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Antimicrobial Resistance of Enteric Pathogens Isolated from Acute Gastroenteritis Patients in Gaza strip, Palestine

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Background: Acute gastroenteritis is a common infection among the children of Gaza. The emergence of antibiotic resistance in enteropathogenic bacteria has profound clinical implications on the acute gastroenteritis disease. This study is a matched case-control and aims to determine the incidence of enteropathogenic bacteria, antibiotic resistance and associated-risk factors in diarrheal patients in Gaza Strip.

Methods: A total of 132 patients with acute diarrhea were investigated along with data collected from 132 healthy controls having the same demographic characteristics of patients except they didn't suffer from diarrhea within the last three months. The same data were collected through completing a questionnaire form for the cases and healthy controls. Stool samples were collected from six Primary Health Care Clinics and the samples were inoculated, isolated and identified using standard bacteriological methods in the Remal clinic- Microbiology laboratory.

Results: A total of 12 (9.1%) enteropathogenic bacteria spp. were isolated from 132 stool samples. *Salmonella, Campylobacter coli/jejuni,* and *Aeromonas hy-drophilia* were isolated in equal numbers from samples (3/12; 25% each), *Shigella* 2/12 (16.7%), and *Yersinia enterocolytica* 1/12 (8.3%). The two *Shigella* isolates were *Shigella boydii*. The antimicrobial profile of all isolated enteropathogenic bacteria showed high resistance rates against the tested antimicrobials. *Campylobacter coli/jejuni* (61.1%), followed by *Y. enterocolytica* (57.1.7%), *A. hydrophilia* (54.7.1%), *Shigella* (28.5%) and *Salmonella* spp. (9.5%). The highest antimicrobial resistance rates were found against erythromycin (75%) and Amoxicillin (65%).

Conclusions: Most enteropathogenic bacteria isolates showed high resistance rate to several antimicrobials. This study demonstrated that *C. coli/jejuni*, *A. hy-drophilia*, and *Y. Enterocolytica* are detected as causative agents of diarrhea in Gaza Strip.

Keywords: Gastroenteritis, Diarrhea, Enteropathogenic bacteria, Antibiotic resistance, Gaza strip.



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Introduction

Acute gastroenteritis is a common infection among children in Gaza, and it is know to cause high morbidity and mortality among them if they have not received proper treatment on time (1). The antibiotic resistance of enteric bacteria has profound clinical features on children because it cause sever dehydration and threats their life (2).

It is well established that antibiotic resistance increased with uncontrolled use of antibiotics in any community, especially when antibiotics are used by population without prescription, and this trend has become a major concern in Gaza (3). Moreover, there is few data about the spread of antibiotic resistance in enteropathogenic bacteria causing diarrhea in children of Gaza. The results of this investigative work would provide useful data for Gaza health authorities in order to control the incidence of infectious diarrhea among children.

Material and methods

A total of 132 stool samples were collected from patients with acute diarrhea during the period from Jan 2010 to Feb 2010. These patients were attended the Primary Health Care Clinics (PHCC) at Ministry of Health (MOH) in Gaza **Strip be**cause they have acute diarrhea (defined as the passage of 3 or more loose or liquid stools per day). The age of patients was between one month to 60 years old. In addition, personal data were collected from 132 healthy controls visited the same PHCC at the same period of study. Healthy controls had the same demographic characteristics of patients such as age, gender, except the fact they don't suffer from diarrhea within the last three months.

A questionnaire was administered and data was collected through personal interview of subjects or their guardians. The questionnaire included personal, socioeconomic and past history of diseases. An ethical approval from health authority in Gaza was obtained and informed consent from each participant was also obtained.

Isolation procedures of enteropathogenic bacteria

Each stool sample was directly inoculated onto Xylose Lysine Deoxycolate agar (XLD) agar, *Salmonella Shigella* (SS) agar, Hektoen enteric (HE) agar, *Campylobacter* Blood-Free Selective Agar Base (Modified CCDA) and *Yersinia* selective agar. Approximately 1 g of each sample inoculated into 10 ml of Selenite F broth, alkaline peptone water, and phosphate buffer saline (PBS). The cold enrichment method was used to isolate *Yersinia enterocolitica* (4-6).

Identification procedures of enteropathogenic bacteria

Suspected colonies of enteropathogenic bacteria were identified by colony morphology, Gram stain and biochemical tests including oxidase, catalase, API20E and API campy systems. Confirmation of *Salmonella* and identification of *Shigella* strains carried out by polyvalent antisera (7, 8).

Antimicrobial susceptibility for the bacterial isolates

Antimicrobial susceptibility for the isolated enteropathogenic bacteria was done by disk diffusion method on Muller-Hinton agar. The procedures and zone of inhibition interpretation was done according to Clinical Laboratory Standards Institute (CLSI, 2011) (9). For *Campylobacter*, all strains were suspended in phosphate buffer saline (PBS) pH 7.4 and inoculated on Muller-Hinton agar supplemented with 5% of whole blood according the method described by Miranda et al. (10).

Results

This study included (56%) female and (44%) males. A total of 75% of diarrhea cases and controls were aged less than 5 years. It was found that (15.2%) of cases live in houses with < 5 household members, while (19.7%) of controls live in houses with < 5 household members (P = 0.209). An equal number of enteropathogenic bacteria were isolated from both male and female. From male: *Salmonella* and *A. hydrophilia* account for (1.5%), whereas *Shigella* and *Campylobacter coli/jejuni* account for (0.76%). However, for female the percentage was (0.76%) for *Salmonella*, *Shigella*, *A. hydrophilia* and *Y. enterocolytica*, and (1.5%) for *C. coli/jejuni*, (P = 0.736) (**figure 1**).

The antimicrobial profile of all isolated enteropathogenic bacteria showed high resistance rates for *Campylobacter colil jejuni* (61.1%), followed by *Y. enterocolytica* (57.1.7%), *A. hydrophilia* (54.7.1%), *Shigella* (28.5%) and *Salmonella* spp. (9.5%) as shown in **Table 1**.

Table 2 shows the relation between diarrhea cases and presence of poultry in houses, and **table 3** show the distribution of diarrhea cases in association of available drinking water in houses.

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|-------------------------------------|-------------|----|------|----|------|------|------|-------------------------------------|------|
| Enteropathogenic bacteria (n=12) | Antibiotics | | | | | | | Total Percentage of resistance % | |
| | AM | E | CU | GM | TE | SXT | NA | CIP | |
| Salmonella spp. | 0 | - | 0 | 0 | 33.3 | 0 | 33.3 | 0 | 9.5 |
| Shigella spp. | 50 | - | 0 | 0 | 50 | 50 | 50 | 0 | 28.5 |
| Campylobacter coli/jejuni | 75 | 75 | - | - | - | - | - | 33.3 | 61.1 |
| Aeromonas hydrophilia | 100 | - | 75 | 0 | 75 | 100 | 33.3 | 0 | 54.7 |
| Yersinia enterocolytica | 100 | - | 0 | 0 | 100 | 100 | 100 | 0 | 57.1 |
| Total Percentage of resistance | 65 | 75 | 18.7 | 0 | 64.5 | 62.5 | 54.1 | 6.6 | 40.1 |

Table 1. Antimicrobial susceptibility of isolated enteropathogenic bacteria.

Amoxicillin (AM), Erythromycin (E), Cefuroxime (CU), Gentamicin (GM), Tetracycline (TE), Trimethoprim/sulfamethoxazole (SXT), Nalidixic Acid (NA), Ciprofloxacin (CIP),.

Table 2. Domestic animals in the house associated with diarrhea cases.

| P value | Control | (n=132) | Case (| n=132) | (n=264) | | |
|---------|---------|---------|--------|--------|---------|---------|--|
| | % | N0. | % | N0. | | | |
| 0.001* | 11.4 | 15 | 26.5 | 35 | Yes | Poultry | |
| | 88.6 | 117 | 73.5 | 97 | No | | |
| 0.111 | 9.1 | 12 | 4.5 | 6 | Yes | Rabbit | |
| | 90.9 | 120 | 95.5 | 126 | No | | |
| 0.086 | 5.3 | 7 | 10.6 | 14 | Yes | Sheep | |
| | 94.7 | 125 | 89.4 | 118 | No | | |
| 0.006* | 8.3 | 11 | 19.7 | 26 | Yes | Pigeon | |
| | 91.7 | 121 | 80.3 | 106 | No | | |
| 0.500 | 9.1 | 12 | 8.3 | 11 | Yes | Donkey | |
| | 90.9 | 120 | 91.7 | 121 | No | | |

*Significance P < 0.05

Table 3. Presence of drinking water in the houses of investigated cases

| P value | Control(n=132) | | Case(I | n= 132) | | |
|---------|----------------|-----|--------|----------------|--------------|----------------------|
| | % | N0. | % | N0. | Total(n=264) | |
| 0.421 | 9.8 | 13 | 11.4 | 15 | Yes | Tap water |
| | 90.2 | 119 | 88.6 | 117 | No | |
| 0.500 | 89.4 | 118 | 88.6 | 117 | Yes | Filtered water |
| | 10.6 | 14 | 11.4 | 15 | No | |
| 0.142 | 6.8 | 9 | 11.4 | 15 | Yes | Boiled water |
| | 93.2 | 123 | 88.6 | 117 | No | |
| 0.286 | 13.6 | 18 | 10.6 | 14 | Yes | Bottle water |
| | 86.4 | 114 | 89.4 | 118 | No | |
| 0.042* | 2.3 | 3 | 7.6 | 10 | Yes | Domestic wells |
| | 97.7 | 129 | 92.4 | 122 | No | |
| 0.247 | 25.8 | 34 | 30.3 | 40 | Yes | Water access changes |
| | 74.2 | 98 | 69.7 | 92 | No | |

Significance P < 0.05 *

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Figure 1. Enteropathogenic bacteria isolated from the stool samples.

Note: An equal number of enteropathogenic bacteria were isolated from both male and female. From male: Salmonella and A. hydrophilia account for (1.5%), whereas Shigella and Campylobacter coli/jejuni account for (0.76%). However, for female the percentage was (0.76%) for Salmonella, Shigella, A. hydrophilia and Y. enterocolytica, and (1.5%) for C. coli/jejuni

Discussion

This study shows that children aged less than 5 years old are significantly more susceptible (75%) to develop infectious diarrhea than older children, and enteropathogenic bacteria were isolated with a higher frequency (66.7%) from this age group. However, this study can't exclude that many other diarrheal cases might have been caused by enteric viruses or pathogenic intestinal parasites. According to recent reports of WHO, diarrheal diseases caused by viruses, parasites and enteropathogenic bacteria are a leading cause of morbidity and mortality in children; with 1.5–2.5 million deaths estimated to occur annually among children aged < 5 years and mostly in developing countries (11).

The results of this study is limited only to bacterial causative agents of diarrhe and these are nearly similar to previous study conducted in Gaza by Abu Elamreen *et al.,* (2007) where they have reported that (10%) of their stool samples had enteropathogenic bacteria using conventional culture method (12). Other studies from our Arab countries reported higher percentage of enteropathogenic bacteria (17%) in Egypt (13) and (40%) in Jordan (14). In addition, our findings showed that rate of detection of enteropathogenic bacteria is higher than in a local study in El-Naser Hospital carried during the period 1999 to 2006, which has showed that the isolation frequency of *Salmonella* and *Shigella* was 1.8% and 0.8%), respectively (15).

This study suspected that contamination of groundwater in Gaza Strip with sewage water may be an important source of diarrhea caused by *A. hydrophila*, especially in Khan Yunis area, where *A. hydrophila* is frequently present in its sewage (16, 17). Nearly 10% of the population of the Gaza Strip (over 100,000 people) had no proper tap water supply in February 2009, following the last Israeli military attacks on Gaza (18, 19).

Multidrug resistance was commonly observed among enteropathogenic bacteria isolates, especially in *Campylobacter coli/jejuni* (75% resistant to erythromycin and ampicillin). Studies from neighbouring Arab countries (Jordan, Egypt, Lebanon) showed also wide spread of multidrug resistance among enteric bacteria isolated from human and water sources (14, 20, 21, 22). There was statistically significant correlation between the presence of poultry in the houses and occurrence of diarrhea cases (P < 0.05). This is probably because those who live in rural areas have more close contact with livestock and have a greater risk of *Campylobacter* infection (23).

In Conclusion, this study demonstrates a relatively low percentage of enteropathogenic bacteria detected among diarrheal patients, and most of these bacteria isolates were multidrug resistance.

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