Comparative Study of the Antibiotic Resistance Exhibited by Fecal Coliforms Obtained from Dighi Talab and Bisar Talab of Gaya, Bihar (India)

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Abstract: The antibiotic resistance pattern of four types of fecal coliform bacteria obtained from two fresh water ponds of Gaya was examined during present research work. Data obtained showed that cent per cent of the total isolates (83) of E. coli obtained from water sample of Dighi Talab showed resistance against Ampicillin as well as 83.13%, 71.08%, 69.88% and 32.53% remained resistant against Erythromycin, Gentamycin, Tetracycline and Chloramphenicol respectively. Cent per cent of the total isolates (27) of K. pneumoniae obtained from the water sample of this pond remained resistant against Tetracycline as well as 96.30%, 92.59%, 25.93% and 18.52% of total isolates showed resistance against Erythromycin, Ampicillin, Chloramphenicol and Gentamycin respectively. Out of the total (26) isolates of P. aeruginosa obtained from water sample of this pond 96.15%, 88.46%, 80.77%, 58.55% and 34.62% exhibited resistance against Tetracycline, Amplicillin, Erythromycin, Gentamycin and Chloramphenicol respectively. Cent per cent of the total isolates (06) of Sammonella sp. obtained from the water of this pond remained resistant against Teracycline as well as 83.33% remained resistant against Amplicillin, Gentamycin and Chloramphenicol. Only 66.67% of Salmonella sp. obtained from this pond remained resistant against Enthromycin. Out of the total (40) isolates of E. coli obtained from water sample of Bisar Talab, 97.50%, 95%, 87.50%, 75% and 60% showed resistance against Amplicillin, Tetracycline, Erythromycin, Gentamycin and Chloramphenicol respectively. Whereas, out of the total (36) isolates of K. pneumoniae obtained from this pond, cent per cent exhibited resistance against Amplicillin and Gentamycin. But 91.67%, 83.33% and 19.44% of the total isolates of this bacteria obtained from this pond and tested during present research work exhibited resistance against Tetracycline, Erythromycin and Chloramphenicol respectively. Out of the total (27) isolates of P. aeruginosa obtained from water of this pond, 24,22,17 and 13 exhibited resistance against Amplicillin & Tetracycline, Erythromycin, Chloramphenicol and Gentamycin respectively. Only two isolates of Salmonella sp. were obtained from the water sample of this pond, out of which one remained resistant against Gentamycin and Chloramphenicol whereas both remained resistant against. Amplicillin, Erythrimycin and Tetracyline. The main cause of antibiotic resistance exhibited by different bacterial isolates should be examined by future research.

Keyword: Antibiotic Resistance, Fecal coliforms, Dighi Talab, Bisar Talab

1. Introduction

Antibiotics discharges in water body remain resistant to biodegradation. More investigations in different parts of world are conducted with regard to antibiotic resistance among fecal coliforms present in different aquatic ecosystems. Antibiotic resistance gene transfer mainly takes place through E. coli of fecal origin because this bacteria remain abundant in aquatic ecosystems. Grabow and Prozesky (1973) pointed out that 30% of all coliform bacteria of fecal origin have transferable antibiotic resistance through cell to cell contact, through R-feactor or R-plasmid, and extachromosomal genetic material. Mudryk and Skorcewski (2009) analysed the frequency of antibiotic resistance in bacteria inhabiting water of a pond and obtained significant result. Toroglu et al. (2005) examined the antibiotic resistance pattern in Gram negative bacteria obtained from a river of Turkey. Salmonella bacteria is also called as target bacteria which acquire resistance genes horizontally and develops multi-drug resistance. Selective pressure on pathogenic fecal bacteria increased due to increasing use of different antibiotics by humans. They pointed out that E. coli, Salmonella, K. pneumoniae and P. aeruginosa are used as indicators of water quality. In above mentioned perspective the antibiotic resistance pattern of four types of fecal coliforms obtained from two ponds of Gaya were evaluated.

2. Materials and Method

Water samples from both ponds under study were collected in the month of July, 2022 in sterilized bottles, properly capped and transported to the laboratory. Bacterial isolates were cultured on nutrient agar media. Identification of bacterial isolates was conducted on the basis of characteristics morphological and biochemical characterization with the help of standard literature. Antibiotic resistance properties of each bacterial isolated obtained from both ponds were examined separately by Kirby-Bauer Disc Diffusion method (1966). Amplcillin and Gentamycin with disc concentration of 10µg, Chloramphenicol and Tetracydine with 10disc concentration of 30 µg as well as Erythromycin with disc concentration of 15µg were used during present experimentation. Bacteria exhibiting zone of inhibition as $\leq 11(mm)$ with Ampicillin, \leq 12 (mm) with Gentamycin and Chloramphenicol < 13 (mm) with Erythromycin and ≤ 14 (mm) with Tetracycline were considered as resistant.

3. Result and Discussion

It became evident from the data presented in Table-01 that out of 83 isolates of *E. coli* obtained from Dighi Talab, all showed resistance against Ampicillin and 69, 59, 58 and 27 isolates showed resistance against Erythromycin, Gentamycin, Tetracycline and Chloramphenical

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respectively. Out of the 27 isolates of *K. preumoniae* obtained from water sample of this pond, all showed resistance against Tetracycline as well as 26, 25, 07 and 05 isolates exhibited resistance against Erythromycin, Ampicillin, Chloramphenicol and Gentamycin respectively

Total 06 isolates of *Salmonella* sp. were isolated from this pond, among them, all 06 exhibited resistance against Tetracycline; 05 against Amplicillin, Gentamycin and Chloramphenicol as well as 04 against Erythromycin.

Table 1: Antibiotic Resistance Pattern of Bacterial Isolates obtained from Dighi Talab

	S. No.	Bacterial Isolates	Antibiotic Resistance										
			Ampicillin		Erythromycin		Gentamycin		Chloramphenicol		Tetracycline		Total Isolates
			No.	%	No.	%	No.	%	No.	%	No.	%	
	1	E.coli	83	100	69	83.13	59	71.08	27	32.53	58	69.88	83
	2	K. pneumoniae	25	92.59	26	96.30	05	18.52	07	25.93	27	100	27
	3	P. aeruginosa	23	88.46	21	80.77	14	53.85	09	34.62	25	96.15	26
	4	Salmonella sp.	05	83.33	04	66.67	05	83.33	05	83.33	06	100	06

Table 2: Antibiotic Resistance Pattern of Bacterial Isolates obtained from Bisar Talab

	Bacterial Isolates	Antibiotic Resistance										
S. No.		Ampicillin		Erythromycin		Gentamycin		Chloramphenicol		Tetracycline		Total Isolates
		No.	%	No.	%	No.	%	No.	%	No.	%	
1	E.coli	39	97.50	35	87.50	30	75.00	24	60.00	38	95.00	40
2	K. pneumoniae	36	100	30	83.33	36	100	07	19.44	33	91.67	36
3	P. aeruginosa	24	88.89	22	81.48	13	48.15	17	62.96	24	88.89	27
4	Salmonella sp.	02	100	02	100	01	50.00	01	50.00	02	100	02

Data mentioned in Table-02 indicates that out of the 40 isolates of *E. coli* obtained from water sample of Bisar Talab, 39, 38, 35, 30 and 24 exhibited resistance against Ampicillin, Tetracycline, Erythromycin, Gentamycin and Chloramphenicol respectively. Altogether 36 isolates of *K. pneumonia* were obtained from the water sample of this

sample of this pond, out of which 24, 22, 17 and 13 remained resistant against Ampicillin & Tetracycline, Erythnomycin, Chlormaphenicol and Gentamycin respectively. Out of the 02 isolates of *Salmonella* sp. obtained from the water sample of this pond, one remained resistant against Gentamycine and Chlororamphenicol. Both isolates of *Salmonella* sp. obtained from water of this pond showed resistance against Ampicillin, Erythromycin and Tetracycline.

Antibiotic resistance pattern of different fecal bacteria and presence of resistance gene were observed in waste water discharged in water bodies by Andersen (1993), Iwane *et al.* (2001), Schwartz *et al.* (2003) and Luczkiewicz *et al.* (2010). They pointed out that antibiotic resistance genes can spread by fecal contamination leaking from septic tanks and sewage as well as sanitary pipes. Micro-biological quality of the water of any pond requires periodic monitoring because presence of fecal coliforms often remain associated with presence of other pathogenic bacteria (Abdul Salam *et al.*, 2012). Karaca *et al.* (2023) observed 40% antibiotic susceptibility of *E. coli* obtained from Orontes river (Turkey) against Ampicillin. They conducted their study by Kirby Bauer Method.

Americo-Pinheiro *et al.* (2021) pointed out that indiscriminate use of antibiotics in human medicine remain responsible for the emergence of resistant bacterial strains against different antibiotics in the water bodies. Ozgumus *et al.* (2009) also isolated Ampicillin resistant *E.coli* while examining the water samples obtained from ten rivers of Turkey. The results obtained during present research work remain in conformity with results obtained by Abdulsalam pond, out of which 33,30 and 07 remained resistant against Tetracycline, Erythromycin and Chloramphenicol respectively. But all strains of *K. pneumonia* remained resistance against Amplicillin and Gentamycin. Altogether 27 isolates of *P. aeruginosa* were obtained from the water

et al. (2012), Karaca et al. (2023) and Ozgumes et al. (2009).

4. Conclusion

Out of the total isolates (83) of E. coli obtained from the water sample of Dighi Talab, 32.53% showed multi-drug resistance against all tested antibiotics. 18.52% of K. pneumniae, 34.62% of P. aeruginosa and 66.67% of Salmonella sp. obtained from the water sample of this pond showed multi-drug resistance against all tested antibiotics. Out of the total isolates (40) of E.coli obtained from water sample of Bisar Talab, 60% showed multi-drug resistance against all five tested antibiotic drugs. As well as, 19.44% of total K. pneumonia, 48.15% of P. aeruginosa obtained from this pond showed multidrug resistance and one out of two isolates of Salmnella sp. showed multi-drug resistance against all tested antibiotics. Thus it became evident that water of Dighi Talab contains high number of antibiotic resistant bacteria as compared to Bisar Talab. This may be due to high level of anthropogenic activities with Dighi Talab and low level of anthropogenic activities with Bisar Talab of Gaya.

References

- [1] Abdulsalam I., Rafida, Altabet A. and Guuma A.R., 2012, The presence of total coliform and fecal coliform bacteria in rainwater harvesting in Jefara district of Libya, *Int. Cont. Ecol. Environ. Biol. Sci.*,7-8: 340-342.
- [2] Americo-Pinheiro J.P.H. *et al.*, 2021, Monitoring microbial contamination of resistant *Escherichia coli* isolated from the surface water of Urban park in South

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Eastern Brazil, *Environmental Nanotechnology Monitoring and Management*, 15: 100438.

- [3] Anderson S.R, 1993, Effects of waste water treatment on the species composition and antibiotic resistance of coliform bacteria, *Current Microbiology*, 26:97.
- [4] Bauer A.W., Kirby W.M., Sherris J.C. *et al.* 1966, Antibiotic susceptibility testing by a standardized single disc method, *American Journal of Clinical Pathology*, 45(4): 493-496.
- [5] Grabow W.O. and Prozesky O.W., 1973, Drug resistance of coliform bacteria in hospital and city sewage, *Antimicrobial Agents and Chemotherapie*, 3(2): 175-180.
- [6] Iwane T., Urase T. and Yamamoto K., 2001, Possible impact of treated wastewater discharge on incidence of antibiotic resistant bacteria in river water, *Wat. Sci. Technol.*, 43(2):91.
- [7] Karaca C., Hiiner T. and Taker H.A.M., 2023, Antibiotic susceptibility profiles of *Escherichia coli* strains and fecal contamination in Orontes River, Turkey, *International Journal of Life Sciences* and Biotechnology, 6(2): 155-165.
- [8] Luczkiewicz A., Fudala K.S., Jankows K., Quant B. and Olanczuk N.K., 2010, Diversity of fecal coliforms and their antimicrobial resistance patterns in wastewater treatment model plant, *Water Sci. Technol.*, 61(6): 1383.
- [9] Ozgumus B.B. *et al.*, 2009, Class 1 and Class 2 integrons and plasmid mediated antibiotic resistance in coliforms isolated from ten rivers in Northern Turkey, *Journal of Microbiology*, 47(1): 19-27.
- [10] Schwartz T., Kohnew W., Jansen B. and Obst U., 2003, Detection of antibiotic- resistant bacteria and their resistance genes in wastewater, surface water and drinking water biofilms, *FEMS Microbiology Ecology*, 43(3): 325.
- [11] Toroglu S., Dincer S. and Korkmaz H., 2005, Antibiotic resistance in Gram negative bacteria isolated from Aksu river in Turkey, *Annals of Microbiology*, 55(3): 229-233.