

ADVANCING THE LABORATORY PROFESSION AND NETWORKS IN AFRICA

ANTIMICROBIAL RESISTANCE (AMR) COMMUNITY OF PRACTICE (CoP)



The Rise of Resistance: How Bacteria "Outsmart" Antibiotics

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Outline

- Background
- Antimicrobials overview
- Antibiotics mechanisms of action
- Acquisition of resistance
- Antibiotics mechanisms of resistance
- Beta-lactamases
 - Classification
 - Detection methods



- The discovery of antibiotics, made us think the battle against infectious diseases has been won
- However, as many bacteria have developed resistance to multiple antimicrobial agents, the war now seems to be shifting in favour of the bacteria
- Antimicrobial Resistance (AMR) is a major threat to global health
- The World Health Organization (WHO) has declared it as one of the <u>top 10</u> <u>global public health threats</u> confronting humanity



 Globally, an estimated 4.95 million deaths were associated with drug-resistant infections, both directly and indirectly, with 1.27 million of these deaths directly attributable to drug resistance

• This indicates that over one million people died because their infections could not be treated with any available medicines due to resistance

 Seventy percent of these deaths resulted from severe infections that no longer responded to first-line antibiotics used for empirical treatment

Antimicrobials





Mechanisms of action





Causes of resistance





Over-prescribing of antibiotics



Poor infection control in hospitals and clinics



Patients not finishing their treatment



Lack of hygiene and poor sanitation



Over-use of antibiotics in livestock and fish farming



Lack of new antibiotics being developed

www.who.int/drugresistance

#AntibioticResistance



Acquisition of resistance genes

AFRICAN SOCIETY FOR LABORATORY MEDICINE

VERTICAL GENE TRANSFER

HORIZONTAL GENE TRANSFER



Trends in Microbiology

Mechanisms of resistance



INTRINSIC RESISTANCE

Natural

Wild-type

ACQUIRED RESISTANCE



Classification of betalactams





Betalactamases-Ambler classification



Classification of β -Lactamases



Ambler class A



Penicillinases (Subgroup 2a)

- β-lactamases with a relatively limited spectrum of hydrolytic activity
- Predominant β-lactamases in Gram-positive cocci, including the staphylococci (blaZ)
- Hydrolyze benzyl-penicillin and many penicillin derivatives, with poor hydrolysis of cephalosporins, carbapenems, or monobactams
- Inhibited by beta-lactamase inhibitors

• Subgroup 2b β-lactamases

- Hydrolyze penicillins and early cephalosporins, such as cephaloridine and cephalothin
- Strongly inhibited by clavulanic acid and tazobactam.
- They include the TEM-1, TEM-2, and SHV-1 enzymes, the most common plasmid-mediated β -lactamases identified in the 1970s and early 1980s
- Mainly found in gram negative bacteria

Phenotypic β-lactamase detection

- Phenotypic Methods
- 1. Nitrocefin Test:
 - **Procedure**: Use a chromogenic cephalosporin (nitrocefin) which changes colour when hydrolyzed by beta-lactamase
 - Interpretation: A colour change (yellow to red) indicates beta-lactamase activity
 - QC: S.aureus 29213-Positive
 - S.aureus 25923 Negative
- 2. Combination Disk Test:
 - **Procedure**: Use disks containing a beta-lactam antibiotic alone and in combination with a beta-lactamase inhibitor.
 - Interpretation: An increase in the zone of inhibition with the inhibitor indicates the presence of beta-lactamase
- NOTE: A positive test means penicillin, ampicillin, amoxicillin and piperacillin

Tests indicated for *S. aureus* isolates that test susceptible to Penicillin before reporting





AMBLER CLASS A-Broad spectrum



Group 2be

- Produce extended spectrum of activity
- Also known as ESBL
- Inactivates penicillins,1st to 3rd gen cephalosporins, monobactams but not carbapenems
- ESBL types are diverse, encoded by TEM,SHV,CTX-M variants
- Inhibited by beta-lactamase inhibitors



Phenotypic β-lactamase detection



- Double-Disk Synergy Test (DDST)/Clavunate inhibition Test:
 - **Procedure**: Place a beta-lactam antibiotic disk and a beta-lactamase inhibitor disk close to each other on an agar plate inoculated with the test organism
 - Interpretation: Enhanced inhibition near the inhibitor disk indicates beta-lactamase production



Quality control for ESBL

- Escherichia coli ATCC 25922 ≤ 2 mm increase in zone diameter for antimicrobial agent tested in combination with clavulanate vs the zone diameter when tested alone.
- Klebsiella pneumoniae ATCC 700603 ≥5mm increase in zone diameter of ceftazidimeclavulanate vs ceftazidime alone.
- ≥3mm increase in zone diameter of cefotaxime-clavulanate vs cefotaxime alone.

AMBLER CLASS C



- AmpC beta-lactamases are classified as Group 1 according to the Bush-Jacoby-Medeiros classification
- Provide resistance to most penicillins, cephalosporins (except for fourth-generation cephalosporins like cefepime), cephamycins (e.g., cefoxitin), and monobactams (e.g., aztreonam)
- They do not hydrolyze carbapenems, but AmpC-producing organisms may have additional resistance mechanisms that confer resistance to these drugs
- The genes encoding AmpC beta-lactamases are either chromosomal or plasmid-mediated
 - Chromosomal AmpC genes are typically inducible and are found in species such as Serratia marcescens, Providencia spp, Indole(+)Proteus, Citrobacter freundii, Enterobacter spp, and Morganella morganii.
 - Plasmid-mediated AmpC genes, such as blaCMY, blaACT, and blaDHA, can spread between different bacterial species.

Phenotypic β-lactamase detection



- There are no CLSI guidelines for detecting AmpC-mediated resistance in Gram-negative clinical isolates, which often leads to misleading results, particularly with phenotypic tests
- Detection of AmpC production in organisms with inducible chromosomal AmpC β -lactamase is unnecessary
- Phenotypic Methods:
 - Testing for AmpC production can include the use of cefoxitin disks in disk diffusion tests, which can indicate resistance due to AmpC production
- Molecular Methods:
 - PCR and sequencing can identify specific genes encoding AmpC beta-lactamases

Mapping of Carbapenem resistant Enterobacteriales









- Carbapenemase producing isolates of Enterobacterales usually test intermediate or resistant to one or more carbapenems
- Ertapenem non susceptibility is the most sensitive indicator of carbapenemases production
- Imipenem or meropenem MICs 2-4 ug/ml or Ertapenem MIC 2 ug/ml
- Quality control strains
 - E.coli ATCC 25922
 - K. pneumoniae ATCC BAA 1705 KPC positive
 - *K. pneumoniae* ATCC BAA 1706: Carbapenemase negative
 - K. pneumoniae ATCC BAA 2146:NDM positive (For use with eCIM)

Carbapenemases detection



| | Tests | Used for Epidemiological or Infec | tion Prevention-Related Te | sting |
|-------------|---|--|--|--|
| | CarbaNP | mCIM | mCIM With eCIM | |
| | (Table 3B) | (Table 3C) | (Table 3C) | Other (eg, molecular assays) |
| Organisms | Enterobacterales and <i>P. aeruginosa</i> that are not susceptible to one or more carbapenems | Enterobacterales and <i>P. aeruginosa</i> that are not susceptible to one or more carbapenems | Enterobacterales that are positive by mCIM | Enterobacterales and <i>P. aeruginosa</i> that are not susceptible to one or more carbapenems to determine the presence of a carbapenemase, or to determine carbapenemase type in isolates positive by CarbaNP or mCIM. |
| Strengths | Rapid | No special reagents or media necessary | No special reagents or media necessary | Determines type of carbapenemase in addition to absence or presence of the enzyme |
| Limitations | Special reagents are needed, some of which necessitate in- house preparation (and have a short shelf life). Invalid results occur with some isolates. Certain carbapenemase types (eg, OXA-type, chromosomally encoded) are not consistently detected. | Requires overnight incubation | Requires overnight incubation | Special reagents and equipment are needed. Specific to targeted genes; false-negative result if specific carbapenemase gene present is not targeted. |

Abbreviations: eCIM, EDTA-modified carbapenem inactivation method; mCIM, modified carbapenem inactivation method, MIC, minimal inhibitory concentration.

Carbapenemase detection methods





Result: Negative for carbapenemases Report: Carbapenemase not detected







Result: Positive mCIM and eCIM Report: Metallo-b-lactamase

CLSI 2023 http://clsi.org>

Carbapenemase detection methods









| Organism Quantity: | Selected Organism: Klebsiella pneumoniae |
|--------------------|--|
| Comments: | |

| Identification Information | | | | |
|----------------------------|------------|-----------------------|-----|----------|
| Organism Origin | MYLA® | | | |
| Selected Occurring | Klebsiella | pneumoniae | | |
| Selected Organism | Entered: | Jul 9, 2024 12:23 CAT | By: | Download |
| Analysis Messages: | 1 | | | |

| Susceptibility | Card: | AST-N | 255 Lot Numbe | r: 6552721403 | Expires: | May 9, 2 | 2025 12:00 CAT | |
|-----------------------------------|---------------|-----------|-----------------|-----------------------------------|--------------------------|----------------------------|----------------|--|
| Information | Status: | Final | Analysis Ti | me: 9.18 hours | Complete | ted: Jul 9, 2024 21:52 CAT | | |
| Antimicrobial | | MIC | Interpretation | Antimicro | bial | MIC | Interpretation | |
| Ampicillin | | >= 32 | R | lmipenem | | >= 16 | R | |
| Amoxicillin/Clavulanic A | cid | >= 32 | R | Meropenem | _ | >= 16 | R | |
| Piperacillin/Tazobactam | | >= 128 | R | Amikacin | | >= 64 | R | |
| Cefuroxime | | >= 64 | R | Gentamicin | | >= 16 | R | |
| Cefuroxime Axetil | | >= 64 | R | Ciprofloxacin | | >= 4 | R | |
| Cefoxitin | | >= 64 | R | Tigecycline | | 4 | I | |
| Cefotaxime | and the state | >= 64 | R | Nitrofurantoin | | >= 512 | R | |
| Ceftazidime | 1979 E. | >= 64 | R | Colistin | | <= 0.5 | S | |
| Cefepime | | >= 64 | R | Trimethoprim/ Sulfamethoxazole | | >= 320 | R | |
| Ertapenem | | >= 8 | R | | | | | |
| AES Findings: | Last N | 1odified: | Oct 24, 2023 12 | :59 CAT Param | eter Set: | NHLS TSH | WANE | |
| Confidence Level: | Consis | tent | | | | | | |
| Phenotypes flagged for review: | BETA- | LACTAMS | | CARBAPENEM/ CARBA (+ESBL | ASE (+ OR - OR +HL An | ESBL),IMPE npC) | RMEABILITY | |



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Selected Organism: Klebsiella pneumoniae

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|-----------------------|--|
| Comments: | |
| comments. | |
| | |

| Identification Information | | | | |
|----------------------------|---------------------------------|-----|----------|--|
| Organism Origin | MYLA® | | | |
| | Klebsiella pneumoniae | | | |
| Selected Organism | Entered: Jun 26, 2024 11:45 CAT | By: | Download | |
| Analysis Messages: | | | | |

| Susceptibility | Card: | AST-N | 1255 Lot Number | r: 6552721403 | Expires: | May 9, | 2025 12:00 CAT | | |
|--------------------------|---------|--------|-----------------|-----------------------------------|----------|-----------------------------------|----------------|--|--|
| Information | Status: | Final | Analysis Ti | Analysis Time: 8.73 hours | | Completed: Jun 26, 2024 20:48 CAT | | | |
| Antimicrobial | 5.5 | MIC | Interpretation | Antimicro | bial | MIC | Interpretation | | |
| Ampicillin | | >= 32 | R | Imipenem | | <= 0.25 | S | | |
| Amoxicillin/Clavulanic A | cid | 16 | *R | Meropenem | | <= 0.25 | S | | |
| Piperacillin/Tazobactam | 100 100 | 8 | S | Amikacin | | <= 2 | S | | |
| Cefuroxime | | >= 64 | R | Gentamicin | | <= 1 | S | | |
| Cefuroxime Axetil | | >= 64 | R | Ciprofloxacin | | 1 | R | | |
| Cefoxitin | | <= 4 | S | Tigecycline | | 2 | S | | |
| Cefotaxime | | >= 64 | R | Nitrofurantoin | | 64 | I | | |
| Ceftazidime | | 16 | R | Colistin | | <= 0.5 | S | | |
| Cefepime | | 2 | *R | Trimethoprim/ Sulfamethoxazole | | >= 320 | R | | |
| Ertapenem | | <= 0.5 | S | | | | | | |

| AES Findings: | Last Modified: | Oct 24, 2023 | 12:59 CAT | Parameter Set: NHLS TSHW | ANE |
|------------------------|----------------|--------------|-----------|----------------------------|-----|
| Confidence Level: | Consistent | | | | |
| Phenotypes flagged for | BETA-LACTAMS | | EXTEN | DED SPECTRUM BETA-LACTAMAS | SE |
| review: | | | - | | |



| Organism Quantity; | Selected Organism: Staphylococcus aureus | |
|--------------------|--|---|
| | | _ |
| Comments: | | |
| | | |

| Identification Information | | | | |
|----------------------------|--------------------------------|-----|----------|--|
| Organism Origin | MYLA® | | | |
| 51 | Staphylococcus aureus | | | |
| Selected Organism | Entered: Jul 9, 2024 12:01 CAT | By: | Download | |
| | | | | |

Analysis Messages:

The following antibiotic(s) are not claimed: Ampicillin, Gentamicin High Level (synergy), Streptomycin High Level (synergy).

| Susceptibility | Card: AST- | P603 Lot Number | r: 4832749403 | Expires: | Jun 6, 2 | 2025 12:00 CAT | | |
|--------------------------------------|---------------|-----------------|-----------------------------------|----------|--------------------------------|----------------|--|--|
| Information | Status: Final | Analysis Ti | Analysis Time: 12.40 hours Con | | pleted: Jul 10, 2024 00:45 CAT | | | |
| Antimicrobial | MIC | Interpretation | Antimicro | bial | MIC. | Interpretation | | |
| Cefoxitin Screen | NEG | - | Clindamycin | | <= 0.25 | S | | |
| Benzylpenicillin | 0.12 | S | Linezolid | | 2 | S | | |
| Ampicillin | | | Teicoplanin | | <= 0.5 | S | | |
| Oxacillin | 0.5 | S | Vancomycin | | 1 | S | | |
| Gentamicin High Level (synergy) | | | Tetracycline | | >= 16 | R | | |
| Streptomycin High Level (synergy) | | | Tigecycline | | <= 0.12 | S | | |
| Gentamicin | <= 0.5 | S | Fusidic Acid | | <= 0.5 | S | | |
| Ciprofloxacin | <= 0.5 | S | Mupirocin | | <-2 | | | |
| Moxifloxacin | <= 0.25 | S | Rifampicin | | <= 0.5 | S | | |
| Inducible Clindamycin Resistance | NEG | | Trimethoprim/ Sulfamethoxazole | | >= 320 | R | | |
| Erythromycin | <= 0.25 | S | | | | | | |



| omments: | |
|---|--|
| dentification Information | |
| Organism Origin MYLA® | |
| Staphylococcus aureus | |
| Entered: Jul 9, 2024 12:01 CAT By: Download | |

Analysis Messages:

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The following antibiotic(s) are not claimed: Ampicillin, Gentamicin High Level (synergy), Streptomycin High Level (synergy),

| Susceptibility | Card: AST-F | 603 Lot Number | 4832749403 | Expires: | Jun 6, 2 | 025 12:00 CAT | |
|--------------------------------------|---------------|----------------|-----------------------------------|----------|-----------------------------------|----------------|--|
| Information | Status: Final | Analysis Tin | Analysis Time: 12.65 hours | | Completed: Jul 10, 2024 01:00 CAT | | |
| Antimicrobial | MIC | Interpretation | Antimicro | bial | MIC | Interpretation | |
| Cefoxitin Screen | NEG | - | Clindamycin | | <= 0.25 | S | |
| Benzylpenicillin | >= 0.5 | R | Linezolid | | 2 | S | |
| Ampicillin | | | Teicoplanin | | <= 0.5 | S | |
| Oxacillin | 0.5 | S | Vancomycin | | <= 0.5 | S | |
| Gentamicin High Level (synergy) | | | Tetracycline | | <=] | S | |
| Streptomycin High Level (synergy) | | | Tigecycline | | <= 0.12 | S | |
| Gentamicin | 4 | •R | Fusidic Acid | | <= 0.5 | S | |
| Ciprofloxacin | <= 0.5 | S | Mupirocin | 1 | <= 2 | | |
| Moxifloxacin | <= 0.25 | S | Rifampicin | | <= 0.5 | S | |
| Inducible Clindamycin Resistance | NEG | | Trimethoprim/ Sulfamethoxazole | | 160 | R | |
| Erythromycin | <= 0.25 | S | | | | | |



| Organism Quantity: | Selected Organism: Staphylococcus aureus |
|--------------------|--|
| | |
| Comments: | |
| | |

| Identification Information | -0.00 | |
|----------------------------|---------------------------------|--------------|
| Organism Origin | MYLA® | |
| | Staphylococcus aureus | |
| Selected Organism | Entered: Jul 10, 2024 14:44 CAT | By: Download |

Analysis Messages:

The following antibiotic(s) are not claimed: Ampicillin, Gentamicin High Level (synergy), Streptomycin High Level (synergy),

| Susceptibility | Card | : AST-I | P603 | Lot Number | 4832749403 | Expires: | Jun 6, 2 | 2025 12:00 CAT |
|--------------------------------------|-------|----------|------|----------------------------|-----------------------------------|-----------------------------------|----------|----------------|
| Information | Statu | s: Final | _ | Analysis Time: 12.87 hours | | Completed: Jul 11, 2024 04:00 CAT | | |
| Antimicrobial | | MIC | Inte | erpretation | Antimicro | bial | MIC | Interpretation |
| Cefoxitin Screen | | POS | | + | Clindamycin | | <= 0.25 | *R |
| Benzylpenicillin | | >= 0.5 | | R | Linezolid | | 2 | S |
| Ampicillin | | | | | Teicoplanin | | <= 0.5 | S |
| Oxacillin | | >= 4 | | R | Vancomycin | | <= 0.5 | S |
| Gentamicin High Level (synergy) | | | | | Tetracycline | | >= 16 | R |
| Streptomycin High Level (synergy) | | | | - | Tigecycline | _ | 0.25 | S |
| Gentamicin | | >= 16 | | R | Fusidic Acid | | <= 0.5 | S |
| Ciprofloxacin | | >= 8 | | R | Mupirocin | | >= 8 | |
| Moxifloxacin | - 2 | 1 | | I | Rifampicin | | <= 0.5 | S |
| | | - | | • | Trimethoprim/ Sulfamethoxazole | | >= 320 | R |
| Erythromycin | | >= 8 | | R | | | | |



| Arganism Quantity: | | Selected Organism: Staphylococcus aureus | |
|----------------------|--------|--|--|
| | MRSA | | |
| Comments: | | | |
| Identification Infor | mation | | |
| Organism Origin | | MYLA® | |

By: Download

Selected Organism

Entered: Jul 10, 2024 14:44 CAT

Staphylococcus aureus

Analysis Messages:

The following antibiotic(s) are not claimed:

Ampicillin, Gentamicin High Level (synergy), Streptomycin High Level (synergy),

This isolate is presumed to be resistant based on detection of inducible clindamycin resistance.

| Susceptibility | Card: AST- | P603 Lot Nu | mber: 4832749403 | Expires: | Jun 6, 1 | 2025 12:00 CAT | |
|--------------------------------------|---------------|-------------|-----------------------------------|----------|-----------------------------------|----------------|--|
| Information | Status: Final | Analys | Analysis Time: 12.87 hours | | Completed: Jul 11, 2024 04:00 CAT | | |
| Antimicrobial | MIC | Interpreta | tion Antimic | robial | MIC | Interpretation | |
| Cefoxitin Screen | POS | + | Clindamycin | | <= 0.25 | *R | |
| Benzylpenicillin | >= 0.5 | R | Linezolid | | 2 | 5 | |
| Ampicillin | | | Teicoplanin | | <= 0.5 | S | |
| Oxacillin | >= 4 | R | Vancomycin | | <= 0.5 | S | |
| Gentamicin High Level (synergy) | | | Tetracycline | | >= 16 | R | |
| Streptomycin High Level (synergy) | | | Tigecycline | | 0.25 | S | |
| Gentamicin | >= 16 | R | Fusidic Acid | | <= 0.5 | S | |
| Ciprofloxacin | >= 8 | R | Mupirocin | | >= 8 | | |
| Moxifloxacin | 1 | 1 | Rifampicin | | <= 0.5 | S | |
| Inducible Clindamycin Resistance | POS | + | Trimethoprim/ Sulfamethoxazole | | >= 320 | R | |
| Erythromycin | >= 8 | R | | | | | |

*- AES modified **= User modified

| AES Findings: | Last Modified: | Oct 24, 2023 | 12:59 CAT | Parameter Set: NHLS TSHWANE |
|-----------------------------------|----------------|--------------|-----------|--|
| Confidence Level: | Consistent | | | |
| Phenotypes flagged for review: | BETA-LACTAMS | | MODIF | TCATION OF PBP (mecA) |
| | MUPIROCIN | | LOWL | EVEL RESISTANCE, HIGH LEVEL RESISTANCE |



| rganism Quantity: | Selected Organism: Serratia marces | (cens |
|----------------------------|------------------------------------|--------------|
| Comments: | | |
| Identification Information | | |
| Organism Origin | MYLA® | |
| E-losted Operation | Serratia marcescens | |
| Selected Organism | Entered: Jul 5, 2024 16:26 CAT | By: Download |
| Analysis Messages: | | |

The following antibiotic(s) are suppressed from analysis: Ampicillin, Piperacillin/Tazobactam, Imipenem,

| Susceptibility | Card: | AST-N | 1255 Lot Numb | er: 6552721403 | Expires: | May 9, | 2025 12:00 CAT | |
|-----------------------|----------|--------|----------------|-----------------------------------|----------|----------------------------------|----------------|--|
| Information | Status | Final | Analysis T | Analysis Time: 15.85 hours | | Completed: Jul 6, 2024 08:40 CAT | | |
| Antimicrobial | | MIC | Interpretation | Antimicrobial | | MIC | Interpretation | |
| Ampicillin | | | | Imipenem | | | | |
| Amoxicillin/Clavular | nic Acid | 16 | •R | Meropenem | | <= 0.25 | S | |
| Piperacillin/Tazobact | am | | | Amikacin | | <= 2 | *R | |
| Cefuroxime | 26.0 | 16 | •R | Gentamicin | | <= | S | |
| Cefuroxime Axetil | | 16 | •R | Ciprofloxacin | | <= 0.25 | S | |
| Cefoxitin | | 8 | *R | Tigecycline | | 1 | *R | |
| Cefotaxime | | <= 1 | S | Nitrofurantoin | | 256 | R | |
| Ceftazidime | | <-1 | S | Colistin | | >- 16 | R | |
| Cefepime | | <= 1 | S | Trimethoprim/ Sulfamethoxazole | | <= 20 | S | |
| Ertapenem | | <= 0.5 | S | | | | | |





- Antimicrobial Resistance (AMR) is a public health threat
- It is associated with high morbidity and mortality
- Widespread use of beta-lactams has led to development of resistance
- The choice of testing method to be used is based on the easy of use, and timeliness
- Appropriate use of antibiotic at the appropriate dosage and for the appropriate duration—is one important means of reducing the selective pressure that helps resistant organisms emerge
- Early appropriate laboratory detection of resistance will assist in prompt appropriate management of patients



ADVANCING THE LABORATORY PROFESSION AND NETWORKS IN AFRICA

ANTIMICROBIAL RESISTANCE (AMR) COMMUNITY OF PRACTICE (CoP)



THANK YOU

