

1 Antibiotic Resistance Profile of Waterborne Bacterial Isolates from Drinking Water 2 Sources.

3 Abstract

4 **Background:** Microbial contamination of drinking water remains a significant public
5 health concern, particularly in developing regions. In addition to contamination, the
6 emergence of antibiotic resistance among environmental bacterial isolates has raised
7 serious concerns regarding the potential transmission of resistant pathogens through water
8 sources.

9 **Methods:** A total of 246 drinking water samples were collected from urban and rural areas
10 of Moradabad district and nearby regions. Standard culture-based techniques were used to
11 isolate bacteria, followed by identification using conventional microbiological methods.
12 Antibiotic susceptibility testing of 100 culture-positive isolates was performed using the
13 Kirby–Bauer disc diffusion method against commonly used antibiotics.

14 **Results:** Out of 246 samples, 100 (40.7%) showed bacterial growth. The majority of
15 isolates demonstrated good sensitivity to gentamicin, ciprofloxacin, and imipenem.
16 Moderate resistance was observed against ampicillin (30.0%) and tetracycline (30.0%).
17 Isolates from rural areas and untreated water sources exhibited relatively higher resistance
18 compared to urban and treated sources. Organism-wise analysis revealed that *Escherichia*
19 *coli* showed comparatively higher resistance, while *Vibrio* spp. remained largely
20 sensitive.

21 **Conclusion:** The study highlights the presence of antibiotic-resistant bacteria in drinking
22 water sources, although overall susceptibility patterns remain favorable. Continuous
23 monitoring of water quality and rational antibiotic use are essential to prevent the
24 emergence and spread of resistance in environmental settings.

25 **Keywords:** Drinking water, Antibiotic resistance, Waterborne bacteria, *Escherichia coli*,
26 Water quality.

27 Highlights

- 28 • A considerable proportion (40.7%) of drinking water samples showed bacterial
29 contamination, indicating potential public health risk.
- 30 • Antibiotic susceptibility testing revealed overall good sensitivity, with moderate
31 resistance observed against commonly used antibiotics such as ampicillin and
32 tetracycline.
- 33 • Higher resistance patterns were observed in isolates from rural areas and untreated
34 water sources compared to urban and treated sources.
- 35 • *Escherichia coli* demonstrated relatively higher resistance among isolates, while
36 *Vibrio* spp. remained largely sensitive to most antibiotics.

37 Background

38 In recent years, increasing attention has been directed toward the emergence of antibiotic
39 resistance among environmental bacterial isolates(1). Traditionally, antibiotic resistance

40 has been associated with clinical settings; however, environmental reservoirs such as
41 water sources are now recognized as important contributors to the dissemination of
42 resistant bacteria(2). Drinking water, especially when inadequately treated or improperly
43 stored, can harbor bacteria that have been exposed to antibiotics through human,
44 agricultural, and industrial activities, thereby facilitating the selection and persistence of
45 resistant strains(3). The occurrence of antibiotic-resistant bacteria in drinking water is of
46 particular concern because it poses a dual threat: it increases the risk of infection and
47 limits the effectiveness of commonly used antimicrobial therapies(4). Moreover, these
48 bacteria can act as reservoirs of resistance genes, which may be transferred to other
49 pathogenic organisms, further amplifying the public health burden(5). Rural areas and
50 regions with limited water treatment infrastructure are especially vulnerable, although
51 contamination can also occur in urban and even packaged water sources(6). Given these
52 concerns, evaluating the antibiotic resistance patterns of bacterial isolates from drinking
53 water sources is essential for understanding the extent of the problem and for guiding
54 appropriate public health interventions. Such assessments provide valuable insights into
55 local resistance trends, help in identifying potential risks to human health, and support the
56 development of strategies for improving water quality and antimicrobial stewardship.

57 **Material and Methods**

58 The present cross-sectional laboratory-based study was conducted in Moradabad district,
59 Uttar Pradesh, India, utilizing bacterial isolates obtained from previously analyzed
60 drinking water samples. A total of 100 culture-positive isolates recovered from 246
61 samples collected from both urban and rural areas, including sources such as well water,
62 municipal tap water, rainwater harvesting systems, and bottled water, were included for
63 antibiotic susceptibility analysis. All procedures were carried out in the Department of
64 Microbiology, College of Paramedical Sciences, Teerthanker Mahaveer University,
65 Moradabad. Pure isolates were obtained through subculturing and confirmed by standard
66 microbiological and biochemical methods prior to testing. Antibiotic susceptibility testing
67 was performed using the Kirby–Bauer disc diffusion method on Mueller–Hinton agar,
68 with inoculum standardized to 0.5 McFarland turbidity. The antibiotics tested included
69 ampicillin, ciprofloxacin, ceftriaxone, gentamicin, tetracycline, and imipenem. Plates
70 were incubated at 37°C for 18–24 hours, and zones of inhibition were measured and
71 interpreted as sensitive, intermediate, or resistant according to standard guidelines. All
72 procedures were conducted under aseptic conditions with appropriate quality control
73 measures, and the resulting data were analyzed using descriptive statistics, with findings
74 expressed in terms of frequency and percentage.

75 **Result**

76 A total of 246 drinking water samples collected from urban and rural areas of Moradabad
77 district and nearby regions were analyzed, Out of 246 samples, 100 (40.7%) showed
78 bacterial growth, while 146 (59.3%) showed no growth.

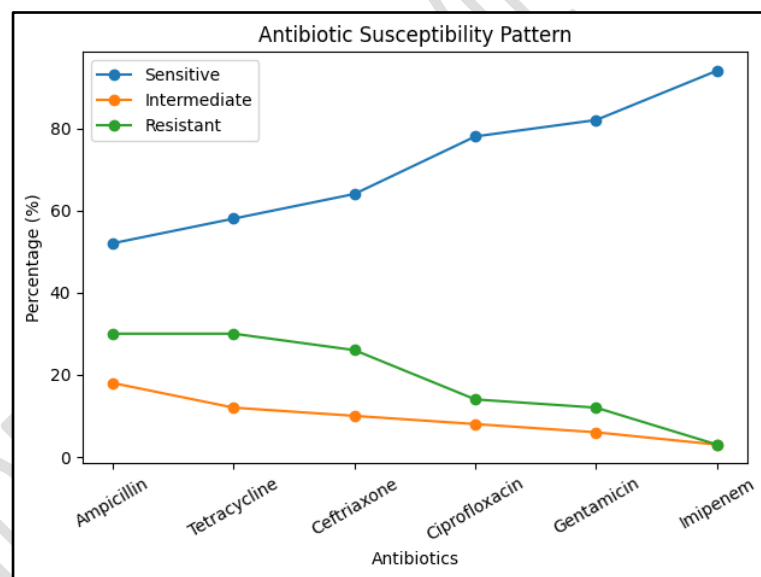
79 **Overall antibiotic susceptibility pattern**

80 The majority of isolates demonstrated good sensitivity to most of the antibiotics tested.
 81 However, moderate resistance was observed against ampicillin and tetracycline. Higher
 82 sensitivity was noted for gentamicin, ciprofloxacin, and imipenem (Table 1).

83 **Table 1. Overall antibiotic susceptibility pattern of isolates (n=100)**

Antibiotic	Sensitive n (%)	Intermediate n (%)	Resistant n (%)
Ampicillin	52 (52.0)	18 (18.0)	30 (30.0)
Tetracycline	58 (58.0)	12 (12.0)	30 (30.0)
Ceftriaxone	64 (64.0)	10 (10.0)	26 (26.0)
Ciprofloxacin	78 (78.0)	8 (8.0)	14 (14.0)
Gentamicin	82 (82.0)	6 (6.0)	12 (12.0)
Imipenem	94 (94.0)	3 (3.0)	3 (3.0)

84 Values expressed as number and percentage of total isolates tested.



85
 86 Figure 1. Antibiotic susceptibility pattern of bacterial isolates from drinking water
 87 sources.

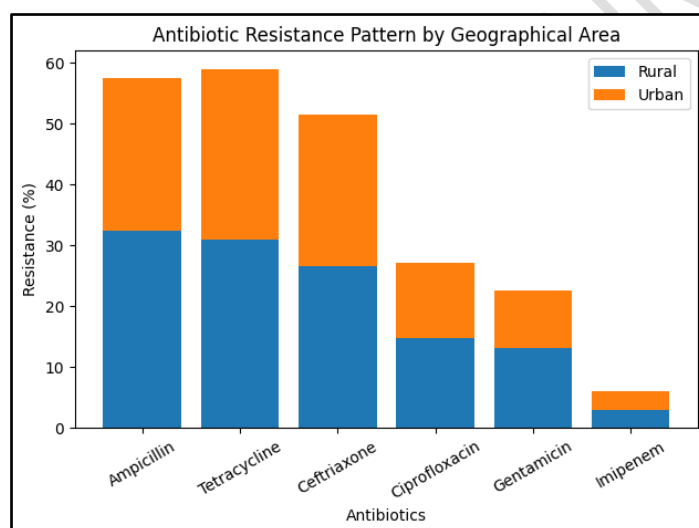
88 **Antibiotic susceptibility according to geographical area**

89 When analyzed based on geographical distribution, isolates from rural areas showed
 90 slightly higher resistance compared to urban isolates, although the overall susceptibility
 91 pattern remained similar across both groups. Ampicillin resistance was more common in
 92 rural samples, while urban isolates showed comparatively higher sensitivity to most
 93 antibiotics (Table 2).

94 **Table 2. Antibiotic resistance pattern according to geographical area**

Antibiotic	Rural Resistant n (%)	Urban Resistant n (%)
Ampicillin	22 (32.4)	8 (25.0)
Tetracycline	21 (30.9)	9 (28.1)
Ceftriaxone	18 (26.5)	8 (25.0)
Ciprofloxacin	10 (14.7)	4 (12.5)
Gentamicin	9 (13.2)	3 (9.4)
Imipenem	2 (2.9)	1 (3.1)

95 Values expressed as number and percentage of total isolates tested. Percentages calculated
 96 within each group.



97
 98 Figure 2. Antibiotic resistance pattern of bacterial isolates according to geographical area.

99 **Antibiotic susceptibility according to water source**

100 Variations in antibiotic resistance were observed among isolates from different water
 101 sources. Isolates from well water and rainwater harvesting systems showed relatively
 102 higher resistance compared to municipal tap water and bottled water. Bottled water
 103 isolates demonstrated the highest sensitivity across all antibiotics tested (Table 3).

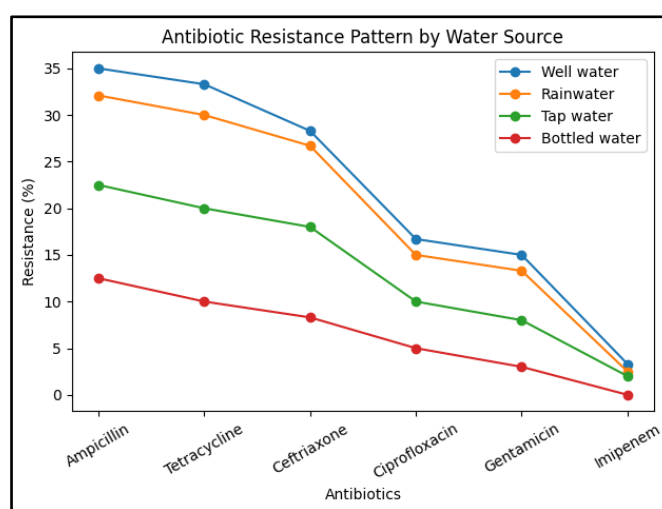
104 **Table 3. Antibiotic resistance pattern according to water source (percentage**
 105 **resistant)**

Antibiotic	Well water (%)	Rainwater (%)	Tap water (%)	Bottled water (%)
Ampicillin	35.0	32.1	22.5	12.5

Tetracycline	33.3	30.0	20.0	10.0
Ceftriaxone	28.3	26.7	18.0	8.3
Ciprofloxacin	16.7	15.0	10.0	5.0
Gentamicin	15.0	13.3	8.0	3.0
Imipenem	3.3	2.5	2.0	0.0

106

Values expressed as number and percentage of total isolates tested.



107

108 Figure 3. Antibiotic resistance pattern of bacterial isolates according to water source.

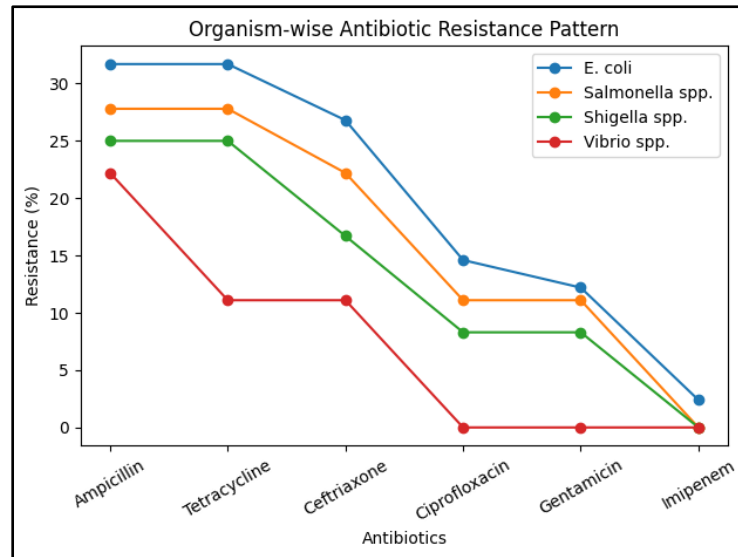
109 **Organism-wise antibiotic susceptibility pattern**

110 Among the identified isolates, *Escherichia coli* showed moderate resistance to ampicillin
 111 and tetracycline, while maintaining good sensitivity to ciprofloxacin, gentamicin, and
 112 imipenem. *Salmonella spp.* and *Shigella spp.* demonstrated similar susceptibility patterns,
 113 with relatively lower resistance levels. *Vibrio spp.* isolates were largely sensitive to most
 114 of the antibiotics tested, with only minimal resistance observed Table 4.

115 **Table 4. Organism-wise antibiotic susceptibility pattern (n, %)**

Organism	Antibiotic	Sensitive n (%)	Intermediate n (%)	Resistant n (%)
<i>Escherichia coli</i> (n=41)	Ampicillin	21 (51.2)	7 (17.1)	13 (31.7)
	Tetracycline	22 (53.7)	6 (14.6)	13 (31.7)
	Ceftriaxone	25 (61.0)	5 (12.2)	11 (26.8)
	Ciprofloxacin	31 (75.6)	4 (9.8)	6 (14.6)

	Gentamicin	33 (80.5)	3 (7.3)	5 (12.2)
	Imipenem	39 (95.1)	1 (2.4)	1 (2.4)
<i>Salmonella</i> spp. (n=18)	Ampicillin	10 (55.6)	3 (16.7)	5 (27.8)
	Tetracycline	11 (61.1)	2 (11.1)	5 (27.8)
	Ceftriaxone	12 (66.7)	2 (11.1)	4 (22.2)
	Ciprofloxacin	14 (77.8)	2 (11.1)	2 (11.1)
	Gentamicin	15 (83.3)	1 (5.6)	2 (11.1)
	Imipenem	17 (94.4)	1 (5.6)	0 (0.0)
<i>Shigella</i> spp. (n=12)	Ampicillin	7 (58.3)	2 (16.7)	3 (25.0)
	Tetracycline	8 (66.7)	1 (8.3)	3 (25.0)
	Ceftriaxone	8 (66.7)	2 (16.7)	2 (16.7)
	Ciprofloxacin	10 (83.3)	1 (8.3)	1 (8.3)
	Gentamicin	10 (83.3)	1 (8.3)	1 (8.3)
	Imipenem	11 (91.7)	1 (8.3)	0 (0.0)
<i>Vibrio</i> spp. (n=9)	Ampicillin	6 (66.7)	1 (11.1)	2 (22.2)
	Tetracycline	7 (77.8)	1 (11.1)	1 (11.1)
	Ceftriaxone	7 (77.8)	1 (11.1)	1 (11.1)
	Ciprofloxacin	8 (88.9)	1 (11.1)	0 (0.0)
	Gentamicin	8 (88.9)	1 (11.1)	0 (0.0)
	Imipenem	9 (100)	0 (0.0)	0 (0.0)



117

118 Figure 4. Organism-wise antibiotic resistance pattern of bacterial isolates from drinking
119 water sources.

120 **Discussion**

121 The present study demonstrated that 40.7% of drinking water samples showed bacterial
122 contamination, indicating a substantial burden of microbiological pollution in the study
123 area. These findings are consistent with recent studies conducted in low- and middle-
124 income countries, where unsafe water sources remain a major contributor to the
125 transmission of waterborne pathogens. **Quansah et al (2025)** conducted in Ghana reported
126 significant contamination of untreated water sources, particularly well and surface water,
127 with *Escherichia coli* and other enteric bacteria (7). Similar observations have been
128 highlighted in **Maurya et al (2025)** contemporary reviews, which emphasize that drinking
129 water systems, especially untreated or poorly managed sources, act as important
130 reservoirs for microbial contamination and antibiotic resistance (8). In the present study,
131 the majority of isolates demonstrated good sensitivity to antibiotics such as gentamicin,
132 ciprofloxacin, and imipenem, whereas moderate resistance was observed against
133 ampicillin and tetracycline. This pattern agrees with recent reports indicating that older
134 and commonly used antibiotics tend to show higher resistance due to their widespread
135 and often indiscriminate use. For instance, **Ahmed et al (2022)** have shown that
136 environmental isolates of *E. coli* from drinking water frequently exhibit resistance to
137 commonly used antibiotics, while retaining susceptibility to higher-generation drugs(9).
138 The continued effectiveness of drugs such as imipenem and gentamicin in the present
139 study may be attributed to their restricted use and limited environmental
140 exposure(10). The slightly higher resistance observed among isolates from rural areas in
141 the present study aligns with **Purohit et al (2017)**, which report increased microbial
142 contamination and resistance patterns in regions with inadequate sanitation and water
143 treatment facilities(11). **Pandey et al (2014)** have demonstrated that rural and untreated
144 water sources are more prone to contamination and may facilitate the persistence and
145 dissemination of antibiotic-resistant bacteria due to environmental exposure and lack of
146 proper infrastructure (12). Furthermore, environmental factors such as agricultural runoff,

147 improper waste disposal, and antibiotic usage in livestock have been identified as key
148 contributors to the development and spread of resistance in water systems (13).Source-
149 wise analysis in the present study revealed higher resistance among isolates from well
150 water and rainwater harvesting systems, whereas municipal tap water and bottled water
151 showed comparatively lower resistance levels. These findings are consistent with **Abebe**
152 **et al (2024)**, which have reported that treated water sources generally exhibit lower
153 microbial contamination and resistance due to effective filtration and disinfection
154 processes(14). In contrast, untreated sources remain vulnerable to contamination and act
155 as reservoirs for resistant organisms .Organism-wise analysis showed that *Escherichia*
156 *coli* exhibited moderate resistance to ampicillin and tetracycline, while *Salmonella* spp.
157 and *Shigella* spp. demonstrated relatively lower resistance levels. *Vibrio* spp. isolates
158 were largely sensitive to most antibiotics tested. These findings are comparable with
159 recent studies where *E. coli* has been identified as a key indicator organism showing
160 higher resistance due to its widespread presence and adaptability in diverse environmental
161 conditions **Quansah et al (2025)**(7). Additionally, several studies have emphasized that
162 environmental *E. coli* strains serve as important indicators for monitoring antibiotic
163 resistance trends in water sources.

164 **Conclusion**

165 The present study demonstrates that drinking water sources in the study area are
166 moderately contaminated with bacterial pathogens, with a subset exhibiting resistance to
167 commonly used antibiotics. Although the overall susceptibility pattern remains favorable,
168 the presence of resistance, particularly to ampicillin and tetracycline, indicates emerging
169 environmental pressure likely associated with widespread antibiotic use. Higher
170 resistance observed in rural and untreated water sources further emphasizes the role of
171 environmental and infrastructural factors in shaping resistance patterns. These findings
172 underscore the need for regular surveillance of drinking water quality, improvement in
173 water treatment and sanitation practices, and promotion of rational antibiotic usage to
174 minimize the risk of dissemination of resistant bacteria through environmental pathways.

175 **Declarations:**

- 176 • **Ethical Approval:** Not Applicable
- 177 • **Informed Consent:** Informed consent was obtained where applicable.
- 178 • **Conflict of Interest:** The authors declare no conflict of interest.
- 179 • **Funding:** The authors received no external funding for this study.
- 180 • **Data Availability:** Data supporting the findings of this study are available from the
181 corresponding author upon reasonable request.
- 182 • **Author Contributions:** SB, conceptualized the study, conducted laboratory analysis,
183 performed data interpretation, and drafted the manuscript. MF, supervised the research
184 work, contributed to study design, critically revised the manuscript, and approved the
185 final version for publication.

186 **Acknowledgment:** The authors acknowledge the support of the Microbiology
187 Laboratory, College of Paramedical Sciences, Teerthanker Mahaveer University, for

188 providing facilities and technical assistance. We also thank the faculty members for their
189 guidance and support.

190 Reference

- 191 1. Larsson DGJ, Flach CF. Antibiotic resistance in the environment. *Nat Rev Microbiol.*
192 2022 May;20(5):257–69. doi:10.1038/s41579-021-00649-x
- 193 2. Selvarajan R, Obize C, Sibanda T, Abia ALK, Long H. Evolution and Emergence of
194 Antibiotic Resistance in Given Ecosystems: Possible Strategies for Addressing the
195 Challenge of Antibiotic Resistance. *Antibiotics (Basel).* 2022 Dec 24;12(1):28.
196 doi:10.3390/antibiotics12010028 PubMed PMID: 36671228; PubMed Central
197 PMCID: PMC9855083.
- 198 3. Kristanti RA, Hadibarata T, Syafrudin M, Yılmaz M, Abdullah S. Microbiological
199 Contaminants in Drinking Water: Current Status and Challenges. *Water Air Soil*
200 *Pollut.* 2022 Jul 22;233(8):299. doi:10.1007/s11270-022-05698-3
- 201 4. Ashbolt NJ. Microbial Contamination of Drinking Water and Human Health from
202 Community Water Systems. *Curr Environ Health Rep.* 2015;2(1):95–106.
203 doi:10.1007/s40572-014-0037-5 PubMed PMID: 25821716; PubMed Central
204 PMCID: PMC4372141.
- 205 5. Shah A, Arjunan A, Baroutaji A, Zakharova J. A review of physicochemical and
206 biological contaminants in drinking water and their impacts on human health. *Water*
207 *Science and Engineering.* 2023 Dec 1;16(4):333–44. doi:10.1016/j.wse.2023.04.003
- 208 6. Niu L, Liu W, Juhasz A, Chen J, Ma L. Emerging contaminants antibiotic resistance
209 genes and microplastics in the environment: Introduction to 21 review articles
210 published in CREST during 2018–2022. *Critical Reviews in Environmental Science*
211 *and Technology.* 2022 Dec 2;52(23):4135–46. doi:10.1080/10643389.2022.2117847
- 212 7. Quansah KE, Ahmed H, Thekkur P, Hedidor GK, Adomako LAB, Banu RA, et al.
213 Antibiotic-Resistant Bacteria in Drinking Water Across Twelve Regions of Ghana:
214 Strengthening Evidence for National Surveillance. *Trop Med Infect Dis.* 2025 Oct
215 14;10(10):291. doi:10.3390/tropicalmed10100291 PubMed PMID: 41150368;
216 PubMed Central PMCID: PMC12567938.
- 217 8. Shubham Kumar Maurya, Himanshu Kumar, Kajal Pal, Chandan Kumar, Harsh Sen
218 Yadav, Sagar Gupta, et al. In Vitro Characterization Of Antimicrobial Resistance And
219 Biochemical Signatures Of Pathogens From Environmental Reservoirs In Pakbara,

220 Moradabad, Uttar Pradesh, India. International Journal of Environmental Sciences.
221 2025 Aug 11;1858–64. doi:10.64252/dwddb56

222 9. Ahmed H, Zolfo M, Williams A, Ashubwe-Jalemba J, Tweya H, Adeapena W, et al.
223 Antibiotic-Resistant Bacteria in Drinking Water from the Greater Accra Region,
224 Ghana: A Cross-Sectional Study, December 2021-March 2022. Int J Environ Res
225 Public Health. 2022 Sep 28;19(19):12300. doi:10.3390/ijerph191912300 PubMed
226 PMID: 36231603; PubMed Central PMCID: PMC9566567.

227 10. Mussema A, Admasu D, Bawore S, Abdo R, Seid A. BACTERIAL PROFILE,
228 ANTIMICROBIAL RESISTANCE, AND FACTORS ASSOCIATED WITH
229 URINARY TRACT INFECTION AMONG PREGNANT WOMEN AT HOSANNA
230 TOWN HEALTH FACILITIES, CENTRAL ETHIOPIA. Georgian Med News. 2023
231 Sep;(342):113–21. PubMed PMID: 37991965.

232 11. Purohit MR, Chandran S, Shah H, Diwan V, Tamhankar AJ, Stålsby Lundborg C.
233 Antibiotic Resistance in an Indian Rural Community: A ‘One-Health’ Observational
234 Study on Commensal Coliform from Humans, Animals, and Water. Int J Environ Res
235 Public Health. 2017 Apr;14(4):386. doi:10.3390/ijerph14040386 PubMed PMID:
236 28383517; PubMed Central PMCID: PMC5409587.

237 12. Pandey PK, Kass PH, Soupir ML, Biswas S, Singh VP. Contamination of water
238 resources by pathogenic bacteria. AMB Express. 2014 Jun 28;4:51.
239 doi:10.1186/s13568-014-0051-x PubMed PMID: 25006540; PubMed Central
240 PMCID: PMC4077002.

241 13. Samreen, Ahmad I, Malak HA, Abulreesh HH. Environmental antimicrobial
242 resistance and its drivers: a potential threat to public health. Journal of Global
243 Antimicrobial Resistance. 2021 Dec 1;27:101–11. doi:10.1016/j.jgar.2021.08.001

244 14. Abebe TA, Gebreyes DS, Abebe BA, Yitayew B. Antibiotic-resistant bacteria and
245 resistance-genes in drinking water source in north Shoa zone, Amhara region,
246 Ethiopia. Front Public Health. 2024 Sep 6;12. doi:10.3389/fpubh.2024.1422137

247