

Research Article

Escherichia coli as a Tool for Disease Risk Assessment of Drinking Water Sources

Stephen T. Odonkor ¹ and Tahiru Mahami²

¹School of Public Service and Governance, Ghana Institute of Management and Public Administration, Accra, Ghana

²Ghana Atomic Energy Commission, Kwabenya, Accra, Ghana

Correspondence should be addressed to Stephen T. Odonkor; stodonkor@gimpa.edu.gh

Received 19 March 2020; Revised 25 May 2020; Accepted 28 May 2020; Published 15 June 2020

Academic Editor: Joseph Falkinham

Copyright © 2020 Stephen T. Odonkor and Tahiru Mahami. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Many diseases have been associated with poor drinking water quality including diseases caused by diarrheagenic pathogens, especially in developing countries where access to a consistent water supply is a problem. The objective of the study was to evaluate the health risks associated with the sources of drinking water in the Dangme West District of Ghana using *E. coli* as a measurement tool, aiming at ascertaining the paths leading to contamination of the water sources. A total of 464 water samples were obtained for analysis. Sampling covered a year across the dry and wet seasons in Ghana. Water samples were obtained from groundwater and surface water sources. *E. coli* counts were determined using the most probable number method (MPN). Disease risk assessment was carried out using the WHO drinking water risk assessment guidelines. Generally, the study revealed significantly higher *E. coli* counts in the wet season than in the dry season. Among the water samples analyzed, surface water, especially from the dams, was found to pose the highest disease risk as compared to the other water sources. Samples from groundwater sources, especially boreholes, posed the lowest disease risk. In conclusion, observations from the study implied that most water sources in the study district are highly polluted with bacteria pathogens beyond recommended safety guidelines. The main causes of faecal contamination in these water sources were purported to be anthropogenic. Therefore, there is a need to formulate a policy aimed at managing and improving rural water sources.

1. Introduction

Water is today a scarce resource, particularly in the developing world. Globally, 663 million people do not have access to safe water [1]. Rural communities in sub-Saharan Africa account for more than 50% of those lacking potable water [2, 3]. Most of these communities, therefore, rely on untreated sources such as streams, dams, boreholes, wells, and rivers to meet fundamental needs such as drinking, sanitation, and cooking and for their sustainable development [4]. However, these untreated sources are contributing significantly to the global burden of disease, as a result of water-related infectious diseases such as cholera, dysentery, and typhoid [5]. Every year, millions of lives are lost worldwide with a chunk in developing countries as a result of waterborne diseases. Diarrhoea is a major cause of the

death of more than 2 million people per year worldwide, mostly children under the age of five. In fact, each day, nearly 1,000 children die due to preventable water and sanitation-related diarrhoeal diseases [6, 7].

Good quality water is a vital deciding factor for human health, which directly relates to the socioeconomic progress of a country [8–10]. Thus, the importance of water cannot be overemphasized. In fact, sustainable socioeconomic development and productive livelihoods can be built on access to water. Productive uses of water are employed at the household levels, including a range of small-scale domestic activities, in agriculture and industrial purposes. Although the quality requirement for water varies from one sector to the other and from one form of usage to the other, many of these activities have implications for public health, because it can result in the transmission of waterborne diseases [11–13].

These water-related infections are a result of the pollution of the water sources, due to anthropogenic influences, population growth, and urbanization [14]. The pollution thus leads to the development and propagation of pathogenic waterborne microorganisms. *Escherichia coli* is one of the major pathogens associated with waterborne diseases. Naturally, *Escherichia coli* (*E. coli*) is a facultative anaerobic bacterium that inhabits the large gastrointestinal tracts of warm-blooded animals and is a major normal flora associated with the human colon [15–17].

Therefore, the existence of *E. coli* in food or water normally signals recent faecal contamination or poor hygienic conditions in food processing facilities [18]. Thus, faecal contamination, poor sanitation measures, and poor storage conditions have a major influence on *E. coli* population [19, 20]. The mere existence of *E. coli* in water does not necessarily imply the presence of disease-causing microbes. However, it gives an indication of the possible existence of faecal-borne microorganisms such as Salmonella and hepatitis A [21, 22]. This accounts for the usage of *E. coli* as an indicator microbe to examine food and water samples aimed at detecting the levels of faecal contaminations [21].

Water safety management depends primarily on hazard identification through drinking water surveillance, which is “the continuous and vigilant public health assessment and review of the safety and acceptability of drinking water supplies” [23, 24]. This surveillance is critical to public health and safety, as it contributes and supports improvements in water quality and accessibility. The present study evaluated the health risks associated with sources of drinking water in the Dangme district in Ghana using *E. coli* as a bacteria source tracking tool measurement tool, aimed at ascertaining the paths leading to contamination of the water sources.

2. Materials and Methods

2.1. Study Area. The Dangme West District (Figure 1) is situated in the southeastern part of Ghana, lying between longitude 5° 45' south and 6° 05' north and longitude 0° 05' east and 0° 20' west. The district has a total land area of 1,442 square kilometers, making it the largest in the Greater Accra Region. The land size represents 41.5% of the regional land area. The district forms part of sixteen (16) metropolitan, municipal, and districts in the Greater Accra Region of Ghana. The district has a 37 km Coastline and a 17 km stretch of the Volta River [25].

2.2. Topography and Drainage. The district forms the central portions of the Accra plains. The relief is generally gentle and undulating, a low plain with heights not exceeding 70 metres. The plains are punctuated in isolated areas by a few prominent inselbergs, isolated hills, outliers, and knolls scattered erratically over the area. Prominent relief features include the Yongua inselberg (427 metres) which appears conical in the air with a number of outliers close to the north of the district around Asutsuare and Osuwem areas, the Krabote inselberg also to the north and the Shai hills (289 metres) found towards the western portions of the district [26]. The mean annual

rainfall increases from 762.5 milliliters on the coast to 1220 milliliters to the North and Northeast close to the foothills of Akwapim range and on the summit [26].

2.3. Climate and Vegetation. The southeastern coastal plain of Ghana, which encompasses the Dangme West District, is one of the hottest and driest parts of the country. Temperatures are, however, subjected to occasional and minimal moderating influences along the coast and altitudinal influences affected by the Akwapim range in the northwest. Temperatures are appreciably high for most parts of the year, with the highest during the main dry season (November–March) and the lowest during the short dry season (July–August). They average a few degrees lower on the coast and close to the Akwapim range than they do over most of the plains. The absolute maximum temperature is 40°C [26]. Along some stream courses, however, higher vegetation type ranging from thickets to light forest is common. Some light forest with tall trees is also found along the foothills of the Akwapim Ridge, especially around Dodowa, Ayikuma, and Agomeda areas. There is a Forest, Game, and Wildlife Reserve around the Shai hills [26].

2.4. Water Sources and Sanitation. The supply of potable water in the district is woefully inadequate, and only a few sections of the district have a regular supply of pipe-borne water. Analysis of the current water and sanitation in the district shows that more effort is needed to meet the 85% water and sanitation coverage. On the basis of the National Community Water and Sanitation Standards of 600 people per standpipe, 350 persons per borehole, and 150 persons per hand-dug well, the district has achieved about 66% coverage with 34% of the population lacking access to a potable water supply. As it stands now, below 37% of the district population has access to pipe-borne water. The remaining population in the district depends on untreated water sources. Visually, the water from the streams and dams are light brownish yellow caused by mostly decayed dead leaves. If turbulently disturbed, it turns to deep brownish yellow and some suspended soils can be seen [26].

2.5. Sample Collection and Analysis. Four hundred sixty-four (464) water samples were obtained from the drinking sources at the study location for analysis. Sampling duration was a year, and sampling was carried out throughout the wet and dry seasons in Ghana. The water samples were obtained from the surface (dams, river, canals, and streams) and underground (hand-dug wells and boreholes) water sources.

Sampling and care of samples were done following guidelines of the WHO [27] while analyses of samples were carried out as described by APHA [28, 29].

Sterile bottles were used to collect the water samples from each of the sites. Samples were taken to the laboratory for analyses. The most probable number method (MPN) illustrated by Prescott et al. [30] was used to determine *E. coli* counts. A total of 15 tubes comprising 5 tubes for each of three dilution factors (0.1 ml, 1 ml, and 10 ml) were used for

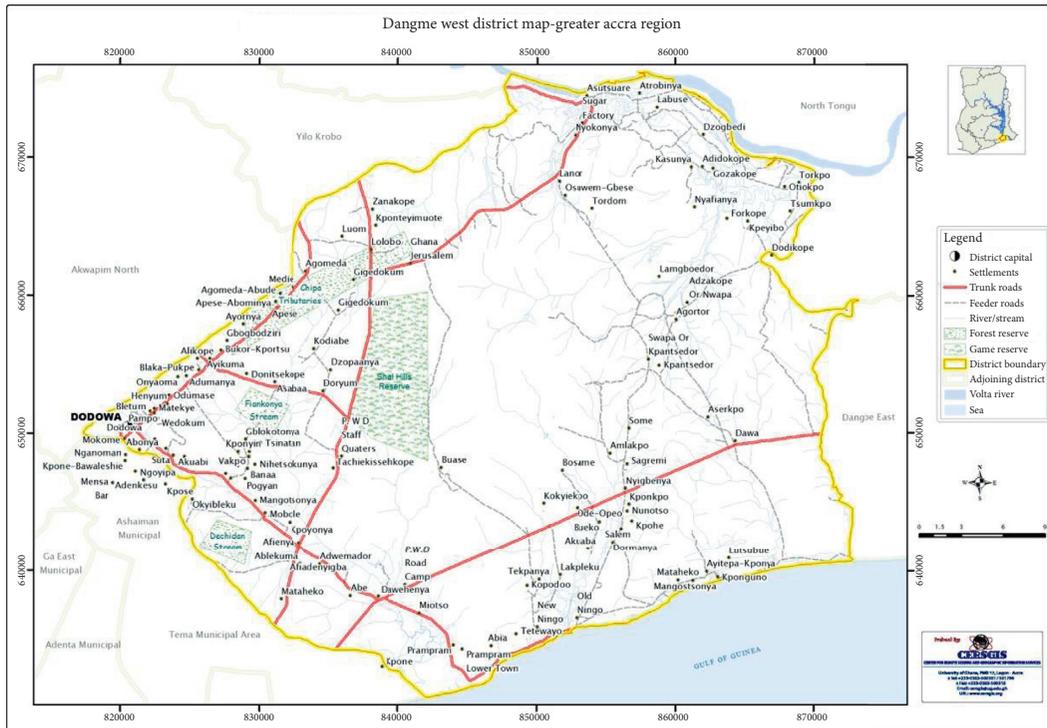


FIGURE 1: A map of the Dangme district of Ghana.

inoculation in the study. As a result, three sets, each containing five tubes, were used in the presumptive test. In the first set, each of the five tubes holding 10 ml of MacConkey Broth with double strength was inoculated with 10 ml of the water sample. The tubes in the second set, which contain 10 ml of single strength MAC broth, were inoculated with 1 ml of subwater samples and the remaining five tubes containing 10 ml of single strength MAC broth were inoculated with 0.1 ml of the sample. After that, the tubes were incubated at 37°C for 24 hours for total coliforms and 44°C for faecal coliforms. The tubes were then examined visually for turbidity. Each tube contains an inverted Durham tube (this is a very small test-tube), which is examined for gas production (bubble) and the media is examined for acid production, i.e, change in colour from pink to yellow.

Confirmation of *E. coli* was carried out by aseptically plating from tubes with acid and gas formation onto L-EMB agar in Petri dishes to obtain pure isolates. The plates were incubated at 35°C with a standard error of 0.5°C for 18–24 hours. The incubated plates were examined for centrally dark and flat colonies having or not having a metallic sheen. These morphological features are associated with *E. coli*. A maximum of 5 colonies from each L-EMB plate was inoculated onto slants of PCA and incubated at 35°C for about 24 hours to prepare stork culture. API20E was used to identify a 24 hr culture of each purified isolate following the manufacturer’s instructions.

3. Results

Table 1 shows the range of seasonal *E. coli* counts (MPN/100 ml) in water. *E. coli* populations in samples sourced from

TABLE 1: The range of seasonal *E. coli* counts (MPN/100 ml) in water.

Water sources	Seasonal counts (MPN/100 ml)	
	Dry season	Wet season
Borehole	0–0.2 × 10 ¹	0–0.2 × 10 ¹
Canal	2.3 × 10 ¹ –1.1 × 10 ²	3.1 × 10 ¹ –6.3 × 10 ¹
Dam	0.7 × 10 ¹ –1.1 × 10 ²	1.2 × 10 ¹ –7.9 × 10 ¹
Hand-dug well	0–0.2 × 10 ¹	0–0.4 × 10 ¹
River	0–0.1 × 10 ¹	0.2 × 10 ¹ –0.7 × 10 ¹
Streams	0–3.3 × 10 ¹	0–3.3 × 10 ¹

hand-dug wells, boreholes, and stream all had a zero-minimum count across both seasons. There was a low count of *E. coli* in water samples obtained from river sources in the wet season. The range of *E. coli* counts (1.2 × 10¹–7.9 × 10¹ MPN/100 ml) was observed in samples collected from dams throughout the wet season.

A summary of comparative analysis between *E. coli* counts measured in MPN/100 ml in samples from the water sources is presented in Table 2. The highest counts for *E. coli* (59.67 ± 45.08) were identified in water samples collected from the canal sources in the wet season. On the other hand, the lowest counts (0–0.1 × 10¹ MNP/100 ml) were identified in the dry season from samples sourced from the rivers.

Analysis of mean variations and confidence intervals for *E. coli* proportions throughout the wet and dry seasons is shown in Tables 3 and 4, respectively. There was a significant difference between mean proportions of *E. coli* from boreholes and canal water sources in the wet season. Mean differences of counts among boreholes, river, and canals in the wet season were also significant.

TABLE 2: Comparison of *E. coli* counts (MPN/100 ml) in the various water sources (SD = standard deviation, df = degree of freedom, Min = minimum, and Max = maximum).

Water sources	Dry season				Wet season				P value
	Mean ± SD	Min	Max	df	Mean ± SD	Min	Max	df	
Bore hole	0.63 ± 0.74	0	2	7	1 ± 0.76	0	2	7	0.090
Canal	47.67 ± 16.04	31	63	2	59.67 ± 45.08	23	110	2	0.280
Dam	38.07 ± 30.71	7	110	14	39.87 ± 19.76	12	79	14	0.350
Hand-dug well	0.93 ± 0.80	0	2	14	1.47 ± 1.06	0	4	14	0.001
River	0 ± 0	0	0	2	4.33 ± 2.52	2	7	2	0.050
Stream	15.06 ± 13.59	0	33	16	20.77 ± 11.05	0	31	16	0.020

TABLE 3: Analysis of mean differences and confidence intervals for *E. coli* levels in the dry season (test value = 6; the mean difference is significant at $P \leq 0.05$).

Water sources	T	df	Sis (2-tailed)	Mean difference	95% confidence interval	
					Upper limit	Lower limit
Borehole	2.376	7	0.049	0.625	0.00	1.25
Canal	5.147	2	0.036	47.667	7.82	87.52
Dam	4.800	14	0.001	38.067	21.06	55.08
Hand-dug well	4.525	14	0.001	0.933	0.49	1.38
River	—	2	—	—	—	—
Stream	4.569	16	0.001	15.059	8.07	22.05

TABLE 4: Analysis of mean differences and confidence intervals for *E. coli* levels in the wet season (test value = 6; the mean difference is significant at $P \leq 0.05$).

Water sources	T	df	Sis (2-tailed)	Mean difference	95% confidence interval	
					Upper limit	Lower limit
Bore hole	3.742	7	0.007	1.000	0.37	1.63
Canal	2.292	2	0.149	59.667	-52.32	171.66
Dam	7.814	14	0.001	39.867	28.92	50.81
Hand-dug well	5.358	14	0.001	1.467	0.88	2.05
River	2.982	2	0.096	4.333	-1.92	10.58
Stream	7.745	16	0.001	20.765	15.08	26.45

Findings of the investigation of the extent of disease risk associated with the drinking water sources in both seasons (wet and dry) are shown in Figures 1 and 2, respectively. This analysis was carried using WHO drinking water risk assessment guidelines. According to the World Health Organization, a zero count of *E. coli* per 100 ml of water is considered safe for drinking. A count of 1–10 MPN/100 ml is regarded as low risk; 11–100 MPN/100 ml is medium risk. Finally, an *E. coli* count greater than 100 MPN/100 ml is adjudged high risk.

With reference to Figure 2, it is evident that the dam and canal water sources revealed intermediate disease risk (100%) throughout the wet season. All water samples (100%) sourced from the canals showed an intermediate risk level followed by that of a dam with an 87% intermediate risk level. An intermediate risk level of 59% was observed in water obtained from streams.

Water from rivers had 100% compliance with WHO standards. However, water from canals and dams showed no conformity with the guidelines, thereby implying the substantial health risks they pose to human health. Water

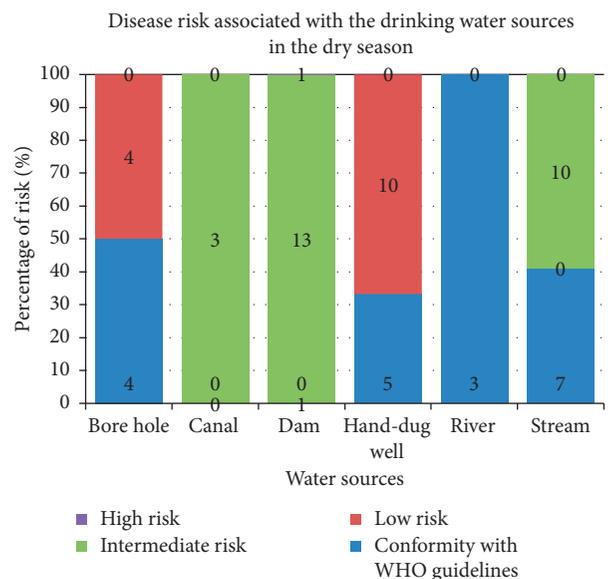


FIGURE 2: WHO disease risk levels of water sources in the wet season.

sources from hand-dug wells revealed a 67% low-risk level. Sources from boreholes showed 50% low risk and 50% conformity with standards set by the WHO. Stream water sources had a 59% intermediate risk level with 41% conformity with the WHO guidelines.

The extent of disease risk of the various water sources during the dry season is presented in Figure 3. Data from Figure 3 show an important trend. All of the sources did not show a high-risk level in the dry season, just as observed in the wet season.

All water sources from rivers, hand-dug wells, and canals had 100% intermediate risk. Boreholes had the lowest disease risk, with a 75% low-risk level and 25% compliance with WHO standards. This was followed by hand-dug wells at 80% low-risk level. On the other hand, dams showed a high disease risk.

4. Discussion

The present study evaluated the health risks associated with sources of drinking water in the Dangme West District in Ghana using *E. coli* as a measurement tool, aimed at ascertaining the paths leading to contamination of the water sources. *E. coli* is best suited as an indicator for faecal coliforms because there are fewer instances of encountering false positives [31, 32]. Also, *E. coli* enumerations are better indicators of faecal pollution than faecal coliform counts [33]. The reason being that certain strains of faecal coliform bacteria are capable of multiplying in the environs which can lead to pseudoelevation of indicator levels [34]. Additionally, recent studies have identified a direct association between the density of *E. coli* microbes in water and the prevalence of water-related gastroenteritis [35]. This, among other reasons, accounts for the use of *E. coli* counts as an indicator of water quality and, by extension, a signal of human, domestic, and natural sources of faecal contamination [18, 36–39].

In this study, we found *E. coli* counts to be predominantly higher in the wet season than the dry season (Table 1). Terrestrial wastes usually budge into most water sources in periods of excessive hailstorms or downpours, which could cause higher levels of bacteria counts during wet seasons than dry seasons [40, 41]. This could account for the higher counts observed in the water samples collected during the wet season than in the dry season. Although there were higher counts in the wet season than in the dry season, several reasons could also explain the counts observed in the dry season. First of all, water shortage and brisk multiplication of bacteria as a result of suitable conditions such as temperature throughout the dry season [42] could account for such a revelation. It is worth noting, rainfalls provided higher water volumes and large surface area for rapid microbial growth, leading to higher microbial counts and subsequently, higher extents of pollution in the wet season. Furthermore, bacterial pollutants in dumpsites, human waste, and heating systems could be drained into the different water sources during the wet season, after which rapid multiplication occurs. These sources are thus, probable sources of transmitting pathogens resulting in more critical health problems in rural neighborhoods.

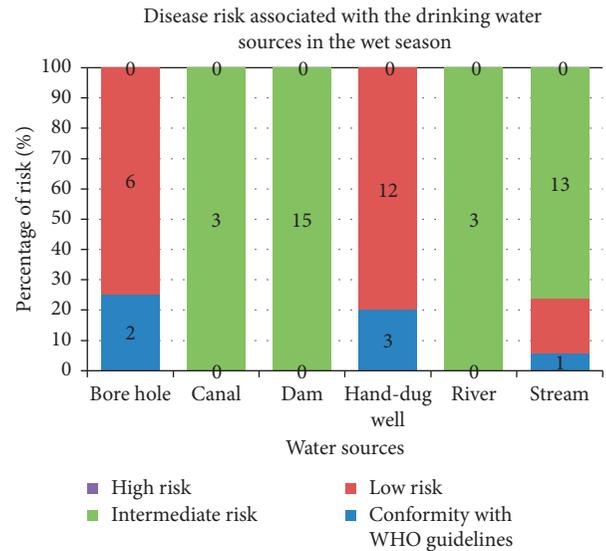


FIGURE 3: WHO disease risk levels of water sources in the dry season.

Findings of the investigation of the extent of disease risk associated with the drinking water sources in both seasons across the seasons (Figures 1 and 2) using WHO drinking water risk assessment guidelines. Water samples from the canals showed an intermediate risk level. An intermediate risk level of 59% was also observed in water obtained from streams. However, in the dry season, none of the water sources showed a high-risk level invariance to observations made in the wet season. This observation is consistent with several similar studies done in the developing world [43].

Undisputedly, anthropogenic activities and faecal releases as preeminent contributing determinants of faecal pollution could explain the high enumerations of *E. coli* in samples collected from various water sources. However, since the comparative significance of precise animals with regard to *E. coli* counts were not befittingly evaluated, it is difficult to confidently determine their contribution to the higher counts observed. Possibly, the higher counts could be associated with other related factors including animal populace density and exploitation of areas close to the water sampling sources or sites. Though the current study did not reveal anthropogenic input as a critical cause of the presence of high faecal coliforms in most water samples investigated, these results hint that human faecal contaminations could be camouflaged by other sources (such as wild and domestic animals) especially when high counts are observed.

It is important to note that a number of observations were made concerning the environmental conditions in the district during the research period, which gives insights of the pollution of the water sources. First, there were limited dumpsites for waste disposal in the district, which could account for the use of streams as alternative waste dumping sites. Moreover, it was observed that inhabitants in some communities in the district adopt unhygienic activities particularly, defecation and urination in the open and in drain ways, which eventually gets into water bodies.

Second, we observed that their groundwater sources had similar characteristics, that is, the absence of efficient physical barriers such as concrete seals, well linings, hygienic covers, and secured aseptic lids among several others capable of avoiding terrestrial runoff comprising anthropogenic and animal waste which will assuredly pollute the sources. This could be linked to the detection of *E. coli* counts in the groundwater sources in the present study. In view of the essence of physical barriers in wells, WHO [44] purported that groundwater is less susceptible to pollution as a result of the presence of barriers. Hence, contamination levels will rise if these barriers are compromised. This is particularly worrisome because, according to Chapman [45] when groundwater is polluted, it can remain polluted for many years. The inference of this result is pivotal and, as such, cannot be overelaborated.

5. Conclusions

The present study showed that most water sources analyzed were contaminated with bacterial pathogens surpassing recommended standards thereby suggesting that residents living in the rural neighborhoods of the study area are exposed to heightened risks and susceptibility to waterborne diseases or health complications. A number of practices are associated with contamination of the water sources especially anthropogenic activities. Therefore, regular tests of water quality in rural vicinities are not only advantageous but necessary.

Therefore, the government, in close collaboration with community chiefs and elders of the district, should develop a keen interest in monitoring water quality in those environments. Government and other stakeholders must also initiate education on environmental awareness including hygienic practices. Additionally, education on proper water storage practices, sanitary ways of handling water, among others, should be done to help to limit water pollution for safe consumption.

Data Availability

The data used to support the findings of this study are included within the article.

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Acknowledgments

The authors' sincere appreciation goes to laboratory technicians who assisted with the water sampling. The authors also want to acknowledge the enormous resources obtained from the thesis "Radiation Sensitivity and Molecular Characterization of Waterborne Multidrug Resistant *Escherichia coli*" towards this work.

References

- [1] H. Chitonge, A. Mokoena, and M. Kongo, "Water and sanitation inequality in Africa: challenges for SDG 6," in *Africa and the Sustainable Development Goals*, pp. 207–218, Springer, Berlin, Germany, 2020.
- [2] C. Osunla and A. Okoh, "Vibrio pathogens: a public health concern in rural water resources in sub-saharan Africa," *International Journal of Environmental Research and Public Health*, vol. 14, no. 10, p. 1188, 2017.
- [3] A. Furtatova and L. Kamenik, "Modeling features of sustainable urban development in modern conditions of water supply," *MATEC Web of Conferences*, vol. 170, p. 04002, 2018.
- [4] S. T. Odonkor and K. K. Addo, "Prevalence of multidrug-resistant *Escherichia coli* isolated from drinking water sources," *International Journal of Microbiology*, vol. 2018, Article ID 7204013, 7 pages, 2018.
- [5] K. O. Odeku and E. L. Meyer, "Socioeconomic implications of energy poverty in South African poor rural households," *Academy of Entrepreneurship Journal*, vol. 25, no. 3, pp. 1–12, 2019.
- [6] E. Bisung and S. Dickin, "Concept mapping: engaging stakeholders to identify factors that contribute to empowerment in the water and sanitation sector in West Africa," *SSM-Population Health*, vol. 9, Article ID 100490, 2019.
- [7] F. M. Gimelli, J. J. Bos, and B. C. Rogers, "Fostering equity and wellbeing through water: a reinterpretation of the goal of securing access," *World Development*, vol. 104, pp. 1–9, 2018.
- [8] C. Anthonj, B. Diekkruiger, C. Borgemeister, and T. Kistemann, "Health risk perceptions and local knowledge of water-related infectious disease exposure among Kenyan wetland communities," *International Journal of Hygiene and Environmental Health*, vol. 222, no. 1, pp. 34–48, 2019.
- [9] X. Xue, S. Cashman, A. Gaglione et al., "Holistic analysis of urban water systems in the greater cincinnati region: (1) life cycle assessment and cost implications," *Water Research X*, vol. 2, Article ID 100015, 2019.
- [10] H. Yousefi, S. Zahedi, and M. H. Niksokhan, "Modifying the analysis made by water quality index using multi-criteria decision making methods," *Journal of African Earth Sciences*, vol. 138, pp. 309–318, 2018.
- [11] M. Pal, Y. Ayele, M. Hadush, S. Panigrahi, and V. J. Jadhav, "Public health hazards due to unsafe drinking water," *Air and Water Borne Diseases*, vol. 7, no. 1, p. 1, 2018.
- [12] A. Efstratiou, J. E. Ongerth, and P. Karanis, "Waterborne transmission of protozoan parasites: review of worldwide outbreaks—an update 2011–2016," *Water Research*, vol. 114, pp. 14–22, 2017.
- [13] J. N. Njeru, "Sanitation," *The Wiley Blackwell Encyclopedia of Urban and Regional Studies*, Wiley, Hoboken, NJ, USA, pp. 1–5, 2019.
- [14] M. V. Yates, "Drinking water microbiology," in *Manual of Environmental Microbiology*, Wiley, Hoboken, NJ, USA, 4th edition, 2015.
- [15] H. T. M. Nguyen, Q. T. P. Le, J. Garnier, J.-L. J. Janeau, and E. Rochelle-Newall, "Seasonal variability of faecal indicator bacteria numbers and die-off rates in the red river basin, North Viet Nam," *Scientific Reports*, vol. 6, no. 1, Article ID 21644, 2016.
- [16] E. Thursby and N. Juge, "Introduction to the human gut microbiota," *Biochemical Journal*, vol. 474, no. 11, pp. 1823–1836, 2017.
- [17] N. R. Krieg and J. G. Holt, *Bergey's Manual of Systematic Bacteriology*, p. 964, Williams and Wilkins, Baltimore, MD, USA, 1984.
- [18] S. T. Odonkor and J. K. Ampofo, "*Escherichia coli* as an indicator of bacteriological quality of water: an overview," *Microbiology Research*, vol. 4, no. 1, p. 2, 2013.

- [19] A. Agensi, J. Tibyangye, A. Tamale, E. Agwu, and C. Amongi, "Contamination potentials of household water handling and storage practices in kirundo subcounty, kisoro district, Uganda," *Journal of Environmental and Public Health*, vol. 2019, Article ID 7932193, 8 pages, 2019.
- [20] J. M. Kayembe, F. Thevenon, A. Laffite et al., "High levels of faecal contamination in drinking groundwater and recreational water due to poor sanitation, in the sub-rural neighbourhoods of Kinshasa, Democratic Republic of the Congo," *International Journal of Hygiene and Environmental Health*, vol. 221, no. 3, pp. 400–408, 2018.
- [21] R. G. Price and D. Wildeboer, "E. coli as an indicator of contamination and health risk in environmental waters," in *Escherichia coli-Recent Advances on Physiology, Pathogenesis and Biotechnological Applications*, IntechOpen, London, UK, 2017.
- [22] H. Brüßow, "Phage therapy: the *Escherichia coli* experience," *Microbiology*, vol. 151, no. 7, pp. 2133–2140, 2005.
- [23] T. Gara, L. Fengting, I. Nhapi, C. Makate, and W. Gumindoga, "Health safety of drinking water supplied in Africa: a closer look using applicable water-quality standards as a measure," *Exposure and Health*, vol. 10, no. 2, pp. 117–128, 2018.
- [24] R. Hope, P. Thomson, J. Koehler, and T. Foster, "Rethinking the economics of rural water in Africa," *Oxford Review of Economic Policy*, vol. 36, no. 1, pp. 171–190, 2020.
- [25] TDWD, "The dangme west district of ghana 2013," 2013, <http://ghanadistricts.gov.gh/>.
- [26] Ministry of Local Government, Rural Development and Environment (MLGRDE), *Dangme West District Water and Sanitation Plan*, Government of Ghana Publication, Accra, Ghana, 2008.
- [27] World Health Organization, *Guidelines for Drinking-Water Quality. Health Criteria and Other Supporting Information*, World Health Organization, Geneva, Switzerland, 1996.
- [28] APHA, *Standard Methods for the Examination of Water and Wastewater*, American Public Health Association, American Water Works Association and Water Environment Federation, Washington, DC, USA, 19th edition, 1995.
- [29] APHA, *Standard Methods for the Examination of Water and Wastewater*, United Book Press, Inc., Baltimore, MD, USA, 20th edition, 1998.
- [30] L. M. Prescott, J. P. Harley, and A. K. Donald, *Microbiology*, W. C. Brown Publishers, McGraw-Hill Companies Inc., New York, NY, USA, 5th edition, 1996.
- [31] K. L. Quiroz, N. G. Rodriguez, S. Murinda, and M. Ibekwe, "Determination of the water quality of a constructed wetland monitoring fecal indicator bacteria," 2018.
- [32] C. Rodrigues and M. Â. Cunha, "Assessment of the microbiological quality of recreational waters: indicators and methods," *Euro-Mediterranean Journal for Environmental Integration*, vol. 2, no. 1, p. 25, 2017.
- [33] E. M. Hachich, M. D. Bari, A. P. G. Christ, C. C. Lamparelli, S. S. Ramos, and M. I. Z. Sato, "Comparison of thermotolerant coliforms and *Escherichia coli* densities in freshwater bodies," *Brazilian Journal of Microbiology*, vol. 43, no. 2, pp. 675–681, 2012.
- [34] S. L. McLellan, A. D. Daniels, and A. K. Salmore, "Clonal populations of thermotolerant enterobacteriaceae in recreational water and their potential interference with faecal *Escherichia coli* counts," *Applied and Environmental Microbiology*, vol. 67, no. 10, pp. 4934–4938, 2001.
- [35] F. Petit, O. Clermont, S. Delannoy et al., "Change in the structure of *Escherichia coli* population and the pattern of virulence genes along a rural aquatic continuum," *Frontiers in Microbiology*, vol. 8, p. 609, 2017.
- [36] S. L. McLellan and A. M. Eren, "Discovering new indicators of fecal pollution," *Trends in Microbiology*, vol. 22, no. 12, pp. 697–706, 2014.
- [37] M. A. Nkansah, N. O. Boadi, and M. Badu, "Assessment of the quality of water from hand-dug wells in Ghana," *Environmental Health Insights*, vol. 4, no. 2, pp. 7–12, 2010.
- [38] M. Odagiri, A. Schriewer, M. E. Daniels et al., "Human fecal and pathogen exposure pathways in rural Indian villages and the effect of increased latrine coverage," *Water Research*, vol. 100, pp. 232–244, 2016.
- [39] M. Pritchard, A. Edmondson, T. Craven, and T. Mkandawire, "Development of sustainable drinking water quality solutions for rural communities in the developing world," in *Sustainable Ecological Engineering Design*, pp. 259–277, Springer, Berlin, Germany, 2016.
- [40] N. Ferrer, A. Folch, G. Masó, S. Sanchez, and X. Sanchez-Vila, "What are the main factors influencing the presence of faecal bacteria pollution in groundwater systems in developing countries?" *Journal of Contaminant Hydrology*, vol. 228, Article ID 103556, 2020.
- [41] M. O. Akrong, F. K. Amu-Mensah, M. A. Amu-Mensah, H. Darko, G. N. D. Addico, and J. A. Ampofo, "Seasonal analysis of bacteriological quality of drinking water sources in communities surrounding Lake Bosomtwe in the Ashanti Region of Ghana," *Applied Water Science*, vol. 9, no. 4, p. 82, 2019.
- [42] H. I. Egilmez, A. Y. Morozov, M. R. J. Clokie, J. Shan, A. Letarov, and E. E. Galyov, "Temperature-dependent virus lifecycle choices may reveal and predict facets of the biology of opportunistic pathogenic bacteria," *Scientific Reports*, vol. 8, no. 1, p. 9642, 2018.
- [43] C. Kostyla, R. Bain, R. Cronk, and J. Bartram, "Seasonal variation of fecal contamination in drinking water sources in developing countries: a systematic review," *Science of the Total Environment*, vol. 514, pp. 333–343, 2015.
- [44] WHO, *Guidelines for Drinking-Water Quality*, WHO, Geneva, Switzerland, 3rd edition, 2006.
- [45] D. Chapman, *Water Quality Assessments. A Guide to Use of Biota, Sediments and Water in Environmental Monitoring*, E. & F. N. Spon, New York, NY, USA, 2nd edition, 1996.