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# Title: Healthcare center wastewaters in Burkina Faso: sources of ESBL, AmpC-β-lactamase and carbapenemase producing Escherichia coli and Klebsiella pneumoniae

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# **Abstract**

#### **Background**

Extended-spectrum β-lactamase (ESBL), plasmid-mediated AmpC-β-lactamase and carbapenemase producing *Escherichia coli* and *Klebsiella pneumoniae* have spread into environment worldwide posing a possible public health threat. However, the prevalence data in low- and middle-income countries are still scarce. The aim of this study was to assess the occurrence of ESBL, AmpC-β-lactamase and carbapenemase producing and multidrugresistant *E. coli* and K. *pneumoniae* in wastewater collected from healthcare centers in Burkina Faso.

#### Methodology

Eighty-four (84) wastewater samples were collected from 5 healthcare centers and bacterial counts on ESBL ChromAgar were performed. *E. coli* and *K. pneumoniae* isolates were identified using API20E. ESBL production was confirmed using the double disc synergy test (DDST). AmpC-β-lactamase detection was performed on Muller Hinton (MH) agar supplemented with cloxacillin (4μg/l). Carbapenemase testing was carried out using O.K.N.V.I. RESIST-5 immunochromatography test.

#### Result

 $E.\ coli$  and/ or  $K.\ pneumoniae$  strains were isolated from 82 wastewater samples (97.6%). In total, 170 strains were isolated,  $E.\ coli$  more commonly (64%). Average concentrations of ESBL producing bacteria per hospital varied from  $1.10 \times 10^5$  to  $5.23 \times 10^6$  CFU/ml. Out of 170 presumptive ESBL producing isolates and 51 presumptive AmpC-β-lactamase producing isolates, 95% and 45% were confirmed, respectively. Carbapenemase production was detected in 10 isolates, 6 were NDM producers, 3 were OXA-48 producers and 1 was NDM and OXA-48 producer. All isolates were multidrug resistant and, furthermore, all of them were resistant to all β-lactams tested. Also, resistance to ESBL inhibitors was common, up to 66%  $E.\ coli$  and 62% in  $K.\ pneumoniae$ . Amikacin, fosfomycin and nitrofurantoin were the antibiotics for which least resistance was detected.

#### Conclusion

This study showed that wastewater from healthcare centers constitutes a reservoir of multidrug-resistant bacteria in Burkina Faso, including those capable of producing carbapenemases, which may disseminate into environment and further back to humans. Therefore, following the microbiological quality of the wastewaters released from healthcare centers is important to include in the future national AMR surveillance program.

# Introduction

The emergence and spread of antimicrobial resistance (AMR) represent a serious threat to human and animal health. In 2014, the number of deaths due to AMR was estimated at 700,000 (1, 2). If no action is taken against AMR, by 2050 this number could reach 10 million per year (1, 2). The economic cost of the AMR will vary from 1.1 to 3.8% of the global GPD (3). The annual shortfall by 2030 was estimated to reach \$3.4 trillion (3). Several causes, such as unreasonable use or overuse of antibiotics, have been speculated to favor the emergence and diffusion of resistance genes and multidrug resistant bacteria, (4–9). In low- and middle-income countries (LMICs), socio-economic and behavioral factors, such as poverty, use of poor quality antibiotics, absence of diagnostic tools, absence of antibiotic stewardship policies and uncontrolled use of antibiotics in animals, have been incriminated (5). The persistence of antibiotic residues, non-degraded antibiotics and disinfectants in the wastewaters contribute to selection of resistant bacteria and their wide spread in environment (4–6, 10–14). Multidrug resistant bacteria harboring extended spectrum β-lactamase genes (*bla*TEM, *bla*SHV, *bla*CTX-M) and carbapenemase genes (*bla*OXA-48, *bla*KPC, *bla*NDM, *bla*VIM and *bla*IPM) have been detected in hospital wastewaters from several countries (15–19). Management of healthcare center wastewaters in LMICs is highly insufficient and sometimes the wastewaters are directly discharged into the environment, drainage, rivers, or lakes without any treatment (20–22). Use of this water for various human activities exposes the population to new infections by multidrug resistant bacteria (23, 24)

In Burkina Faso, data on wastewater contamination are patcy, but recent studies have revealed the abundant presence of resistant bacteria in healthcare center effluents, (25, 26). The present study aimed to assess healthcare center wastewaters contamination specifically by ESBL-producing Gram negative bacilli and to perform phenotypic characterization of ESBL-producing *Escherichia coli* (ESBL-Ec) and *Klebsiella pneumoniae* (ESBL-Kp) in wastewaters from hospitals at different levels of the healthcare system in Burkina Faso.

# Results

# Bacterial concentration in healthcare center wastewaters

The average concentration of bacteria growing on ESBL selective plates from wastewater of each healthcare center varied from  $1.10x10^5$  to  $5.23x10^6$  CFU/ml. The highest bacterial counts were obtained from wastewater drained from Yalgado Ouédraogo teaching hospital (tertiary level hospital), followed by Koudougou regional hospital center and El Fateh SUKA Clinic (both secondary level healthcare facilities) (Table 1).

Table 1
Prevalence of bacteria from wastewaters of five healthcare centers in Burkina Faso growing on ChromAgar™ ESBL plates.

Healthcare centers	No. of samples	Average concentration (CFU/ml)
	(n = 84)	
Yalgado Ouedraogo teaching hospital	28	5.23x10 <sup>6</sup>
Koudougou Regional hospital Center	26	3.37x10 <sup>6</sup>
El Fateh Suka Clinic	14	3.00x10 <sup>6</sup>
Source de Vie medical Center	6	1.10x10 <sup>5</sup>
Saint Camille medical Center in Nanoro	10	1.85x10 <sup>5</sup>

# **Prevalence Of Esbl**

From the 84 healthcare center wastewater samples, suspected ESBL *E. coli* or *K. pneumoniae* isolates were detected in 82 samples (97.62%). In total, 170 strains were isolated (109 *E. coli* and 61 *K. pneumoniae*). ESBL test confirmed 160 (95%) isolates (102 *E. coli* and 58 *K. pneumoniae*) to be ESBL positive. Ten isolates were negative but they were resistant to all the β-lactam + ESBL inhibitors tested.

# Prevalence Of Ampc β-lactamase Producers

Isolates which were resistant or intermediately susceptible to cefoxitin (37 ESBL-Ec and 14 ESBL-Kp) were tested to detect AmpC- $\beta$ -lactamase production by the phenotypic method. 23 of 51 isolates (45%) were AmpC- $\beta$ -lactamase producers (Table 2).

Table 2
Prevalence of AmpC-8-lactamase producers among the cefoxitin resistant or intermediately susceptible isolates from the five healthcare centers

Healthcare centers	E. coli		K. pneumonia		AmpC-β-lactamase producers (%) *	
	tested (n)	AmpC positive (n)	tested (n)	AmpC positive (n)		
Yalgado Ouedraogo teaching hospital	15	3	4	2	9.80	
Koudougou Regional Hospital Center	12	7	5	2	17.65	
El Fateh SUKA Clinic	5	3	1	0	5.88	
Source de Vie medical center	1	1	0	0	1.96	
Saint Camille medical center (Nanoro)	4	1	4	4	9.8	
TOTAL	37	15	14	8	45.09	
* AmpC-β-lactamase producing <i>E. coli</i> and <i>K. pneumonia</i> out of the 51 isolates tested.						

# **Prevalence Of Carbapenemase Producers**

Twenty-one isolates resistant or intermediately susceptible to meropenem (15 ESBL-*Ec* and 6 ESBL-*Kp*) were tested to detect carbapenemase production (OXA-48, KPC, NDM, VIM, and IMP). Ten isolates (47.62%) were carbapenemase producers: 6 were NDM producers, 3 were OXA-48 producers, and 1 was NDM and OXA-48 producer. Carbapenemase producing bacteria were detected among wastewater collected from the tertiary and the secondary level healthcare facilities (Table 3).

Table 3

Prevalence of carbapenemase producers among the meropenem resistant isolates from the five healthcare centers

Healthcare centers	E. coli		K. pneumoniae		
	tested	carbapenemase positive (n)	tested (n)	carbapenemase positive (n)	
	(n)				
Yalgado Ouédraogo teaching Hospital	8	1 OXA-48	4	2 OXA-48	
		1 OXA-48 + NDM		1 NDM	
Koudougou Regional hospital Center	4	1 NDM	1	1 NDM	
El Fateh SUKA Clinic	3	2 OXA-48	0	0	
		1 NDM			
Source de Vie medical center	0	0	0	0	
Saint Camille medical center (Nanoro)	0	0	1	0	

# **Resistance To Antibiotics**

All the isolates from the ESBL selective plates (109 *E. coli* and 61 *K. pneumoniae* isolates) were tested against 31 antibiotics representing different antibiotic categories (Table 4). All the isolates were multidrug resistant. All the isolates (100%) were resistant to aminopenicillins (ampicillin, piperacillin) and cephalosporins except cefoxitin. In case of the ESBL inhibiting combination antibiotics, 65.42% and 65.74% of E. *coli* and 61.67% and 45.76% of *K. pneumoniae* were resistant to amoxicillin + clavulanic acid and to piperacillin + tazobactam, respectively.

Table 4
Antibiotic resistance of ESBL producing *E. coli* and *K. pneumoniae* strains.

Antibiotic group	Antibiotics (concentration in μg)	Resistance and susceptibility to the antibiotic			
		E. coli		K. pneumoniae	
		Resistance	Susceptible	Resistance	Susceptible
		n (%)	n (%)	n (%)	n (%)
Penicillin,	Ampicillin (10)	97(100)	0(0)	61(100)	0(0)
Penicillin and inhibitors	Piperacillin (100)	97(100)	0(0)	60(100)	0(0)
	Amoxicillin + acid clavulanic (30)	71(65.74)	37(34.26)	37(61.67)	24(38.13)
	Piperacillin + Tazobactam (110)	70(65.42)	37(34.58)	27(45.76)	32(54.24)
Cephalosporin	Cefazolin (30)	97(100)	0(0)	60(100)	0(0)
	Cefuroxime (30)	95(100)	0(0)	56(100)	0(0)
	Ceftriaxone (30)	95(100)	0(0)	60(100)	0(0)
	Ceftazidime (30)	95(100)	0(0)	60(100)	0(0)
	Cefepime (30)	95(100)	0(0)	55(98.21)	1(1.79)
	Cefoxitin (30)	43(40.57)	63(59.43)	14(23.73)	45(76.27)
Monobactam	Aztreonam (30)	92(94.85)	5(5.15)	53(94.64)	3(5.36)
Carbapenem	Meropenem (10)	17(15.74)	91(84.26)	5(8.19)	55(91.81)
	Imipenem (10)	22(20.75)	84(79.25)	5(8.19)	55(91.81)
	Ertapenem (10)	35(32.71)	72(67.29)	11(18.33)	49(81.67)
Aminoglycosides	Gentamycin (10)	46(44.66)	57(55.34)	31(5082)	30(49.18)
	Amikacin (30)	7(6.93)	94(93.7)	8(13.11)	53(86.89)
	Tobramycin (10)	74(71.15)	30(28.85)	35(57.37)	26(42.63)
	Kanamycin (30)	65(71.43)	26(28.57)	35(77.77)	10(22.23)
Macrolides	Azithromycin (15)	68(68.69)	31(31.31)	21(35.59)	38(64.41)
Quinolones, Fluoroquinolones	Ciprofloxacin (5)	98(95.15)	5(4.85)	56(91.80)	5(8.20)
	Ofloxacin (5)	59(67.05)	28(32.95)	11(24.44)	34(75.56)
	Levofloxacin (5)	71(71.72)	18(28.28)	32(53.33)	28(46.67)
	Pefloxacin	58(100)	0(0)	57(93.44)	4(6.56)
	Nalidixic acid (30)	99(94.29)	6(5.71)	40(88.89)	5(11.11)
	Norfloxacin (30)	68(80.95)	16(19.05)	30(50.0)	30(50.0)
Cyclins	Tetracycline (30)	80(86.02)	13(13.98)	36(78.26)	10(21.74)
	Doxycycline (30)	70(67.31)	34(32.69)	37(60.66)	24(39.34)
Sulfonamides	Sulfamethoxazole (50)	73(93.59)	5(6.41)	23(100)	0(0)
	Sulfamethoxazole + trimethoprim (25)	94(89.52)	11(10.48)	52(88.14)	7(11.86)
Nitrofurans	Nitrofurantoin (300)	42(40)	63(60)	24(40)	36(60)
Phosphonic acid	Fosfomycin (200)	12(11.43)	93(88.57)	36(61.02)	23(38.98)

High resistance rates were detected against aminoglycoside, quinolone, and fluoroquinolone antibiotic categories. Indeed, in the aminoglycoside family, resistance rates recorded were up to 71.43% in *E. coli* and 77.77% in *K. pneumoniae* against kanamycin. Isolates were more susceptible to amikacin since only 6.93% of resistance in *E. coli* and 13.11% in *K. pneumoniae* were recorded (Table 4). The resistance rates reported against quinolones and fluoroquinolones varied from 67.05–100% in *E. coli* and from 24.44–93.44% in *K. pneumoniae* (Table 4).

In case of carbapenems, 17 E. coli (15.74%) and 5 K. pneumoniae (8.19%) isolates were resistant to meropenem.

Other families of antibiotics commonly used in hospitals in Burkina Faso include cyclins, 86.02% *E. coli* and 78.26% *K. pneumoniae* isolates were resistant to tetracycline. In case of sulfonamides, 88.14% of *K. pneumoniae* isolates were resistant to Sulfamethoxazole + trimethoprim and 100% to

sulfamethoxazole (Table 4).

Azithromycin, an antibiotic widely used in Burkina Faso for Covid19 patient treatment (27, 28), was inactive for 68.69% of *E. coli* isolates and for 35.59% of *K. pneumoniae* isolates (Table 4)

# **Discussion**

β-lactams are widely used in treating patients in healthcare in Burkina Faso, but nowadays bacteria are often highly resistant to these antibiotics. Therefore, we isolated *E. coli* and *Klebsiella pneumoniae* strains from ESBL selective ChromAgar plates inoculated with healthcare center wastewaters to determine their susceptibility to other antibiotics available. Over 97% of the 84 wastewater samples analyzed contained ESBL producing *E. coli* and/or *K. pneumoniae*. The concentrations of ESBL producing Gram-negative bacteria in the healthcare center wastewaters were high, but, our results are comparable to those published in previous studies elsewhere (15, 29–31). For instance, concentrations up 10<sup>7</sup> CFU/ml of ESBL, CARB and OXA producing Enterobacteriaceae were reported from hospital wastewaters in Slovenia and Austria (15). Among our samples, the wastewaters collected from the tertiary and secondary level healthcare centers were the most contaminated with ESBL producers, possibly because these hospitals receive more patients, generally referred from a district level healthcare. Also, antibiotics are used more in terms of both quantity and diversity in tertiary and secondary level hospitals.

In addition to being ESBL producers, many of the isolates characterized in this study were also AmpC-β-lactamase (23 positive out of 51 isolates tested) and carbapenemase (10 positive out of 21 isolates tested) producers. Two types of carbapenemases, OXA-48 and NDM, were detected. Previously, using a metagenomics approach, presence of several carbapenemase genes (*bla*VIM, *bla*IMP, *bla*NDM and *bla*OXA-48) was reported in the wastewaters of some of the same hospitals in Burkina Faso (26). The resistance rate to carbapenems in our study was 15.74% to meropenem, 20.75% to imipenem and 32.71% to ertapenem. Our results differ from the results recently reported from Burkina Faso, where imipenem was the only carbapenem tested and no resistance to it was detected (25, 32). Occurrence and eventual spread of the carbapenem-resistant bacteria into the environment is of a particular concern since carbapenems are currently the antimicrobials of last resort in healthcare.

Wastewaters originating from healthcare centers present a public health concern in Burkina Faso and other countries, where they are discharged directly into the environment or into municipality wastewater channels without any prior treatment (20, 22). Furthermore, in LMICs, hospital wastewater may be used for irrigation of vegetable crops (5). Indeed, ESBL producing bacteria have been isolated on lettuce leaves in Burkina Faso (32). Common intestinal carriage of these bacteria can increase their prevalence in patients visiting healthcare centers, where presence of these bacteria leads to therapeutic treatment complications, prolonged patient hospitalizations and increased hospitalization costs, as well as higher mortality and morbidity (33).

The high level of resistance to the commonly used antibiotics has been reported also by other research groups in West Africa (34, 35). In Nigeria, full resistance to cefotaxime, cefpodoxime, sulfonamide and ertapenem was reported among ESBL producing *E. coli* isolated from wastewater originating from a healthcare facility (35). Likewise, in Côte d'Ivoire, ESBL producing *E. coli* and *K. pneumoniae* isolated from hospital wastewaters were reported to be fully resistant to amoxicilline + clavulanic acid, cefotaxime, ceftriaxone and ceftazidime, *E. coli* were also fully resistant and *K. pneumoniae* 62.5% resistant to ciprofloxacin, 87% resistant to nalidixic acid and cefepime, and furthermore, 76.5% of *E. coli* and 50% of *K. pneumonia* were resistant to gentamycin (34). The high resistance level of bacteria in wastewaters from healthcare centers is the consequence of antimicrobials misuse in hospitals, the discharge at high concentrations of not metabolized antibiotics and antibiotic residues into hospital wastewater, and the fecal contamination by patients (7, 22, 36–38). Furthermore, the high concentration of bacteria in these wastewaters offers an increased chance for horizontal transfer of resistance genes between bacteria (30–32, 37).

Amikacin, fosfomycin and nitrofurantoin were the antibiotics against which we recorded low resistance rates. Also in Mexico, a low resistance rate to amikacin among carbapenemase-producing *Klebsiella* spp. isolated from hospital wastewater was reported recently (40). These antibiotics, mostly used for urinary tract infection treatment, represent a major therapeutic option in case of infection with ESBL producing bacteria.

# Conclusion

This study shows that wastewaters from healthcare facilities represent a reservoir of multidrug-resistant bacteria in Burkina Faso. Wastewaters collected from the healthcare centers representing tertiary and secondary level of the healthcare system were the most contaminated. The ESBL producing *E. coli* and *K. pneumoniae* isolates were resistant to all commonly used antibiotics in Burkina Faso, such as  $\beta$ -lactams,  $\beta$ -lactams combined with ESBL inhibitors (amoxicillin + clavulanic acid and piperacillin + tazobactam), quinolones, fluoroquinolones, aminoglycosides, sulfonamides, cyclins, and macrolides. Only amikacin and fosfomycin showed good activity against these bacteria.

# **Material And Methods**

# Study design, study sites and sampling

A prospective study was carried out in 5 healthcare centers in Burkina Faso representing different healthcare system levels. The samples were collected from the Yalgado Ouédraogo teaching hospital in Ouagadougou (university hospital, tertiary level care), Koudougou regional hospital center in Koudougou and El Fateh SUKA Clinic in Ouagadougou (secondary level care), Source de Vie medical center in Ouagadougou and Saint Camille medical center in Nanoro rural area (primary level).

Three healthcare centers had a sewer system, Yalgado teaching hospital, Koudougou regional hospital center and Saint Camille medical center. Yalgado teaching hospital sewers connected to the city sewage system leading to the city's wastewater treatment plant. Wastewater from the hospital is discharged into the general sewage without any prior treatment. Koudougou regional hospital center has a chemical treatment device. The treated wastewater is discharged into the municipality channel, which is connected to a backwater in the town. Source de Vie medical center and El Fateh SUKA Clinic do not have a sewer system and their wastewater is collected in septic tanks. The management of wastewater in these two healthcare centers and in Saint Camille medical center in Nanoro are not clearly documented. As a rule, there is no wastewater treatment plants in rural areas in Burkina Faso, instead, the wastewaters are directly discharged into the environment without any treatment.

We collected wastewater samples from several sites along the sewers from the healthcare centers with the sewer system and from septic tanks from the healthcare centers without a sewer system. Two rounds of sampling were done, 1) October to December 2019 and 2) October 2020 to March 2021. A total of 84 wastewater samples were collected (Table 1). One liter of wastewater was collected in a sterile glass bottle. The samples were immediately placed in a cooler containing ice blocks and transported to analysis in the microbiology lab of the Nanoro clinical research unit (CRUN) within 12 hours.

# **Bacterial Count, Isolation, And Identification**

Two dilutions were prepared for each sample (1/10 and 1/100) using sterile 0.9% physiological saline water. Following the WHO Tricycle instructions (41), 100  $\mu$ l of each dilution was inoculated on ESBL selective agar plates (ChromAgar<sup>TM</sup> ESBL), which were incubated at 35 ± 2°C for 24 hours. A positive control was carried out for all samples by inoculating a non-selective CLED agar plate with 100  $\mu$ l of the sample. After incubation,, all visible bacterial colonies on the plates were counted, and the results were expressed in colony-forming units per milliliter of wastewater (CFU/ml). Only a plate of one dilution from each sample was considered for the bacterial count.

The agar plates were also inspected for different morphotypes of bacteria, according to the manufacturer's instructions (ChromAgar™ ESBL). Red or pink colonies were considered to be *E. coli*, and blue, green, or blue-green the KESC group (*Klebsiella, Enterobacter, Serratia and Citrobacter*). Five colonies of the same morphotype of *E. coli* or the KESC group were picked for purification on eosin methylene blue agar (EMB). The purified isolates were identified using the API20E (Biomérieux, France).

# **Antimicrobial Susceptibility Testing**

Antimicrobial susceptibility testing of 170 presumptive ESBL producing isolates (109 *E. coli* and 61 *K. pneumoniae* isolates) was performed using the disk diffusion method on Muller Hinton (MH) agar. Thirty-one antibiotic discs were tested (Table 4) and the results were interpreted according to the American Clinical and Laboratory Standards Institute (CLSI) 2021 guidelines.

# Extended Spectrum β-lactamase (Esbl) Confirmation

ESBL confirmation testing was carried out on MH agar using the double disc synergy test (DDST) between a 3rd generation cephalosporin (ceftriaxone or ceftazidime, C3G), a 4th generation cephalosporin (cefepime, C4G) and amoxicillin + clavulanic acid (AMC) following the CLSI 2021 guidelines. The result was interpreted as positive when there was a visible synergy inhibition zone between C3G-AMC-C4G (Fig. 1).

# Phenotypic Ampc-β-lactamase Testing

The 51 isolates (37 *E. coli* and 14 *K. pneumoniae*) which were resistant or intermediately susceptible to cefoxitin were tested for the AmpC-β-lactamase production. A bacterial suspension prepared with fresh colonies (McFarland 0.5) was inoculated on to entire surface of the MH agar supplemented with cloxacillin at  $4\mu g/l$  and a disk of cefoxitin was placed on the plate. The test was positive if the inhibition zone diameter around cefoxitin disc was  $\geq 18$  mm.

# **Carbapenemases Detection Test**

The isolates that were resistant or intermediately susceptible to meropenem were tested for carbapenemase production.with the immunochromatographic test O.K.N.V.I. RESIST-5 (CORIS BioConcept, Belgium), according to the manufacturer's instructions. A total of 21 isolates were tested for the five main carbapenemases (OXA-48-like, KPC, NDM, VIM, IMP) within 15 minutes.

# **Abbreviations**

AMC: amoxicillin + clavulanic acid; AMR: antimicrobial resistance; CAZ: ceftazidime; CFU: colony forming unit; CLED: Cystine Lactose Electrolyte Deficient; CLSI: Clinical and Laboratory Standards Institute; CTX: cefotaxime; CRUN: clinical research unit of Nanoro; C3G: third generation cephalosporin; C4G: fourth generation cephalosporine; DDST: double disc synergy test; EMB: eosin methylene blue; ESBL: Extended-spectrum β–lactamases; KESC: *Klebsiella Enterobacter Serratia* and *Citrobacter*, ITM: Institute of Tropical Medicine; MH: Muller Hinton;

# **Declarations**

## Ethics approval and consent to participate

This study received approval from the health research committee of Burkina Faso (N°153-12-2018/ CE-RS). Authorizations were obtained from all hospitals

## Consent for publication

Not applicable

## Availability of data and materials

All data generated or analyzed during this study are included in this published article

## Competing interests

The authors declare that they have no competing interests.

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## Authors' contributions

ZG, IOJB, KH, LS, and NB conceived and designed the study. ZG, NOM, collected samples. ZG, NOM, PPAV, MAB, IK, ALWT performed bacterial isolation and antimicrobial susceptibility test. HT Contributed for the reagents/materials/analysis tools. ZG, IOJB, HMN, KH and NB were the major contributors in writing the manuscript. All authors read and approved the final manuscript.

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# References

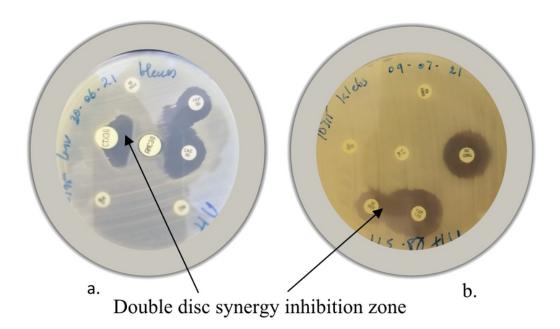
- 1. Sugden R, Kelly R, Davies S. Combatting antimicrobial resistance globally. Vol. 1, Nature Microbiology. Macmillan Publishers Limited; 2016. p. 1-2.
- 2. O'NEILL J. Book review: Tackling drug-resistant infections globally: final report and recommendations. London: Review on Antimicrobial Resistance. Arch Pharm Pract. 2016;7(3):110.
- 3. World Bank. World Bank. 2017. "Drug-Resistant Infections: A Threat to Our Economic Future." Washington, DC: World Bank. License: Creative Commons Attribution CC BY 3.0 IGO. 2017; Available from: www.worldbank.org
- 4. Michael CA, Dominey-Howes D, Labbate M. The antimicrobial resistance crisis: Causes, consequences, and management. Vol. 2, Frontiers in Public Health. 2014.
- 5. Ouedraogo AS, Pierre JH, Banuls AL, Ouedraogo R, Godreuil S. Émergence et diffusion de la résistance aux antibiotiques en Afrique de l'Ouest: facteurs favorisants et évaluation de la menace. Med Sante Trop [Internet]. 2017;27:147–54. Available from: https://pdfs.semanticscholar.org/ee82/5fd8b1d4a75b74ab734acdf136b0324ffffb.pdf
- 6. Hedgespeth ML, Sapozhnikova Y, Pennington P, Clum A, Fairey A, Wirth E. Pharmaceuticals and personal care products (PPCPs) in treated wastewater discharges into Charleston Harbor, South Carolina. Sci Total Environ. 2012 Oct 15;437:1–9.
- 7. Al Aukidy M, Verlicchi P, Voulvoulis N. A framework for the assessment of the environmental risk posed by pharmaceuticals originating from hospital effluents. Sci Total Environ. 2014 Sep 15;493:54–64.
- 8. Porooshat D. Antimicrobial Resistance: Implications and Costs. Infect Drug Resist. 2019;3903-10.
- 9. Chokshi A, Sifri Z, Cennimo D, Horng H. Global Contributors to Antibiotic Resistance. J Glob Infect Dis. 2019;Volume 11(1):36-42.

- 10. Chang X, Meyer MT, Liu X, Zhao Q, Chen H, Chen J an, et al. Determination of antibiotics in sewage from hospitals, nursery and slaughter house, wastewater treatment plant and source water in Chongqing region of Three Gorge Reservoir in China. Environ Pollut [Internet]. 2010;158(5):1444–50. Available from: http://dx.doi.org/10.1016/j.envpol.2009.12.034
- 11. Tariq M, Ali I, Ihsanullah I, Naushad M, Ali S, Hassan S, et al. Journal of Water Process Engineering Hospital wastewater as a source of environmental contamination: An overview of management practices, environmental risks, and treatment processes. J Water Process Eng [Internet]. 2021;41(January):101990. Available from: https://doi.org/10.1016/j.jwpe.2021.101990
- 12. Ekwanzala MD, Lehutso RF, Kasonga TK, Dewar JB, Momba MNB. Environmental dissemination of selected antibiotics from hospital wastewater to the aquatic environment. Antibiotics. 2020;9(7):1–16.
- 13. Chunhui Z, Liangliang W, Xiangyu G, Xudan H. Antibiotics in WWTP discharge into the Chaobai River, Beijing. Arch Environ Prot. 2016;42(4):48-57.
- 14. Aydin S, Aydin ME, Ulvi A, Kilic H. Antibiotics in hospital effluents: occurrence, contribution to urban wastewater, removal in a wastewater treatment plant, and environmental risk assessment. Environ Sci Pollut Res. 2019;26(1):544–58.
- 15. Rozman U, Duh D, Cimerman M, Turk SŠ. Hospital wastewater effluent: Hot spot for antibiotic resistant bacteria. J Water Sanit Hyg Dev. 2020 Jun 1;10(2):171–8.
- 16. Paulus GK, Hornstra LM, Alygizakis N, Slobodnik J, Thomaidis N, Medema G. The impact of on-site hospital wastewater treatment on the downstream communal wastewater system in terms of antibiotics and antibiotic resistance genes. Int J Hyg Environ Health [Internet]. 2019;222(4):635–44. Available from: https://doi.org/10.1016/j.ijheh.2019.01.004
- 17. Zurfluh K, Bagutti C, Brodmann P, Alt M, Schulze J, Fanning S, et al. Wastewater is a reservoir for clinically relevant carbapenemase- and 16s rRNA methylase-producing Enterobacteriaceae. Int J Antimicrob Agents [Internet]. 2017;50(3):436–40. Available from: http://dx.doi.org/10.1016/j.ijantimicag.2017.04.017
- 18. Tesfaye H, Alemayehu H, Desta AF, Eguale T. Antimicrobial susceptibility profile of selected Enterobacteriaceae in wastewater samples from health facilities, abattoir, downstream rivers and a WWTP in Addis Ababa, Ethiopia. Antimicrob Resist Infect Control. 2019;8(1):1–11.
- 19. Rodríguez EA, Garzón LM, Gómez ID, Jiménez JN. Multidrug resistance and diversity of resistance profiles in carbapenem-resistant Gram-negative bacilli throughout a wastewater treatment plant in Colombia. J Glob Antimicrob Resist [Internet]. 2020;22:358–66. Available from: https://doi.org/10.1016/j.jgar.2020.02.033
- 20. Al Aukidy M, Al Chalabi S, Verlicchi P. Hospital wastewater treatments adopted in Asia, Africa, and Australia. In: Handbook of Environmental Chemistry. Springer Verlag; 2018. p. 171–88.
- 21. Josué llunga Mubedi a, Naresh Devarajan b, c, Séverine Le Faucheur c, John Kayembe Mputu a, Emmanuel K. Atibu e, Periyasamy Sivalingam d, Kandasamy Prabakar b, Pius T. Mpiana e, Walter Wildi c J. Effects of untreated hospital effluents on the accumulation of toxic metals in sediments of receiving system und ... Related papers. 2013;
- 22. Hocquet D, Muller A, Bertrand X. What happens in hospitals does not stay in hospitals: antibiotic-resistant bacteria in hospital wastewater systems. Vol. 93, Journal of Hospital Infection. W.B. Saunders Ltd; 2016. p. 395–402.
- 23. Bougnom BP, Zongo C, McNally A, Ricci V, Etoa FX, Thiele-Bruhn S, et al. Wastewater used for urban agriculture in West Africa as a reservoir for antibacterial resistance dissemination. Environ Res [Internet]. 2019;168(September 2018):14–24. Available from: https://doi.org/10.1016/j.envres.2018.09.022
- 24. Bougnom BP, Thiele-Bruhn S, Ricci V, Zongo C, Piddock LJV. Raw wastewater irrigation for urban agriculture in three African cities increases the abundance of transferable antibiotic resistance genes in soil, including those encoding extended spectrum β-lactamases (ESBLs). Sci Total Environ [Internet]. 2020;698:134201. Available from: https://doi.org/10.1016/j.scitotenv.2019.134201
- 25. Abasse O, Boukaré K, Sampo E, Bouda R, CISSE H, Stéphane K, et al. Spread and antibiotic resistance profile of pathogens isolated from human and hospital wastewater in Ouagadougou. Microbes Infect Dis. 2021;0(0):0–0.
- 26. Markkanen MA, Haukka K, Pärnänen KMM, Dougnon VT, Bonkoungou IJO, Garba Z, et al. Metagenomic analysis of antimicrobial resistance genes in wastewaters in Benin and Burkina Faso indicates a serious health risk from untreated hospital wastewaters in low-income countries. medRxiv [Internet]. 2022;2021.10.19.21265183. Available from: https://www.medrxiv.org/content/10.1101/2021.10.19.21265183v7.abstract
- 27. Rouamba T, Barry H, Ouédraogo E, Tahita MC, Yaméogo NV, Poda A, et al. Safety of chloroquine or hydroxychloroquine plus azithromycin for the
- treatment of COVID-19 patients in Burkina Faso: An observational prospective cohort study. Ther Clin Risk Manag. 2021;17(August):1187–98.
- 28. Adama B, Armel P, Kadari C, Apoline K S, Boukary O, Abdoul Risgou O, et al. Effect of Hydroxychloroquine or Chloroquine and Azithromycin on COVID-19 Patients' Recovery and Mortality: Evidence from a Hospital Based Retrospective Cohort Study Conducted in Burkina Faso. J Infect Dis Epidemiol. 2021;7(2):1–9.
- 29. Le TH, Ng C, Chen H, Yi XZ, Koh TH, Barkham TMS, et al. Occurrences and characterization of antibiotic-resistant bacteria and genetic determinants of hospital wastewater in a tropical country. Antimicrob Agents Chemother. 2016;60(12):7449–56.
- 30. Galvin S, Boyle F, Hickey P, Vellinga A, Morris D, Cormican M. Enumeration and characterization of antimicrobial-resistant escherichia coli bacteria in effluent from municipal, hospital, and secondary treatment facility sources. Appl Environ Microbiol. 2010 Jul;76(14):4772–9.
- 31. Drieux L, Haenn S, Moulin L, Jarlier V. Quantitative evaluation of extendedspectrum β-lactamase-producing Escherichia coli strains in the wastewater of a French teaching hospital and relation to patient strain. Antimicrob Resist Infect Control. 2016;5(1).

- 32. Soré S, Sawadogo Y, Bonkoungou JI, Kaboré SP, Béogo S, Sawadogo C, et al. Detection, identification and characterization of extended-spectrum beta-lactamases producing Enterobacteriaceae in wastewater and salads marketed in Ouagadougou, Burkina Faso. Int J Biol Chem Sci. 2020;14(8):2746–57.
- 33. French GL. Clinical impact and relevance of antibiotic resistance. Adv Drug Deliv Rev. 2005;57(10):1514-27.
- 34. Guessennd N, Ouattara M, Ouattara N, Nevry R, Gbanon V, Tiekoura K, et al. Étude des bactéries multirésistantes des effluents hospitaliers d'un centre hospitalier et universitaire (CHU) de la ville d'Abidjan (Côte d'Ivoire). J Appl Biosci. 2013;69:5456.
- 35. Adekanmbi AO, Akinpelu MO, Olaposi AV, Oyelade AA. Diversity of Extended Spectrum Beta-lactamase (ESBL) genes in Escherichia coli isolated from wastewater generated by a Sick Bay located in a University Health Care Facility. Gene Reports. 2020 Sep 1;20.
- 36. Ye Q, Wu Q, Zhang S, Zhang J, Yang G, Wang H, et al. Antibiotic-resistant extended spectrum β-lactamase- and plasmid-mediated AmpC-producing enterobacteriaceae isolated from retail food products and the pearl river in Guangzhou, China. Front Microbiol. 2017;8(FEB):1–12.
- 37. sangadah khotimatus, Kartawidjaja J. Hospital wastewater treatment scenario around the globe. Orphanet J Rare Dis. 2020;21(1):1-9.
- 38. Egbule OS. Detection and Transfer of Extended Spectrum Beta Lactamase Enzymes from Untreated Hospital Waste Water. Adv Microbiol. 2016;06(07):512–20.
- 39. Czatzkowska M, Wolak I, Harnisz M, Korzeniewska E. Impact of Anthropogenic Activities on the Dissemination of ARGs in the Environment—A Review. Int J Environ Res Public Health. 2022;19(19).
- 40. Galarde-López M, Velazquez-Meza ME, Bobadilla-Del-valle M, Carrillo-Quiroz BA, Cornejo-Juárez P, Ponce-De-león A, et al. Surveillance of Antimicrobial Resistance in Hospital Wastewater: Identification of Carbapenemase-Producing Klebsiella spp. Antibiotics. 2022;11(3).
- 41. WHO. Global Tricycle Surveillance ESBL E.coli Integrated Global Surveillance on ESBL-producing E. coli Using a "One Health" Approach: Implementation and Opportunities [Internet]. World Health Organization, Global Antimicrobial Resistance Surveillance System (GLASS), Surveillance Prevention and Control. 2021. 76 p. Available from: http://apps.who.int/bookorders.

# **Figures**

Figure 1



ESBL confirmation test for a *K. pneumoniae* strain showing a double disc synergy inhibition zone.