Epidemiology and Antibiotic Susceptibility of Vibrio cholerae Associated with the 2017 Outbreak in Kasese district

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Abstract

Background: Cholera has continued to be a global threat to public health and a key indicator of lack of social development in developing countries. This report summarizes findings from the epidemiologic investigation, which aimed at identifying the mode of transmission and antibiotic susceptibility patterns of the Vibrio cholerae isolated in Kasese district, Uganda. Materials and methods This was a descriptive cross-sectional study to describe the epidemiology of the 2017 cholera outbreak in Kasese district. Rectal swabs were collected from 69 suspected case-persons and cultured on Thiosulphate-Citrate-Bile-Salts Sucrose (TCBS™; SEIKEN Japan) agar and incubated at 37°C for 18-24 hours. The isolates were serotyped with polyvalent 01 antiserum and monovalent serotype Inaba and Ogawa antisera (Denka Seiken, Tokyo, Japan) to determine which serotype was responsible for the outbreak. Antimicrobial susceptibility testing was performed using the Kirby-Bauer disk diffusion method on Mueller-Hinton agar. Results A total of 222 cases were recorded in the Kasese District outbreak between the month of September 2017 and January 2018 with the case fatality rate (CFR) of 1.4%. Children below the age of 14 years contributed the biggest proportion of the cases (70%) and out of these, 33% were aged below five years. Antibiotic susceptibility using Kirby-Bauer disc diffusion method showed 100% resistance to Ampicillin and over 50% were resistant to trimethoprim/sulfamethoxazole whereas gentamicin showed 100% susceptibility. The general trend also showed increased susceptibility to combination therapy as opposed to mono-therapy.
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Conclusion

Laboratory capacity to detect and monitor the rapidly emerging drug resistance among \textit{V. cholerae} isolates needs to be improved in order to effectively handle the dual challenge of treatment and prevention of Cholera in Kasese District.

Background

Cholera has continued to be a global threat to public health and a key indicator of lack of social development (1, 2). The World Health Organization (WHO) estimates that 3–5 million annual cases of cholera occur worldwide, resulting in 100,000–120,000 deaths (3). Since its introduction to Africa during the seventh cholera pandemic in 1970, \textit{Vibrio cholerae} caused regular vast epidemics across the continent with cumulative 3,762,902 case-notifications (4). In 2013 alone, 22 African countries reported 56,329 cholera cases. However, these numbers are substantially underestimated due to poorly functioning national epidemiological and laboratory surveillance systems, which are not able to detect the majority of mild disease presentations (4). Emergencies often have a direct impact on the health systems and public health systems of an affected region or country, particularly in resource-constrained areas (5, 6). Factors that may lead to outbreaks in endemic areas include wars, civil unrests, flooding, earthquakes, landslide, poor sanitation and improper refuse dumping, open defecation and slumps (7). The mode of transmission of Cholera is through fecal contamination of food or water. Transmission is therefore closely associated to poor environmental management due to water and sanitation issues (1, 2).

In Uganda, which is one of the 51 endemic countries, cholera outbreaks occur mainly along the western border with Democratic Republic of Congo (DRC), in the Karamoja region to the north, Mbale in the Eastern Region and in the slums of Kampala City (8). Kasese district is affected by regular cholera epidemics. The different Serotypes and biotypes have been reported in the previous studies in Uganda (9). However, antibiotic susceptibilities have not been determined. Kasese District has a hilly terrain; hence many areas are supplied by gravitational water flow schemes (GWFS). Poor hygiene and sanitation conditions and regular flooding of the low lands usually create an ideal environment for disease to spread. Consequently, the district is prone to waterborne diseases, including cholera (10). Cholera was eliminated in the developed countries over a century ago largely by simple treatment of domestic and waste water but continues to maintain a home in resource constrained settings especially in Sub-Saharan Africa. One would rightly say that cholera is the disease for the poor. On 2\textsuperscript{nd} October 2017, the Ministry of Health in Uganda issued an alert of a cholera outbreak in Kasese District. This report summarizes findings from the epidemiologic investigation, which aimed at identifying the mode of transmission and antibiotic susceptibility patterns of the \textit{Vibrio cholerae} isolated in Kasese district.
Methods

Study design and study area

This was a descriptive cross-sectional study carried out between September 2017 and February 2018 to describe the epidemiology of the 2017 cholera outbreak in Kasese district, Uganda (Fig. 1). Kasese District is located along the equator and it is bordered by Kabarole District to the north, Kamwenge District to the east, Rubirizi District to the south, and the Democratic Republic of the Congo to the west.

Study population and sampling

The study population comprised of both adults and children suspected to be suffering from cholera admitted in the isolation wards in Kagando hospital and Bwera hospital. In the community, the patients who were discharged from the isolation wards or their parents/guardians formed the study population. Consecutive sampling was carried out throughout the outbreak season until the time when the epidemic was controlled. Sample size calculation was not applicable.

Data collection

A list of discharged patients was obtained from the isolation units of Bwera hospital and Kagando hospital and the individuals were followed to the community where they live. A total of 222 patients were recorded in the discharge book some of whom were children and these were followed to the community. Questionnaires were administered to a total of 75 participants who were either the cases or a relatives to the case. For the cases that were clustered in one place, only one case or relative of the case was selected randomly and considered for interviews. Key questions on symptomatology and environment were asked. We also took GPS points of the homes of the cases and pictures of potential source infection (Fig. 1)

Laboratory detection of *Vibrio cholerae*

Self-collected rectal swabs from suspected case-persons were transported to the laboratory in alkaline peptone water medium for culture and sensitivity. The samples were cultured on Thiosulphate-Citrate-Bile-Salts Sucrose (TCBS™; SEIKEN Japan) agar and incubated at 37°C for 18-24 hours. The isolates were serotyped with polyvalent O1 antiserum and monovalent serotype inaba and Ogawa antisera (Denka Seiken, Tokyo, Japan). Rectal swabs were collected from 69 participants and antibiotic susceptibility done in Bwera hospital and Kagando Hospital laboratories.

Antibiotic susceptibility testing

Antimicrobial susceptibility testing was performed using the Kirby-Bauer disk diffusion method on Mueller-Hinton agar. The Escherichia coli reference strain ATCC 25922 was used as a control. Isolates
were tested against 7 antimicrobial drugs as follows: ampicillin (10 µg), ciprofloxacin (5 µg), chloramphenicol (30µg), gentamycin (30µg), nalidixic acid (30µg), sulfamethoxazole/trimethoprim (1.25 µg + 23.75 µg) and tetracycline (30µg) (all Oxoid, United Kingdom). Zones of inhibition were interpreted according to the 2014 Clinical and Laboratory Standards Institute (CLSI) guidelines as resistant and susceptible (11).

**Potential sources of bias**

Recall bias was the most common potential confounder as recollection of previous events might have been difficult for our respondents. However the authors tried to minimize this by administering the questionnaires during the outbreak season to benefit from fresh memories.

**Ethical considerations**

Ethical approval was obtained from the Makerere University School of Biomedical Sciences Higher Degrees Research and Ethics committee (SBS-292). Written informed consent was obtained from all study participants. Consent from parents/guardians of participants below 18 years was sought and ascent was obtained from all minors who took part in this study. Participation was voluntary.

**Data management and presentation**

Data was checked for completeness to avoid cases of missing data, entered in excel, exported to STATA v14 for analysis and presented in form of Tables and Figures.

**Results**

**Epidemiological description of the 2017 outbreak**

A total of 222 cases were recorded in the Kasese District outbreak between the month of September 2017 and January 2018 with the case fatality rate (CFR) of 1.4%. The patients who died were two females aged 17 and 30 years and one male aged 60 years. The first case was admitted on 24th September, peaking at 87 in the epidemiologic week 39 and the last case was admitted in the epidemiologic week 49 (Fig. 2) before a drastic reduction in the epidemiologic week 41 due to initiation of the interventions to control the disease by the district response team. Few cases continued to be reported in a lower frequency until the outbreak was fully contained by the 8th of January 2018. The mean age of the cases was 16 ± 19.3 years and ranged from 7 months to 75 years. Children below the age of 14 years contributed the biggest proportion of the cases (70%) and out of these, 33% were aged below five years. Katojo primary school had 22 cases among the pupils and their latrine was washed away by the floods into river. The average hospital stay for the admitted cases was 4 days.
The first case was admitted on 24 September 2017 with a peak number of 87 cases in the 39th calendar week (24–29 September) (Fig. 2).

Environmental conditions in the outbreak area

Environmental assessment of the residences of the cases indicated inadequate water supply, poor sanitary conditions and unsafe disposal of solid waste (Fig. 3). All the three case fatalities were reported to consume untreated surface water with no methods employed to make the water safe for human consumption.

The outbreak of the disease was caused by exposure to faecal contaminated water or food, following a series of floods that washed away latrines into water sources in sub-counties of Nyakiyumbu, Bwera and Isango. The most affected villages included; Bunyiswa, Rwehingo, Katholhu, Rusese and Kyanzi.

Other factors responsible for its spread include poor personal hygiene, using contaminated water, poor sanitation as occurs in open defecation, eating food or drinks prepared under unhygienic conditions and poor personal hygiene, especially not washing hands after visiting the toilet.

Water sources and safety

Drinking safe water was not a concern until the time of the outbreak with 49% (37/75) of the homesteads consuming untreated surface water. Only 16% (12/75) of the homesteads reported consuming municipal tap water while the rest, 35% (26/75) consumed borehole water. None of the participants reported boiling drinking water or use of chemi-sterilants before the outbreak period. During the mapping exercise, we did not find any piped (tap) water in the entire outbreak zone and there was only one borehole sited in the whole community. The main source of water was river Kiyansi which is untreated surface water, visibly turbid with on-going sand mining activities.

Other sanitary conditions in the homes of the cases

Whereas most households in the outbreak area had poor pit latrines, there were rampant cases of open defecation. Most of the pit latrines were unimproved (Fig. 4) and some were washed away by heavy floods into the river and this is what sparked of the outbreak. Hand washing facilities were not observed at the pit latrine area a good indication for poor hand hygiene. Most of the toilets and kitchens were built as temporary structures using mad, banana fiber or grass which compromises food hygiene.

Besides the poor management of the human excretor, there was generally poor management of solid waste in the entire community. River banks were the dumping sites for garbage. The cooking areas were not safe for preparation of food and when it rained, the whole area was covered by mud. Most houses of the cases were made of mud walls and most times, especially during flooding, rain water entered the houses.

Symptomatology and clinical picture
The greatest proportion of the cases (92%) were clinically diagnosed and managed without laboratory confirmation for Cholera. All the cases in this outbreak presented with diarrhea as the most common symptom (100), followed by vomiting (92%). Abdominal cramps was however present in only 39% of the cases (Table 1 in the Supplementary Files).

The most affected people were from Bukonzo West constituency, an area that neighbors the Democratic Republic of Congo. This *V. cholerae* outbreak was epidemiologically linked to consumption of untreated surface water after heavy flooding leading to a number of pit latrines washing away into the water.

Antimicrobial susceptibility

Antibiotic susceptibility using Kirby-Bauer disc diffusion method showed 100% resistance to Ampicillin and over 50% were resistant to Trimethoprim/Sulfamethoxazole. Drugs like tetracycline which is among the recommended drugs in the clinical guidelines had close to 50% resistance whereas gentamicin showed 100% susceptibility. In addition, drugs like chloramphenicol and ciprofloxacin showed low resistance rates (11.76% and 5.9%) respectively. The general trend also showed increased susceptibility to combination therapy as opposed to mono-therapy (Fig. 5).

**Discussion**

Kasese suffers frequent annual outbreaks of cholera and in each outbreak, death of a case is the usual trigger for the surveillance activities. A total of 222 cases were recorded in the 2017 with the case fatality rate (CFR) of 1.4%. Similarly, over 100 people were affected by cholera in the same district in the year 2015 (9). The frequent outbreaks in Kasese district and other parts of Uganda have continued to feed into the high density of cholera cases reported in Sub-Saharan Africa.

Environment and sanitation

Kasese District is a hilly area with vast low lands that suffer frequent flooding. The outbreak of 2017 proceeded heavy flooding that washed toilets into the river in the low land. Water from this river is consumed without treatment and therefore exposing people to cholera and other water borne diseases. Consumption of contaminated water has been repeatedly implicated in a similar cholera outbreaks in Kasese (9), other parts of the country like Karamoja in North Eastern Uganda (12) and elsewhere in the world (13-15). Most of the latrines in the affected area were unimproved and this could have encouraged rampant cases of open defecation. While the National Housing and Population census reports that only 10% of the rural population in Uganda lacks access to pit latrines, it also acknowledges that about 58% of the pit latrines are unimproved (16). The quality of latrine facilities needs to be addressed if Uganda is to meet the sanitation SDG targets. Katojo Primary school whose latrine was washed away by the floods also had no hand washing facilities with evidence of open defecation around the latrine which points to the fact that the pupil to stance ratio is higher than the recommended 40:1 (pupil: stance) (17) meaning that the pupils have to queue longer to access the latrine facilities at any given time. Similarly the Uganda
national rural school pupil to stance ration has been reported to have increased from 70:1 to 71:1 between 2014 and 2015 (17).

Access to adequate health care and coordination of response

The persistently high number of cases reported in the frequent cholera outbreaks in Kasese district with some deaths at least reported during each outbreak may reflect more general problems in access to effective health care (9, 12). Uganda enjoys an international applause in management of emergency outbreaks like Ebola and Marburg due to availability of a fully functional state-of-the-art Public Health Emergency Operating Center and an operational public health emergency response plan (18), but more needs to be done in the case of Cholera (19). Bwera hospital and Kagando hospital, like other hospitals in Uganda, have a small health work force (20) that each time needs to heavily rely on other in-country work force in the emergency situations. Access to effective health care also largely depends on the availability to patients often too ill to walk. Cases were located in areas with unpaved roads (Fig. 6) that could not be accessed by motorized transport means. During the mapping of cases, we went sliding and surviving several falls to access homes of cases. The problem of poorly or unpaved roads though seems to be similar across the resource constrained settings across Africa, Asia and Latin America (21).

Even with the poor road infrastructure, the local district team with the benefit of the previous experience of managing cholera outbreaks quickly and effective responded with awareness messages and active identification of cases. At the time of the outbreak, the laboratory lacked culture media and other necessary supplies necessary to detect and confirm the Cholera cases, at that point, anybody who presented with diarrhea was considered a case. A similar study in Kenya also pointed out similar challenges of supply chain and non-preparedness to respond to cholera outbreaks (22).

Antibiotic susceptibility

In this study, we report a high resistance to Ampicillin and trimethoprim/sulfamethoxazole and this compares with the findings of other studies (23). We note with great concern the high resistance to tetracycline of over 50% yet it is one of the main antibiotics recommended in the management of Cholera (Fig.5). Several other studies have reported the rise in resistance to tetracycline by V. cholerae (23, 24). The greatest susceptibility was noted in gentamicin and low resistance against ciprofloxacin was recorded. Similar studies have documented that there has been no notable change in the resistance to ciprofloxacin (23). Multi-drug resistance has also been noted in this study. With the emergence of multidrug resistance among V. cholerae isolates, Uganda now faces a double challenge of both control and management of cholera epidemics.

Conclusion

The Kasese Cholera outbreak 2017 was caused by exposure to faecally contaminated water or food following a series of floods that washed away latrines into water sources in sub-counties of Nyakiyumbu, Bwera and Isango. Laboratory capacity to detect and monitor the rapidly emerging drug resistance
among *V. cholerae* isolates needs to be improved to effectively handle the dual challenge of treatment and prevention of Cholera.

**Generalizability**

Our findings can be generalized to similar settings in Uganda and other developing countries.

**Abbreviations**

OHCEA: One Health Central and East Africa; CFR: case fatality rate; WHO: World Health Organization; CLSI: Clinical and Laboratory Standards Institute; GWFS: Gravitational Water Flow Schemes; SDG: Sustainable Development Goals; GPS: Global Positioning System; IDRC: International Development Research Centre – Canada. Gen = Gentamycin, SXT = Trimethoprim/Sulfamethoxazole, Tet = Tetracycline, Nal = Nalidixic acid, Cip = Ciproflaxacin, Chl = Chloramphenicol, Amp = Ampicillin

**Declarations**

**Acknowledgements**

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**Availability of data and materials**

All data on which the conclusions of this manuscript are drawn are contained in the main manuscript and the tables herein.

**Author Contributions**

IJS participated in conception of the research idea, study design and drafting the first manuscript; CK participated in analysis and interpretation of the data, IBR conducted critical reviews and through his
International Development Research Centre IDRC-Canada Ecohealth project partially funded the project and BBA conducted critical reviews and supervised the study. All authors have read and approved the final version of the manuscript.

**Ethics approval and consent to participate**

Ethical approval was obtained from the Makerere University School of Biomedical Sciences Higher Degrees Research and Ethics committee (SBS-292). Written informed consent was obtained from all study participants. Consent from parents/guardians of participants below 18 years was sought and ascent was obtained from all minors who took part in this study. Participation was voluntary.

**Consent for publication**

Not applicable

Competing interest

The authors declare that they have no conflict of interest

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Fig. 5: Antibiotic Resistance Patterns for *Vibrio cholerae* in Kasese District

Fig. 6: Accessibility to cases

**Table**

Due to technical limitations, table 1 is only available as a download in the supplemental files section.

**Figures**
Figure 1
Figure 2
Figure 5

![Graph showing percentage resistance for different drugs and drug combinations tested.]

Figure 6

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.
• supplement1.jpg