

# Occurrence of Chlorine Resistant Bacteria in Drinking Water Filtration Plants of Rawalpindi City, Pakistan

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**Abstract:** Waterborne bacterial infections are mainly caused by the direct transmission of pathogenic bacteria to the host through DW. Chlorination of DW is critical to prevent the water supplies from bacterial pathogens. In present study, microbiological quality of DW provided through filtration plants to Rawalpindi City was investigated. Experimentations were designed to screen bacterial tolerance to the added Chlorine, and to subsequent study the antibiotic sensitivity of isolated bacteria. A total of 107 water samples were collected. Out of which, 57.95% were found satisfactory for human consumption. pH determination revealed that 18.70% of the samples has pH in the range of 8.0 to 8.9. TDS analysis showed 14.01% of samples above the standard HDL (500 mg/L). Furthermore, 13.34% of the samples showed the highest load (351 to 1800 CFU/100mL) of fecal coliforms. Ciprofloxacin and Tetracycline were found to be the most effective antibiotics against the isolated pathogens. These bacterial strains were also able to tolerate 1mg/L of Chlorine. Present study revealed the presence of Chlorine and antibiotic resistant bacteria in DW, which poses a great health risk to the consumers. In conclusion, it is recommended to boil the DW, as it effectively kills all the bacteria in addition to chlorination.

**Keywords:** Chlorination, Chlorine Resistance, Drinking Water, Filtration Plants, Waterborne Bacteria

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## 1. Introduction

Water is an essential component for all forms of life and is vital to sustain life on earth [1, 2]. Accessibility and availability of fresh and clean water is a key to sustainable development and an essential element in health [3]. Water free from microorganisms especially pathogenic bacteria is a fundamental requirement for human life, personal hygiene and for drinking. However, all available water is not fit for human consumption [2, 4–6]. As per WHO reports the mortality rate of water associated diseases exceeds 5 million people per year. Among these, > 50% of intestinal infections are caused by the consumption of drinking water (DW) possessing microbial contamination [5, 7].

Pollution of water occurs when water sources such as ground water, ponds, rivers, and water distribution channels are contacted by pathogenic bacteria, viruses, fungi, protozoans and industrial toxic chemicals [8]. Coliform bacteria in DW are the indication of animal and human fecal

contamination [9]. Water contamination rates are highest in the rainy seasons when microorganism rapidly grow and disperse in the water bodies. Lack of improper water treatment, facilities attribute to the spread of water borne diseases. In addition to this, DW distribution system is very poor as water channels and drainage lines run in parallel which hinder the provision of the good quality water [10, 11].

As far as DW quality is concerned, Pakistan ranks at number 80 among 122 nations. Thus proving that pollution of DW is one of the major threats for the public health in Pakistan. DW sources, both surface and groundwater are found contaminated with coliforms, toxic metals and pesticides throughout the country. In other words microbial and chemical pollutants are the main sources responsible exclusively and/or in combination for various public health problems [12, 13]. This is also supported by a study conducted by Pakistan Council for Research in Water Resources covering 64 Tehsils in four provinces. The results of DW quality from 23 major cities revealed that the water resources of Pakistan are 27 –

100% contaminated with bacteria [13, 14].

Chlorination is practiced at most of the filtration plants as a mean of water disinfection, before it is supplied to the public via distribution network [15, 16]. Low cost and effectiveness of chlorine against pathogenic microorganisms has made it a chemical of choice in many countries including Pakistan [17]. Although, addition of chlorine is a common practice however, it is not sufficient to ensure the safety of water. Detection of any disinfectant residue reduces the microorganism count and frequency at the consumer's tap. Furthermore, DW Chlorine residues have long been recognized as an excellent indicator for accessing water quality in the distribution networks [16–18]. Therefore, the maintenance of chlorine residues is needed at all points in any distribution system [16, 17, 19].

The current study was designed to assess the quality of chlorinated drinking water supplied to the different parts of the city through the bacterial tolerance to the added Chlorine.

## 2. Materials and Methods

### 2.1. Study Area

Rawalpindi is the fourth largest city of Pakistan and is the third largest metropolitan city of Pakistan. It is situated on the Pothohar Plateau, adjacent to Islamabad, the capital city of Pakistan. It has GPS coordinates of 33°37' 33.8052" N and 73°4' 17.1912" E and elevation of 508 m (1,667 ft). Population of this city according to the 2017 census is 2,098,231. The weather is highly variable due to the proximity of the city to the foothills of Himalayas. The city features a humid subtropical climate (Köppen: Cwa) with hot and wet summers, a cooler and drier winter. It experiences on an average 91 thunderstorms per year. Strong windstorms are frequent in the summer during which wind gusts 176 km/h (109 mph) have been reported by Pakistan Meteorological Department. Such conditions (wind/thunder storm) often lead to damage of infrastructure. The main source of water for the city are ground and surface water, Khanpur and Rawal Dams, and 270 tube wells [20, 21].

### 2.2. Sample Collection

A total of 107 water samples were randomly collected from different Water and Sanitation Agency (WASA) filtration plants. Water sampling was carried out during October, 2016 to May, 2017 in the presence of WASA inspector. Approximately 100 mL of water was collected in sterile 300 mL glass bottles. Taps were turned on full and water was allowed to flow for at least 1 minute. Subsequently, taps were sterilized with cotton soaked in spirit, water was again allowed to flow for few seconds. Sample bottles were filled with gentle flow of water and then capped immediately. Collected samples were then transported to the laboratory for microbiological processing.

### 2.3. Media & Chemicals

Unless otherwise stated, all the bacteriological media, and antibiotic discs were purchased from Oxoid, Basingstoke, Hampshire, UK. Chemicals used for the bacterial

identification were obtained from Merck, Darmstadt, Germany.

### 2.4. Physiochemical Analysis

All collected samples were processed for determination of pH and total dissolved solids (TDS) using pH meter (Oakton, U- Tech Instruments Malaysia) and observations were recorded for further analysis.

### 2.5. Bacteriological Analysis

#### 2.5.1. Enumeration of Coliforms by Most Probable Number (MPN) Test

In order to screen the presence of total and fecal coliforms, collected water samples were processed through MPN method. The test procedure included three phases namely presumptive, confirmative and completed test.

*Presumptive test:* Each water sample was divided into 3 tubes of each containing 10, 1 and 0.1 mL respectively. 10 mL sample was inoculated in double strength Lactose broth (LB-2X) while, 1 and 0.1 mL of the samples were inoculated in the single strength Lactose broth (LB-1X). All the inoculated tubes were incubated at 37°C for 24 – 48 hrs. Media tubes showing the presence of growth in terms of turbidity (with or without gas) were subjected to the confirmed test.

*Confirmed test:* All test tubes found turbid were gently shaken and one loopful of this culture was inoculated on to EMB (Eosin Methylene Blue) agar. Media plates were incubated at 37°C for 24 hrs. Subsequently, plates were observed for the growth and development of green metallic sheen.

*Completed test:* To confirm the presence of *E. coli*, single colony showing green metallic sheen was selected. Half of the selected colony was aseptically inoculated in Lactose broth containing Durham tube to reconfirm the positive lactose fermentation. While Nutrient agar slants inoculated with rest of the selected colony and incubated at 37°C were further proceeded for the identification of isolates. Finally, the isolated organisms were identified by performing standard microbiological methods (Cellular morphology including Gram reaction & standard biochemical tests).

#### 2.5.2. Detection of Other Gram Negative Bacteria

For the isolation of other Gram negative pathogens, 0.1 mL of each water sample was separately inoculated on the sterile MacConkey agar plates. All inoculated plates were incubated at 37°C for 24 hrs. Bacterial identification was carried out by performing standard microbiological methods including Gram reaction and biochemical tests. All experiments were run in duplicates.

### 2.6. Antimicrobial Susceptibility Testing

Antimicrobial susceptibility testing was done on Muller-Hinton agar using disk diffusion technique according to Kirby–Bauer method [22]. Antimicrobial agents tested were: Augmentin (30µg), Cefixime (5µg), Ceftriaxone (30

$\mu\text{g}$ ), Tetracycline (30 $\mu\text{g}$ ), Ciprofloxacin (5 $\mu\text{g}$ ), Sulphamethox/trimethoprim (25 $\mu\text{g}$ ), Meropenem (10 $\mu\text{g}$ ), Oxacillin (30  $\mu\text{g}$ ), Nalidixic Acid (30  $\mu\text{g}$ ) and Chloramphenicol (30  $\mu\text{g}$ ). After 18 h incubation at 37°C, the size of the zone of inhibition was measured and interpreted by comparing with the standard antibiotic sensitivity chart to determine their resistance patterns. All experiments were run in duplicates.

### 2.7. Screening of Chlorine Resistant Bacteria

To check the chlorine tolerance of the bacteria isolated from aforementioned samples. All bacterial strains were separately inoculated on nutrient agar supplemented with pure grade Sodium hypochlorite (Daejung, Korea) (1 mg/L) plates. Incubation of these plates was carried out at 37°C for 24 hrs. Subsequently, plates were observed for bacterial growth. The experiment was run in triplicates. Chlorine concentration used in this study was the actual concentration used to disinfect water before its supply to households of the city.

### 2.8. Statistical Analysis

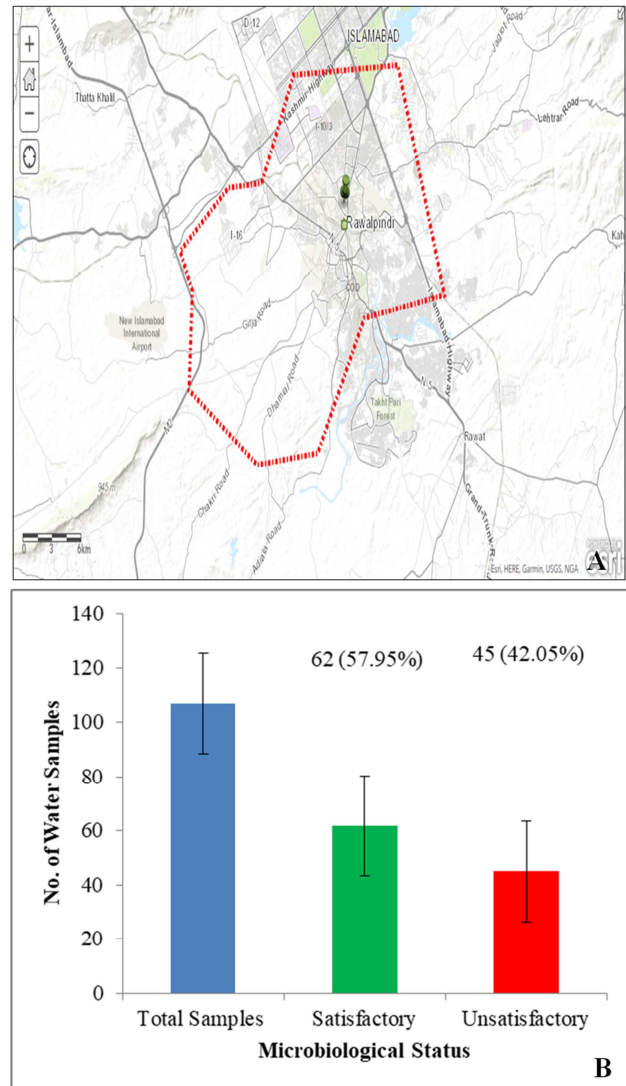
Data obtained was analyzed through calculation of mean, proportions, percentages and graphs and tables were prepared for data visualization. The bacteriological counts recorded were compared with the WHO guidelines for drinking water.

## 3. Results and Discussion

### 3.1. Collection and Water Potability Status

Rawalpindi is the third largest city of Pakistan, located near the capital city of Islamabad, in the province of Punjab. Sufficient quantity of DW supply in this city is the responsibility of WASA. WASA (Rawalpindi) was created with the mandate to provide sufficient quantity of quality DW, sewerage treatment services and for maintaining drainage facilities of the city. There are two main sources of DW available in the city i.e. ground water and surface water. Ground water is obtained with the help of 270 tube wells, supplying approximately 26 million gallon of water per day (MGD) to the city. Whereas surface water is supplied from Khanpur Dam through Sangjiani Water Treatment Plant and Rawal Lake through a 23-MGD Rawal Lake Filtration Plant [13, 23].

The study reported herein is the first study in Pakistan that focuses on the possible applications of excessive usage of Chlorine to disinfect DW directly on bacteria residing in it. It further provides an overview of the impacts on human health due to consumption of Chlorinated water. These include the infections caused by Chlorine resistant bacteria and other possible biochemical manifestations in human host caused by continuous intake of Chlorine rich water. Literature search revealed various studies on both Gram positive and Gram negative bacteria isolated from DW [14-18]. Contrary to that, current study was focused on the Gram negative (Gram -ve) bacteria including total and fecal coliforms.



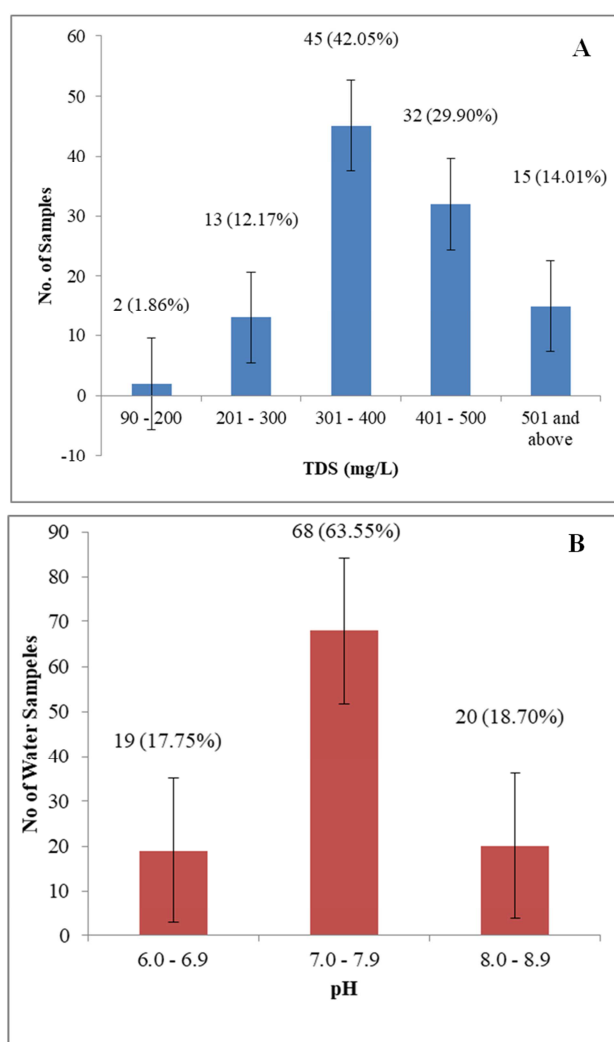
**Figure 1.** (A) Location map of Rawalpindi city for water sample collection; (B) Microbiological Status of the collected water samples.

To check the quality of DW, a total of 107 filtration plants of Rawalpindi city were considered (Figure 1A). The map for sampling sites was drawn using ArcGIS tools. Out of these, 62 samples (57.95%) were found satisfactory for human consumption. However, 45 samples (42.05%) were found microbiologically contaminated either with fecal coliforms and/or with other Gram -ve bacteria, thus unfit for the drinking purpose (Figure 1B). The results of the present study are in line with previous findings in which bacterial contamination of DW was reported in Rawalpindi and Islamabad cities [13, 23-25]. Thus, the present study highlights the significance of the microbiological examination of the DW before human consumption. For the reason that, fecal coliforms and most of the other Gram -ve bacteria isolated in present study are well-known principle pathogens of human gastrointestinal tract (GIT) infections. These results obtained can be further explained by the fact that these water borne pathogens are not only responsible for causing infections in some individuals. Nevertheless, there are reports for sudden increase in number of coliform bacteria in DW

distribution systems in various parts of the world that have led to the occurrence of an infectious outbreak in human population [26-29].

### 3.2. Physiochemical Quality

This set of experiment was conducted to evaluate the pH of the collected water samples. Results of this experiment indicated that 17.75% water samples showed pH ranging 6.0 to 6.9, whereas 68 samples (63.55%) showed pH from 7.0 to 7.9. It is interesting to mention that 18.70% of samples yielded pH in the range of 8.0 to 8.9 (Figure 2B). Therefore, the later cohort of water samples was found beyond the highest desirable limit (HDL) set by WHO i.e. 8.5.



**Figure 2.** (A) Total dissolved solids (TDS) profile of water samples; (B) pH profile of water samples.

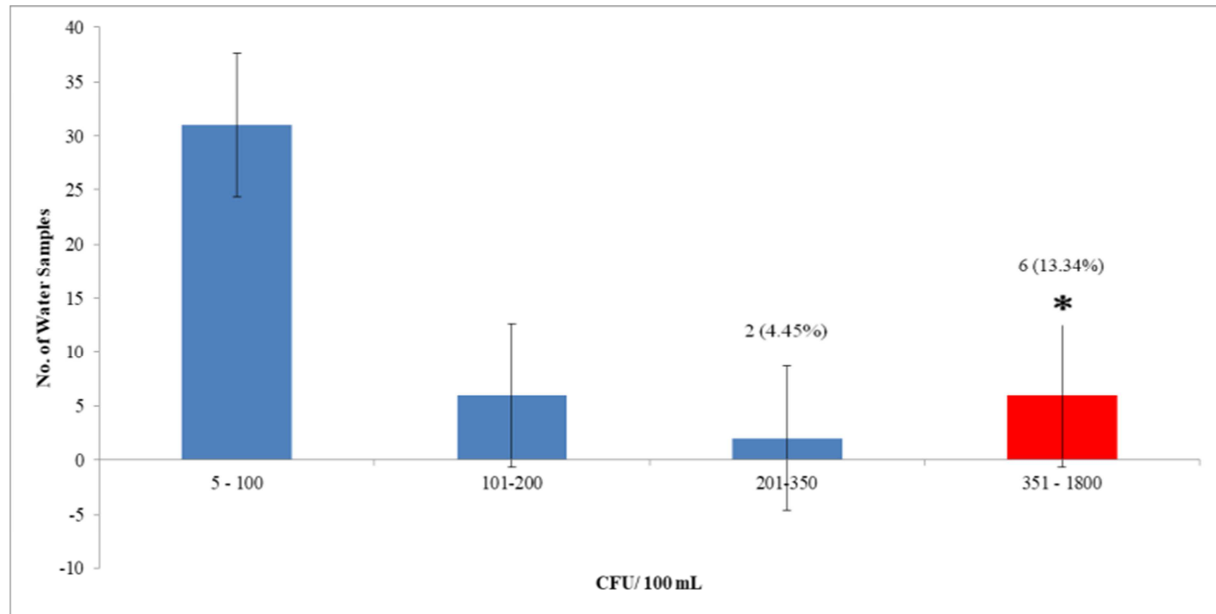
In order to determine the TDS in the collected DW samples, another set of experiment was performed. Out of 107 collected water samples, 15 (14.01%) samples were found above the standard HDL of 500 mg/L (Figure 2A). In contrast, 2 (1.86%) of the samples yielded the lowest TDS values i.e. 90 to 200 mg/L. Furthermore, 201 to 300 mg/L and 401 to 500 mg/L of

TDS were found in 13 (12.17%) and 32 (29.90%) water samples respectively. Interestingly, 45 (42.05%) of the samples indicated the presence of TDS as 301 to 400 mg/L.

It is striking that, 14.01% of the DW samples yielded total TDS ranging > 500 mg/L which is beyond the standard HDL of 500 mg/L (Figure 2A). In addition, the results of the physiochemical analysis highlighted that 18.70% of DW samples has elevated pH i.e. 8.6 to 8.9 which is also higher than WHO standard limit i.e. 8.5 (Figure 2B). The possible reason for the elevations in the physiochemical aspects of DW may be due to bacterial growth and multiplication in it. It is worth to mention herein that temperature monitoring at these filtration plants was beyond the scope of the present study, nevertheless temperature, in combination with TDS, pH and other physical factors like (infrastructure, water pipeline conditions and rusting) also strongly aid in the survival of water borne pathogens in DW. Evidence also suggest that DW physiochemical imbalance also leads to the corrosion of water mains and interior plumbing systems. This happens when biofilms influence the local chemistry of distribution system surfaces and leads to physiochemical imbalance [30]. Consequently, failure to control to these can result in the survival of different bacterial pathogens in the systems [23, 30]. Research studies have shown that biofilms developed more rapidly on pipes composed of iron, it hence proves that pipe material could be a source of nutrition for certain bacteria in DW [41, 42]. Growth of *Lagionella spp* was found to be associated with rubber gaskets used in water mains [43]. Microorganisms such as *Pseudomonas aeruginosa*, *Chromobacter spp*, *Enterobacter aerogenes* and *Klebsiella pneumoniae* have dependent growth pattern on joint packing materials in water distribution system [44]. In the present study, it can be concluded that bacteria may be using the elevated solids as nutrients to multiply in the suitable pH. Moreover, it definitely adversely affects taste, odor and appearance of the DW [32].

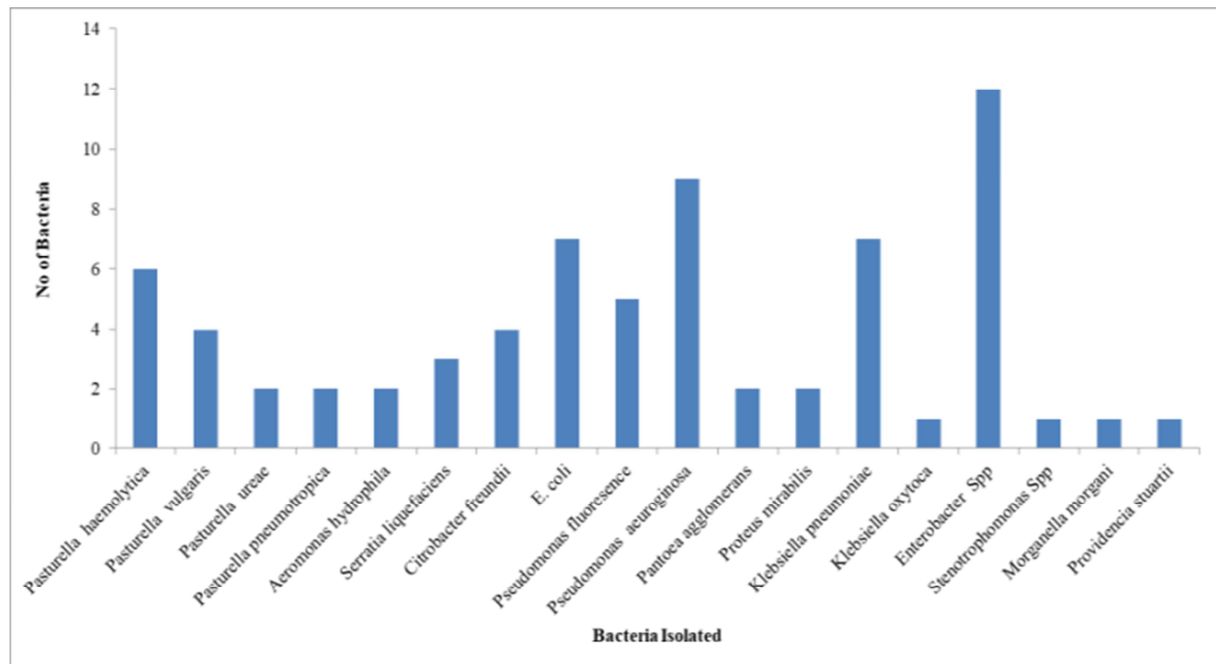
### 3.3. Bacteriological Quality

In this experiment, water samples were evaluated for the presence of fecal coliforms and other Gram negative bacteria. Results of MPN test indicated that 45 (42.05%) samples showed the presence of fecal contamination (Figure 1B). Out of these 45 contaminated water samples, 31 (68.89%) samples yielded the bacterial load as 5 to 100 CFU/ 100mL. Whereas, 6 (13.34%) samples showed coliforms ranged from 101 to 200 CFU/ 100mL (Figure 3). Further analysis of the results of the present study indicated that, 2 (4.45%) of DW samples revealed the presence of coliforms bacteria ranging from 201 to 350 CFU/100mL. It is significant to mention herein that 6 (13.34%) of the DW samples of Rawalpindi city filtration plants yielded the highest load of the fecal coliforms in range of 351 to 1800 CFU/100mL (Figure 3). Nonetheless, such high numbers of coliforms bacteria significantly highlight the extreme health concern for the population of the city.



**Figure 3.** Quantification of fecal contamination (CFU/100ml) of water samples.

Furthermore, the most frequently isolated bacterial strains (i.e. both coliforms and other Gram -ve bacteria) are summarized in Figure 4.



**Figure 4.** Pictorial representation of the total bacterial strains isolated.

The MPN analysis showed diversified response. It was interesting to note that 13.34% of DW samples showed the highest bacterial load i.e. ranging from 351 to 1800 CFU/mL (Figure 3). Whereas, other DW sample cohorts revealed the presence of fecal coliforms load in the range from 5 to 350 CFU/mL. Thus, it is justified to conclude that elevated number of bacterial pathogen in DW is an extreme health concern for the consumers. Although, no visible microbial contamination was detected at any filtration plant taps during the sampling period; however, results of the MPN experiment significantly

highlights the deterioration of the DW quality as it flowed through the distribution network. The detection of fecal coliforms in DW is mainly associated with the unhygienic conditions, parallel lining of the DW and sewerage and insufficient water supply. Our findings are in accordance to other reports [6, 12, 13, 23-25, 31] that indicated the presence of fecal coliforms in combination with other Gram -ve bacteria in the DW. Results of the present study are in part consistent with the previous research, which reported fecal contamination in distribution channels of Rawal Lake [32] and



other water distribution channels and treatment plants of Rawalpindi city [33, 34].

The culturable count of other Gram –ve water borne bacteria was enumerated additionally (Figure 4). Further analysis of the results directed us to report the presence of a number of water borne bacteria that may also serve as possible pathogens to human host. It was significant to note that, *Enterobacter Spp*, *Pseudomonas aeruginosa*, *Klebsiella pneumonia* and *E. coli* are the most isolated bacterial strains from the DW samples. These results can be explained by the fact that aforementioned bacteria are known to cause vomiting, diarrhea, dysentery, gastroenteritis, other GIT illnesses and kidney problems [35-37]. Therefore, consumption of DW contaminated with these pathogens poses serious health risk to the consumers.

### 3.4. Antimicrobial Susceptibility Profile

The above-mentioned bacterial isolation prompted us to evaluate the antibiotic resistance profile of the isolated DW borne bacterial pathogens, results of which are summarized in Table 1. This analysis indicated that, *Pasturella ureae*, *S. liquefacientes*, *Citrobacter freundii*, *Proteus mirabilis*,

*Klebsiella pneumonia*, *Morganella morganii* and *Providencia stuartii* were found moderately resistant to Ceftriaxone. However, >80% of isolated bacterial pathogens were found resistant to Cefixime. In contrast, Sulphamethox/trimethoprim was effective against *Pasturella vulgaris*, *Aeromonas hydrophila*, *S. liquefacientes*, *Proteus mirabilis*, *Stenotrophomonas Spp* and *Providencia stuartii*. Augmentin showed inhibitory activity specifically against bacterial spp of *Aeromonas hydrophila*. Oxacillin and Meropenem showed moderate inhibition of *Citrobacter freundii*, *E. coli* and *Ps. fluorescences*. It is worth mentioning here that Ciprofloxacin and Tetracycline were the most effective antibiotics against the isolated waterborne pathogens. In contrast, Nalidixic acid was ineffective against *P. pneumotropica*, *E. coli*, *Pantoea agglomerans* and *Providencia stuartii*. Chloramphenicol, on the other hand revealed the sensitivity against *Pasturella ureae*, *Pseudomonas fluorescences*, *Klebsiella oxytoca*, *Enterobacter Spp* and *Morganella morganii* only (Table 1). Thus, these results highlight that Ciprofloxacin and Tetracycline are drug of choice against the infections caused by any of the bacterial pathogen isolated in present study.

Table 1. Antibiotic sensitivity profile of bacterial strains isolated from water.

|                           | <i>Pasturella haemolytica</i> | <i>Pasturella vulgaris</i> | <i>Pasturella ureae</i> | <i>P. pneumotropica</i> | <i>Aeromonas hydrophila</i> | <i>S. liquefacientes</i> | <i>Citrobacter freundii</i> | <i>E. coli</i> | <i>Ps. fluorescences</i> |
|---------------------------|-------------------------------|----------------------------|-------------------------|-------------------------|-----------------------------|--------------------------|-----------------------------|----------------|--------------------------|
| Total No of Isolates (n)  | 06                            | 04                         | 02                      | 02                      | 02                          | 03                       | 04                          | 07             | 05                       |
| Ceftriaxone               | 100                           | 100                        | 50                      | 100                     | 100                         | 34                       | 25                          | 100            | 100                      |
| Cefixime                  | 67                            | 0                          | 0                       | 0                       | 0                           | 0                        | 0                           | 0              | 40                       |
| Sulphamethox/trimethoprim | 34                            | 100                        | 50                      | 0                       | 100                         | 100                      | 75                          | 85             | 0                        |
| Augmentin                 | 34                            | 0                          | 0                       | 0                       | 100                         | 34                       | 25                          | 0              | 0                        |
| Oxacillin                 | 0                             | 0                          | 0                       | 0                       | 0                           | 0                        | 25                          | 28             | 0                        |
| Ciprofloxacin             | 100                           | 100                        | 100                     | 100                     | 100                         | 100                      | 75                          | 85             | 100                      |
| Tetracycline              | 50                            | 100                        | 100                     | 100                     | 100                         | 100                      | 75                          | 85             | 100                      |
| Meropenem                 | 0                             | 0                          | 0                       | 0                       | 0                           | 0                        | 0                           | 0              | 20                       |
| Nalidixic Acid            | 17                            | 100                        | 50                      | 0                       | 100                         | 100                      | 75                          | 0              | 20                       |
| Chloramphenicol           | 0                             | 0                          | 50                      | 0                       | 0                           | 0                        | 0                           | 0              | 100                      |

Table 1. Continued.

|                           | <i>Ps. aeruginosa</i> | <i>Pantoea agglomerans</i> | <i>Proteus mirabilis</i> | <i>K. pneumoniae</i> | <i>K. oxytoca</i> | <i>Enterobacter Spp</i> | <i>Stenotrophomonas Spp</i> | <i>Morganella morganii</i> | <i>Providencia stuartii</i> |
|---------------------------|-----------------------|----------------------------|--------------------------|----------------------|-------------------|-------------------------|-----------------------------|----------------------------|-----------------------------|
| Total No of Isolates (n)  | 09                    | 02                         | 02                       | 07                   | 01                | 12                      | 01                          | 01                         | 01                          |
| Ceftriaxone               | 100                   | 100                        | 50                       | 71                   | 100               | 100                     | 100                         | 0                          | 0                           |
| Cefixime                  | 0                     | 0                          | 0                        | 0                    | 0                 | 0                       | 0                           | 0                          | 0                           |
| Sulphamethox/trimethoprim | 23                    | 0                          | 100                      | 71                   | 0                 | 59                      | 100                         | 0                          | 100                         |
| Augmentin                 | 0                     | 0                          | 0                        | 29                   | 0                 | 9                       | 0                           | 0                          | 0                           |
| Oxacillin                 | 0                     | 0                          | 0                        | 0                    | 0                 | 0                       | 0                           | 0                          | 0                           |
| Ciprofloxacin             | 100                   | 100                        | 100                      | 100                  | 100               | 100                     | 100                         | 100                        | 100                         |
| Tetracycline              | 13                    | 0                          | 100                      | 100                  | 100               | 25                      | 100                         | 100                        | 100                         |
| Meropenem                 | 0                     | 0                          | 0                        | 0                    | 0                 | 9                       | 0                           | 0                          | 0                           |
| Nalidixic Acid            | 100                   | 0                          | 100                      | 15                   | 100               | 75                      | 100                         | 100                        | 0                           |
| Chloramphenicol           | 0                     | 0                          | 0                        | 15                   | 100               | 100                     | 0                           | 100                        | 0                           |

\*Numbers represent percent susceptible

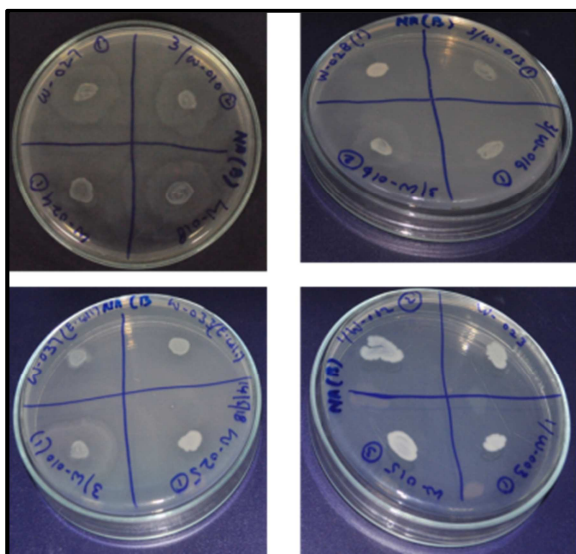
Interestingly, our results revealed that Ciprofloxacin, Ceftriaxone and Tetracycline can effectively kill all the reported water borne bacteria isolated in this study. In contrast, all the rest of the antibiotics either showed moderate or no activity against these bacteria. Our findings are in part

consistent with another study which reported the sensitivity of water borne thermo tolerant *E. coli* against Nalidixic acid and resistance against Augmentin [38]. The possible explanation of these findings could be the excessive and uncontrolled use of antibiotics significantly contributes in the dissemination of

antibiotic residues in fresh water environment. Nonetheless, the antibiotic sensitivity pattern reported here uncovers a potential link and a greater risk of resistance genes transfer to other pathogenic bacteria. Taken together, this may worsen the diagnosis and treatment of infections caused by aforementioned bacteria.

### 3.5. Prevalence of Chlorine Resistant Bacterial Pathogens

This experiment was outlined to evaluate the ability of isolated pathogens tolerance against most commonly used disinfectant worldwide i.e. Chlorine. It is worth to mention herein that all the isolated bacterial strains were able to tolerate 1 mg/L of Chlorine. All isolates showed the growth in 24 hrs incubation in the presence of Chlorine. These results highlight the ineffectiveness of chlorine to be used as disinfectant in DW (Figure 5).



**Figure 5.** Chlorine resistance (1 mg/L) pattern in water isolates.

These findings are in line with another study in which Chlorine resistance of different bacteria isolated from water distribution system in Duhok, province of Iraq was reported [39]. Another possible explanation for the present findings may be that DW borne bacteria definitely has developed some underline mechanism to survive in the mentioned concentration of chlorine. Extensive literature search has revealed that very less attention has given to study the possible effects of excessive chlorinated water intake on human health. Therefore, the scientific community may also focus on this issue. Furthermore, booster chlorination doses to attain the bacterial survival mechanism in chlorinated water may be the reason for the compromised purity of DW. Evidence suggested that adverse effects related persistent intake of high concentrations of Chloride is attributed to increase the number of polymorphonuclear leukocyte and disturbed blood cell counts in full blood cell count analysis. Chlorine is one of the oxidizing substances that reduce the level of oxygen in the cells, increasing the risk of heart disease. Furthermore, teeth weakness and decay, inflammation of mouth lining, respiratory system illnesses including bronchitis, digestive

system irritations including vomiting, chest pain, inflammation of pharynx, esophagus, and liver and intestinal tissues are the cardinal related risks of prolonged consumption of chlorinated water. In addition to that presence of positive correlation of developing colon, stomach, pancreas, liver, bladder and anal cancer and the exposure to chlorinated byproducts in drinking water is also a major point of concern [40].

## 4. Conclusion

Since sustained chlorinated DW consumption is attributed to the mild to severe complications of the cardinal systems of human body. Therefore, in conclusion it is recommended to regularly monitor Chlorine dosage to attain the DW disinfections. Furthermore, health risk to humans upon exposure to Chlorine tolerant bacterial contaminated DW may also be considered as development of resistance mechanism in bacteria. This may also contribute in alteration of antibiotic sensitivity profile of these bacteria. Thus serving as an additional factor to increase the pathogenicity of bacteria. In contrast, collectively these factors may narrow the treatment options for patients developed infection by these chlorine and antibiotic resistant DW bacteria. Lastly, it is strongly recommended to boil the DW, as it effectively kills all the bacteria in addition to chlorination.

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