water in Gusau metropolis to determine its potability. Triplicate samples were collected from fifteen (15) different boreholes and analyzed. The total bacterial load, total coliform, fecal coliform, and Salmonella-Shigella count were determined using the membrane filtration technique. The average counts were as follows: total bacteria ranged from  $6 \times 10^{-1}$  cfu/ml to  $39 \times 10^{-1}$  cfu/ml, total coliform count from  $3 \times 10^{-1}$  cfu/ml to  $65 \times 10^{-1}$  cfu/ml, fecal coliform count from  $1 \times 10^{-0}$  cfu/ml to  $4 \times 10^{-0}$  cfu/ml, and Salmonella-Shigella count from 0 cfu/ml to  $4 \times 10^{-0}$  cfu/ml. The isolated organisms were identified as Escherichia coli, Salmonella typhi, and Shigella spp. The predominant bacterial isolate was Escherichia coli. Our study indicated that the bacteriological quality exceeded the World Health Organization (WHO) allowable limits of 0 cfu/100ml for total bacterial load, total coliform count, fecal coliform count, and Salmonella-Shigella count due to the underground aquifers. Therefore, we recommend that borehole water sources be adequately treated before consumption to reduce the risk of waterborne diseases.

Keywords: Bacteriological, Quality, Borehole, Water and Coliform.

# **INTRODUCTION**

Water is a ubiguitous substance essential for life on Earth, composed of the chemical elements hydrogen and oxygen. Its molecular formula is  $H_2O$ , indicating that each water molecule consists of two hydrogen atoms bonded to one oxygen atom (Theodore *et al.*, 2017). It is one of the most essential compounds that is colorless, tasteless, and odorless liquid. Under room temperature, it possesses the ability to dissolve many other substances (Zumdahl, 2023). Water from both surface and groundwater sources plays vital roles in meeting diverse human needs, including domestic, industrial, agricultural, and environmental purposes (United Nations, 2019). Surface water includes water that flows across the land in the form of streamlets, springs, streams, and rivers or it collects to form ponds, lakes, and seas (Niyogi, 2011). In contrast, groundwater is located in aguifers underground and links with surface water through penetration and springs (Gleeson et al., 2012). The underground water supplies are usually considered safe provided they are

located, constructed, and operated according to the guidelines for drinking water (WHO 2018).

Owing to the government's failure to meet the ever-increasing water demand, many individuals turn to groundwater sources like boreholes as an alternative water source (Harvey, 2014). Therefore, people can obtain groundwater through boreholes drilled into aquifers for industrial, agricultural, and domestic purposes. Despite access to water, it does not guarantee access to safe water; as per the United Nations International Children Emergency Fund (UNICEF) and World Health Organization (WHO), Nigeria is classified among countries lacking access to improved water sources (UNICEF and WHO, 2019).

Lack of basic sanitation and poor hygienic practices in the supply of water for drinking and other domestic uses have been associated with a high morbidity and mortality rate globally, with diarrhea and cholera being reported as killer diseases (WHO 2018). These issues have been persistent in Gusau metropolis (NCDC, 2021). Therefore, this study aimed to assess the bacteriological quality of borehole water in Gusau metropolis.

# MATERIALS AND METHODS

#### Study Area

This study was conducted in Gusau local government area, Zamfara state. The area is situated between latitude  $12.09^{\circ}$ N- $12.5^{\circ}$ N and longitude  $6.67^{\circ}$ E, covering an area of 3,364 km<sup>2</sup> with an estimated population of 582,100 as of March 21, 2019 (NPCN, 2019). The samples were gathered from 15 distinct boreholes in Gusau metropolis and conveyed to the microbiology laboratory at Federal University Gusau for bacteriological analysis.

#### **Boreholes location**

Fifteen boreholes situated in the following locations within the study settings were sampled in triplicate:

Danbedi, Maryam hall, Ugwuan Dallatu, Gidan Musa, Gidan Mohammed, Gidan Kabiru, Bayan NTA, Tankin Ruwa, Bayan NEPA, Makaranta, Dan Buba, Living Faith, Gidan Duchi, Gidan Ahmed, Gidan Kema

# Sample collection

Samples for bacteriological analysis were aseptically collected in sterilized 250ml screwcapped plastic containers. All plastic containers used for sample collection were sterilized, rinsed with 70% ethanol, distilled water, and the water samples. Borehole water samples were collected into sterile plastic containers by allowing the water to flow for 5 minutes, then the containers were uncapped, filled with the water samples, and recapped. Care was taken to prevent air bubbles from entering the container. The samples were transported to the laboratory in an ice box and analyzed within 24 hours of collection following the method outlined by Onuorah *et al.*, 2022.

#### Determination Of Bacteriological Quality

The total bacterial, total coliform, fecal coliform, salmonella, and shigella counts were determined as described by APHA, (2017).

# **Total Bacterial Count**

Nutrient agar was weighed and prepared according to the manufacturer's instructions. It was sterilized in an autoclave at 121°C for 15 minutes, allowed to cool, and then aseptically dispensed into Petri dishes. The membrane filter paper containing the bacteria was carefully placed with the grid side facing up on the nutrient agar. Duplicate plates were prepared and labeled for the fifteen water samples. Incubation was carried out in an inverted position at 28°C for 24 hours, after which the bacterial colonies that developed were counted, and the results were recorded. The colonies were subcultured and stored on a sterile nutrient agar slant for characterization and identification (APHA, 2017).

#### **Total Coliform Count**

MacConkey agar was weighed and prepared according to the manufacturer's instructions. It was sterilized in an autoclave at 121°C for 15 minutes, allowed to cool, and then aseptically dispensed into Petri dishes. The membrane filter paper containing the bacteria was carefully placed with the grid side facing up on the MacConkey agar. Triplicate plates were prepared and labeled for the fifteen water samples. Incubation was carried out in an inverted position at 28°C for 48 hours, after which the bacterial colonies that developed were counted, and the results were recorded. The colonies were subcultured and stored on a sterile nutrient agar slant for characterization and identification (APHA, 2017).

#### Faecal Coliform Count

Endo agar was prepared according to the manufacturer's instructions, sterilized in an autoclave at 121°C for 15 minutes, cooled, and then dispensed aseptically into Petri dishes. The membrane filter paper with bacteria was placed on the endo agar with the grid side up. Duplicate plates were prepared and then placed in an incubator at 28°C for 48 hours. The developed faecal coliform bacteria were counted, and a few colonies were subcultured and stored on a sterile nutrient agar slant for additional studies (APHA, 2017).

#### Salmonella-Shigella count

The Salmonella-Shigella agar (SSA) was prepared following the manufacturer's instructions, and a 1ml aliguot of each water sample was transferred onto the dried and sterilized SSA plates. The plates were inoculated, evenly spread, and then incubated at 37°C for 48 hours. Subsequently, pure cultures were obtained by sub-culturing onto freshly prepared SSA plates, and pure colonies were identified based on their biochemical and morphological characteristics (Onuorah et al., 2022).

# Sample Preparation

The membrane filtration apparatus was utilized to filter the water samples. The samples were adequately mixed by inverting the container multiple times. A sterile forceps was used to place the filter paper in the apparatus. Subsequently, 100ml of borehole water sample was slowly filtered through the membrane filter in the funnel. The membrane filter paper was then delicately removed with a sterile smoothtipped forceps and placed grid side up on the culture medium in the Petri dish, ensuring no air bubbles were trapped underneath the membrane (APHA, 2017).

# Characterization and Identification of the Isolates

The bacterial isolates were characterized based on their morphological and biochemical characteristics. Gram staining, catalase, **urease**, indole, coagulase, citrate utilization, Carbohydrate utilization test, motility test, Methyl Red, Voges-Proskauer **test**, and Hydrogen Sulphide production tests were all conducted following the procedures outlined by Willey *et al*. (2020). **The** isolates were identified according to the scheme **proposed by** Patel (2020).

### RESULTS

The total bacteria, total coliforms, faecal coliforms and Salmonella Shigella count per 100ml of the borehole water in Gusau metropolis are shown in Table 1. The Total bacterial count ranges from 6 to 39 cfu/100ml,Total coliform count 3 to 65 cfu/100ml while the faecal coliform count range is 1 to 4 cfu/100ml, also the total *Salmonella Shigella* count range is from 0 to 4 cfu/100ml, (Table 1).

The morphological and biochemical characteristics of the bacterial isolates from borehole water samples are depicted in Table 2. The isolated bacteria include *Salmonella* typhi, *Shigella* spp, and *Escherichia* coli.

The frequency of occurrence of the bacterial isolates in the borehole water in Gusau metropolis is depicted in Table 3. *Escherichia coli* exhibited the highest frequency of occurrence, with 15 instances (65.21%), while *Salmonella typhi* and *Shigella* spp had the lowest frequency of occurrence, with 4 instances each (17.39%), in the borehole water samples analyzed.

Table 1: Total bacterial	count, T	Total coliform	count, feacal	coliform	count,	and Salmonella-
Shigella count.						

Sample	Borehole Location	Total Bacterial Count (CFU/ml)	Salmonella- Shigella Count (CFU/ml)	Total Coliform Count (CFU/ml)	Feacal Coliform Count (CFU/ml)
1	Danbedi	33×10 <sup>1</sup>	0×10 <sup>0</sup>	10×10 <sup>1</sup>	1×10 <sup>0</sup>
2	Maryam Hall	26×10 <sup>1</sup>	0×10 <sup>0</sup>	20×10 <sup>1</sup>	1×10 <sup>0</sup>
3	UgwanDallatu	29×10 <sup>1</sup>	0×10 <sup>0</sup>	31×10 <sup>1</sup>	1×10 <sup>0</sup>
4	Gidan Musa	39×10 <sup>1</sup>	0×10 <sup>0</sup>	13×10 <sup>1</sup>	2×10 <sup>0</sup>
5	GidanMuhammadu	35×10 <sup>1</sup>	0×10 <sup>0</sup>	19×10 <sup>1</sup>	1×10 <sup>0</sup>
6	GidanKabiru	13×10 <sup>1</sup>	3×10 <sup>0</sup>	6×10 <sup>1</sup>	1×10 <sup>0</sup>
7	Bayan NTA	29×10 <sup>1</sup>	2×10 <sup>0</sup>	8×10 <sup>1</sup>	2×10 <sup>0</sup>
8	TankinRuwa	15×10 <sup>1</sup>	0×10 <sup>0</sup>	5×10 <sup>1</sup>	3×10 <sup>0</sup>
9	Bayan NEPA	22×10 <sup>1</sup>	0×10 <sup>0</sup>	4×10 <sup>1</sup>	2×10 <sup>0</sup>
10	Makaranta	33×10 <sup>1</sup>	1×10 <sup>0</sup>	3×10 <sup>1</sup>	1×10 <sup>0</sup>
11	Dan Buba	9×101	0×10 <sup>0</sup>	65×10 <sup>1</sup>	4×10 <sup>0</sup>
12	Living Faith	15×10 <sup>1</sup>	4×10 <sup>0</sup>	17×10 <sup>1</sup>	1×10 <sup>0</sup>
13	GidanDuchi	6×10 <sup>1</sup>	0×10 <sup>0</sup>	22×10 <sup>1</sup>	2×10 <sup>0</sup>
14	Gidan Ahmed	12×10 <sup>1</sup>	3×10 <sup>0</sup>	30×10 <sup>1</sup>	1×10 <sup>0</sup>
15	GidanKema	7×10 <sup>1</sup>	1×10 <sup>0</sup>	22×10 <sup>1</sup>	3×10 <sup>0</sup>
	WHO Standard	0	0	0	0

E-ISSN: 2814 – 1822; P-ISSN: 2616 – 0668

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Isolates	1	2	3
Gram Staining	-	-	-
Form	Rod	Rod	Rod
Indole	+	-	-
Methyl Red	+	+	+
Voges Proskaer	-	-	-
Citrate	-	-	-
Urease	+	-	-
Glucose	+	+	+
Lactose	+	-	-
Sucrose	-	-	-
H <sub>2</sub> S	+	-	-
Catalase	+	+	+
Motility	+	+	-
Suspected Organisms	Escherichia coli	Salmonella typhi	Shigella spp

Table 2: Morphological and Biochemical characteristics of the Bacteria Isolates fromBoreholewater samples in Gusau Metropolis.

**Key;** + = Positive - = Negative

Table 3: Occurrence of the Bacterial Isolates in the Borehole water investigated in Gusau Metropolis

Sample	Borehole Location	Salmonella typhil	Escherichia coli	Shigella spp
1	Dambedi, Hayan Buba	-	+	-
2	Maryam Hall, Hayan Buba	-	+	-
3	Ugwan Dallatu Hayan Buba	-	+	-
4	Gidan Musa Low-cost	-	+	-
5	Gidan Muhammadu Low- cost	-	+	-
6	Gidan Kabiru Low-Cost	-	+	-
7	Bayan NTA Samaru	-	+	+
8	Takin Ruwa Samaru	+	+	-
9	Bayan NEPA Samaru	+	+	-
10	Makaranta Premier	-	+	+
11	Dan Buba Premier	-	+	-
12	Living Faith Premier	+	+	-
13	Gidan Duchi Gadabiyu	-	+	+
14	Gidan Ahmed Gadabiyu	-	+	+
15	Gidan Kema Gadabyu	+	+	-

Key; + = Present

– = Absent

# DISCUSSION

The quality of the borehole water from the various sample points varies as follows: The total bacterial count ranges from  $6 \times 10^1$  to  $39 \times 10^1$ cfu/ml (Table 1). The sample from Gidan Musa, Low cost had the highest count of  $39 \times 10^{1}$  cfu/ml. while the sample from Gidan Duchi Gadabiyu had the least count of  $6 \times 10^1$  cfu/ml, indicating a high level of pollution of borehole water due to human and animal activities. The values obtained from all the water samples were higher than the WHO standard (Table 1). This is in agreement with the previous work by Fardami et al. (2020), who had earlier reported high microbial counts in drinking water in Zamfara North Senatorial District, Nigeria. The result of the total coliform counts ranges from 3×10<sup>1</sup> to  $65 \times 10^{1}$  cfu/ml (Table 1). The results showed the total coliform counts for all the various water sources, where the least coliform counts were 3×10<sup>1</sup> cfu/ml at Gidan Kabiru Low-cost, and the highest total coliform counts were 65×10<sup>1</sup> cfu/ml at Dan Buba, Premier borehole water. This corresponds with the work of Onwughara et al., (2013), who recorded a high coliform count above the WHO standard in Umuahia North Local Government Area, Abia State, Nigeria. The fecal coliform counts ranged from 1×10° cfu/ml to  $4 \times 10^{0}$  cfu/ml (Table 1) and this exceeded the WHO standard for coliform bacteria in water, which is zero total coliform per 100 ml of water (Table 1). Eight locations had  $1 \times 10^{\circ}$  cfu/ml, while Dan Buba had the highest fecal coliform count of  $4 \times 10^{\circ}$  cfu/ml. The presence of coliforms in the samples is an indication of fecal contamination. This is in agreement with the findings of Kumarasamy et al. (2009), who reported fecal coliform counts ranging from 1 to 6 CFU/100ml. The Salmonella-Shigella count ranges from 0 to  $5 \times 10^{\circ}$  CFU/ml, with all the samples analyzed within the WHO standard except for Gidan Kabiru, Bayan Nepa, Makaranta, Living Faith, Gidan Ahmed, and Gidan kema, which exceeded the WHO standard of 0 CFU/ml (Table 1). This is in correspondence with the findings of Fardami et al. (2019), who reported Salmonella-Shigella counts ranging from 0 to 4 CFU/ml in borehole water samples analyzed in Zamfara, North Senatorial District, Nigeria. Water samples from all the analyzed sample locations are only fit for domestic purposes due to the fact that the water samples total coliform, analyzed contained fecal coliform, total bacterial count, and some Salmonella-Shigella count above the WHO standard limits of 0 CFU/ml, making them unfit for human consumption. Further analysis of the bacteriological quality of water samples collected in this study revealed the presence of three dominant bacterial genera, namely; *Escherichia coli, Salmonella* typhi, and *Shigella* spp (Table 2).

The most frequently isolated bacterium in this study was Escherichia coli (Table 3), commonly found in the lower intestine of warm-blooded organisms. lts presence indicates fecal contamination of the water due to human and animal waste, which can lead to gastrointestinal disorders (Kuta, 2008; Onwughara et al., 2013). Escherichia coli is known to cause various illnesses, including watery and bloody diarrhea, dysentery, urinary tract infections (WHO, 2018), and bacteremia if it enters the bloodstream. Moreover, certain strains of E. coli have the ability to produce enterotoxins in the human small intestine, leading to diarrhea (Obioma et al., 2017).

Salmonella typhi, which was also predominant in the samples analyzed, can be acquired by ingestion of food and water contaminated by feces of infected humans or through person-toperson contact. The symptoms include fever, headache, abdominal pains, anorexia, and malaise. The bacteria can reinfect the gastrointestinal tract, producing abdominal pain and diarrhea (Willey *et al.*, 2008).

Shigella, which was also predominant in the shigellosis samples, causes or bacillary dysentery, a diarrheal illness resulting from an acute inflammatory reaction of the intestinal tract caused by the four species of the genus Shigella. The organism is transmitted by the fecal-oral route, primarily through food, feces, fingers, and flies (the four fs) (Willey et al., 2008). This is in agreement with the findings of Fardami et al., (2019), who isolated similar microorganisms in their analysis of drinking water.

#### CONCLUSION

The bacteriological parameters affecting groundwater originate from various sources, including human waste, agricultural waste, and industrial waste. The borehole water in Gusau metropolis that was studied exhibited poor quality concerning the microbiological parameters assessed. Most of the microbiological parameters exceeded WHO standards, except for some of the Salmonella-Shigella counts that fell within the WHO specified limit. The presence of indicator bacteria such as Escherichia coli, Salmonella typhi, and Shigella spp (Table 2) in high

concentrations indicates the potential presence of pathogenic bacteria in the borehole water analyzed. Therefore, it is imperative to treat borehole water before human consumption.

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