

Antimicrobial Resistance in Zambia: A Systematic Review

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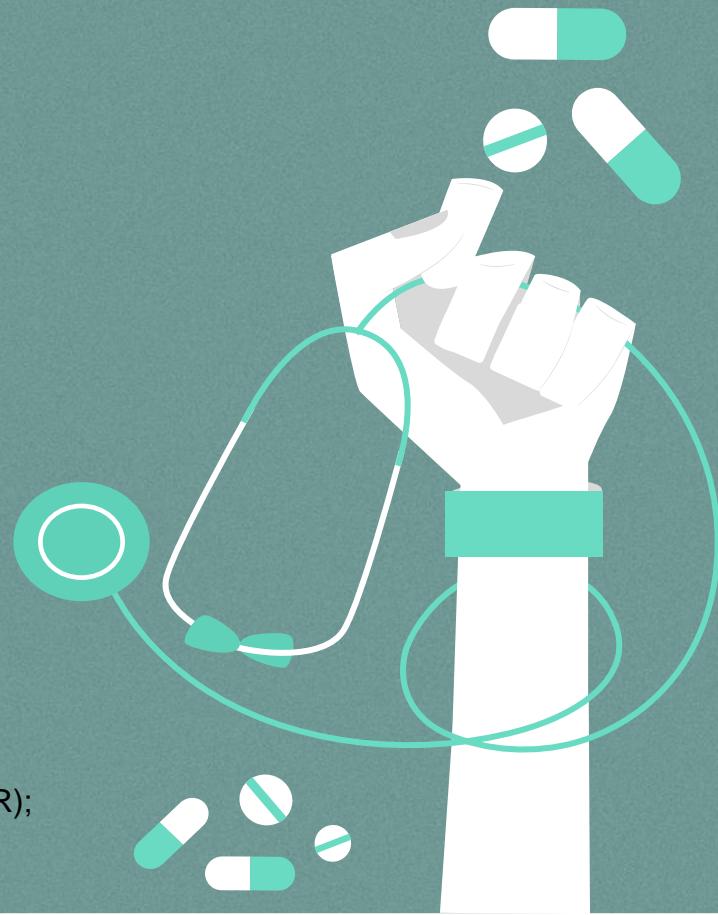
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BACKGROUND



Antimicrobials

- Antibiotics
- Antivirals
- Antifungals
- Antiparasitics



Antimicrobial Resistance (AMR)

Microorganisms **change over time** and no longer respond to medicines

Minor injuries and infections = deadly



Global Health Crisis

As dangerous as COVID-19 Pandemic²

Threatens to erode centuries of progress³



Zambia

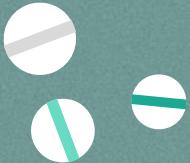
Epidemics: malaria, HIV, TB, COVID-19⁴

Access, use, control of antimicrobials in humans, animals and environment remain unknown



AIM

This study reviewed published studies of AMR prevalence across **human-animal-environmental** domains in Zambia.



METHODOLOGY

Search Terms:

“multiresistant OR multi-drug resistant OR antimicrobial resistance OR drug resistance OR bacterial resistance” “AND Zambia”, with spelling variations included

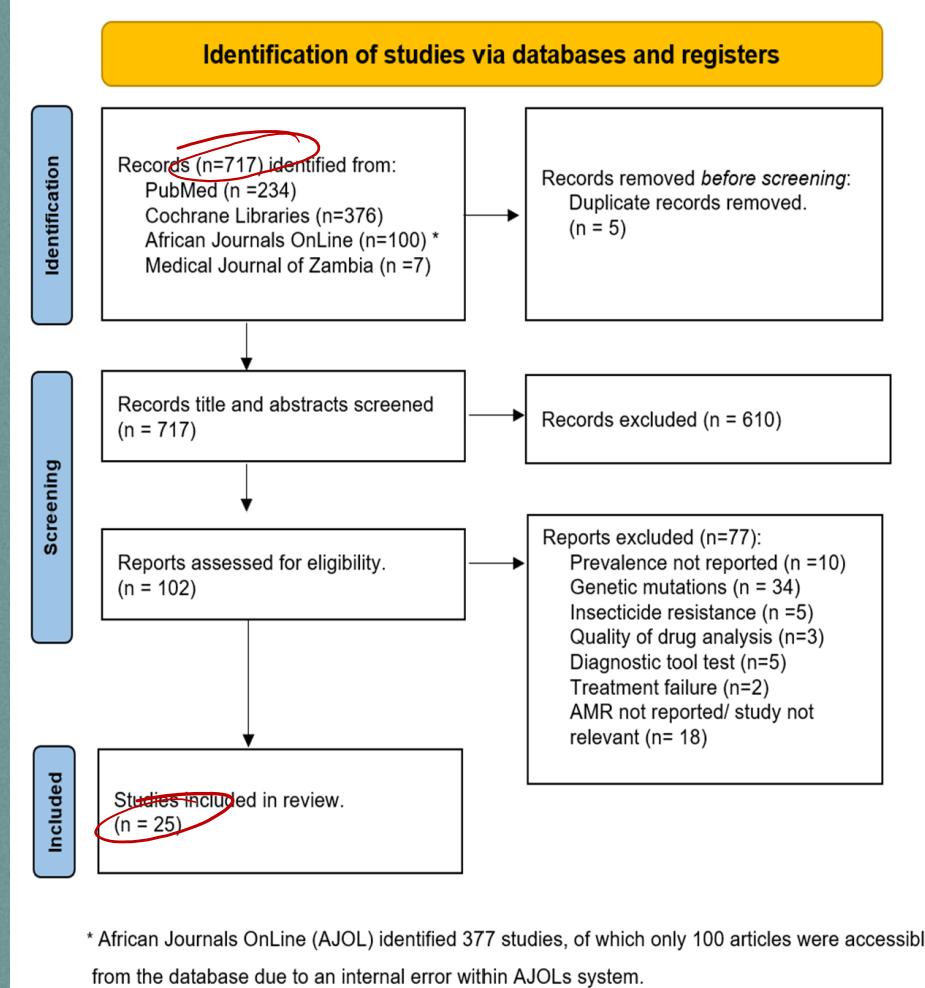


Figure 1: A PRISMA diagram illustrating the search strategy for the review

RESULTS

- Lusaka: more than half (16/25) of the studies.
- 6 of the 10 provinces in Zambia had no AMR data.
- 15 of the 25 studies were conducted only on the human sector

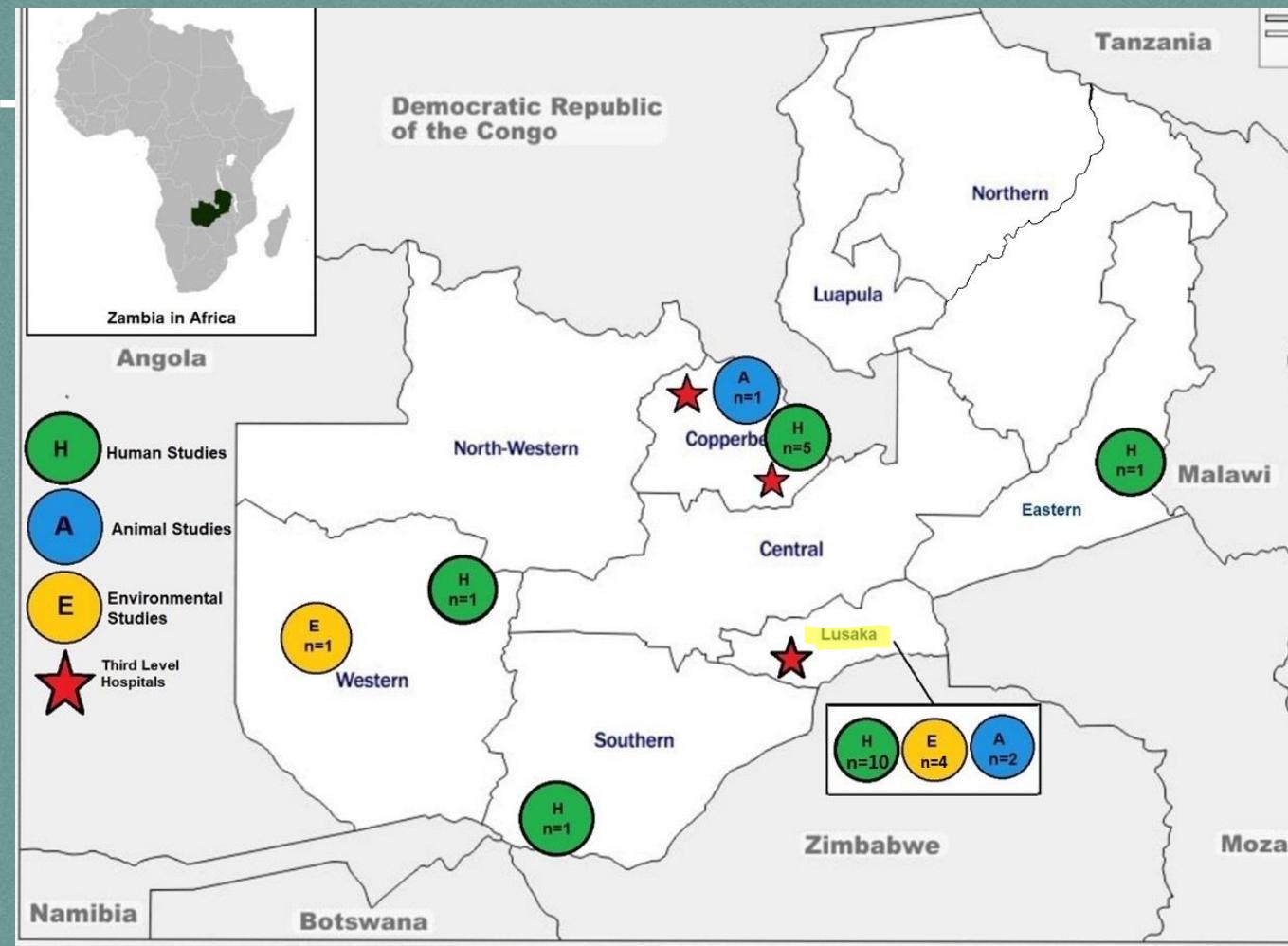


Figure 2: Distribution of studies in Zambia



RESULTS

21 microbes:

- *S. aureus* : most common
- *E. coli* : 11 studies
- *Klebsiella species* : 7 studies
- *M. tuberculosis* : 5 studies
- HIV : 3 studies

Antibacterial studies: 21/25

Antiviral studies: 3/25

Antiparasitic: 1/25

No fungal conditions

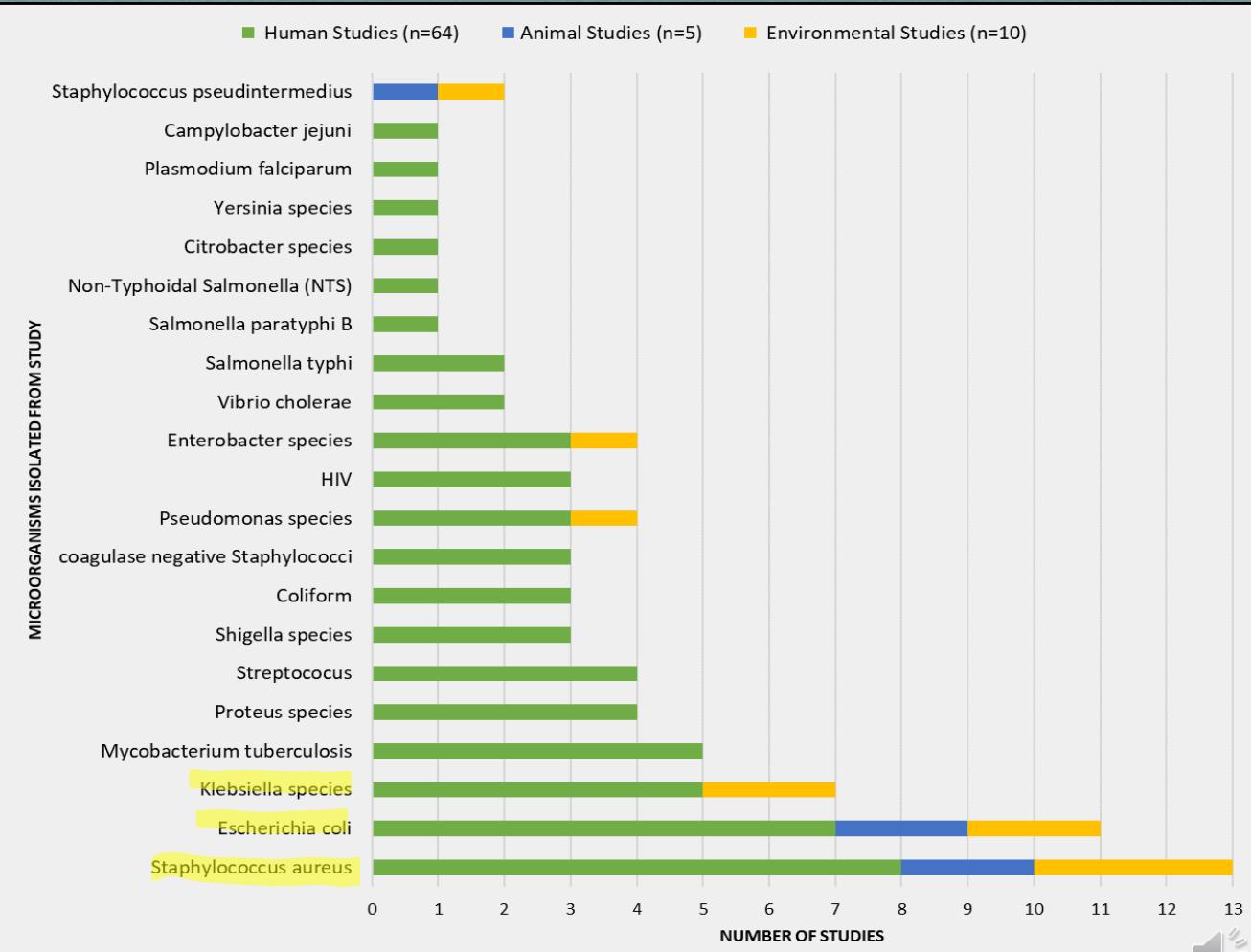


Figure 3: Microorganisms Studied by Sector



RESULTS

Table 1: Resistance rates against antibiotics used for two widely surveyed pathogens: *E. coli* and *Klebsiella spp.*

Legend	Gram Negative Bacteria																	
	<i>E. coli</i>										<i>Klebsiella</i>							
	Human					Animal		Environment			Human				Environment			
	Mainda et al.	Chiyangi et al.	Mainda et al.	Kabwe et al.	Chanda et al.	Chanda et al.	Chanda et al.	Mainda et al.	Songe et al.	Mwamun- gule et al.	Kabwe et al.	Chanda et al.	Chanda et al.	Chanda et al.	Ziwa et al.	Ziwa et al.	Mwamun- gule et al.	
100%																		
50-99%																		
10-49%																		
0-9%																		
Gentamycin	27%			1%	100%				1%	91%	4%	96%				30%	30%	0%
Streptomycin		34%								88%								
Kanamycin	99%																	
Amikacin	47%															20%	20%	
Cefpodoxime	29%	67%	1%					2%								100%	100%	
Cefotaxime	25%	67%		100%					100%		96%					100%	100%	
Cefalexin										100%								50%
Ceftriaxone	24%			100%							94%					40%	40%	
Ceftazidime	28%	67%		100%	40%		50%		95%		97%		69%			100%	100%	
Ampicillin	61%	67%	8%	100%				6%	100%	50%	99%					100%	100%	
Amoxicillin-clavulanic acid	46%			50%							93%							
Trimethoprim-Sulfamethoxazole			5%	100%				4%	74%		100%							
Co-trimoxazole	57%	100%			67%	100%						100%	0%			90%	90%	100%
Ciprofloxacin	20%		0%	80%	57%	63%	100%	0%	95%	0%	71%	33%	68%	100%	28%	28%		
Nalidixic acid		16%			43%	76%			93%			33%	56%					
Ceftazidime						61%						50%						
Norfloxacin					30%	62%	100%		91%			33%	40%					
Erythromycin				100%							92%					32%	32%	
Imipenem			0%								1%					0%	0%	
Nitrofurantoin					33%	48%	0%					33%	75%					
Tetracycline	51%	50%	14%					11%	91%	0%						40%	40%	50%
Chloramphenicol	22%	16%		60%	67%	26%			81%	0%	71%	0%	44%	100%	21%	21%	50%	



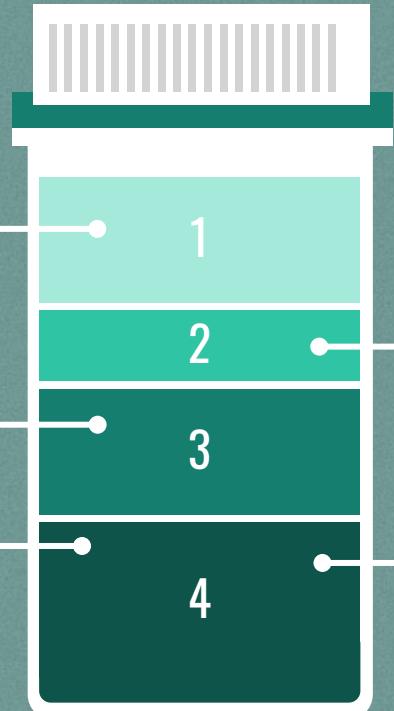


CONCLUSION

The level of resistance to commonly prescribed antibiotics is often **extremely high**.

There is a **bias** across many reference centers that don't reflect the disease condition.

Address the gaps in AMR diagnostic standardization and reporting.



More AMR data is needed for many provinces.

Improve surveillance, stewardship, infection control, and implementations of updated treatment guidelines and monitoring.



Thank you!



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