

JASPI

Journal of Antimicrobial Stewardship Practices
and Infectious diseases

VOLUME 2 | ISSUE 3 | SUPPL. | 30 SEPTEMBER 2024

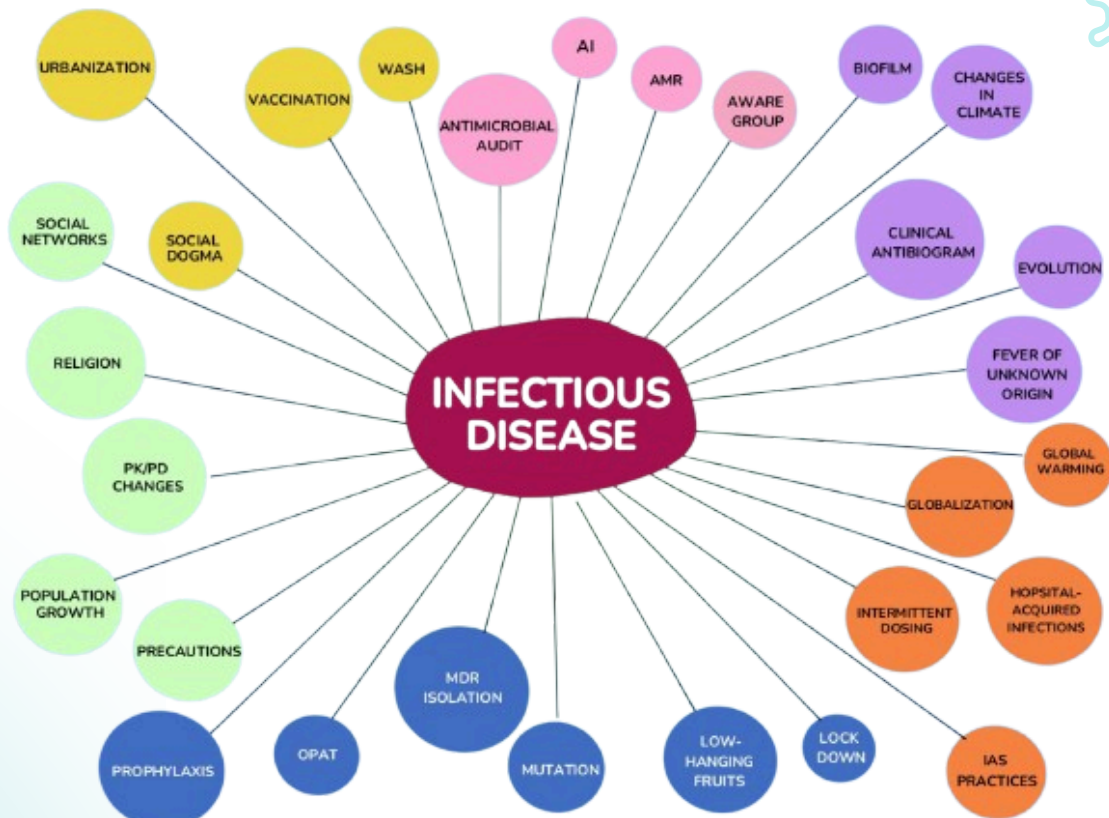
ASPICON 2024 | AUGUST 22 -25, 2024

6th National Conference and Workshop on Antimicrobial Stewardship

PGIMER, Chandigarh

Theme: “Contextualized Approach to Antimicrobial Stewardship”

ABSTRACT BOOK



Publisher

Society of Antimicrobial Stewardship and Practices (SASPI)

Evaluation of Drug Use and Audit of Prescription of Restricted Antibiotics in the Intensive Care Unit of Tertiary Care Hospital

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Background

Antimicrobial agents are the biggest contribution to the 20th century, which are used to prevent and cure infections. Reserve antibiotics are the last-choice of antibiotics to treat multidrug-resistant infections. The inappropriate use & irrational prescriptions of these drugs is a global health problem, and has an ill effect on health as well as on healthcare expenditure. To prevent it, regular and timely prescription audits are an important tool to improve the quality of healthcare.

Aim: To rationalize the overuse/misuse of the reserve drugs in intensive care units and to avoid emergence of drug resistance.

Material & Method

This was a cross-sectional observational study done in Dayanand Medical College and Hospital from March To May 2024 in patients admitted in ICUs. In our hospital Reserve Drug policy six drugs were included: Linezolid, Polymyxin B, Colistin, Fosfomycin, Tigecycline & Ceftazidime-avibactam. The data was collected on reserve drug audit proforma. All the forms were analyzed by a team of Microbiologist, Pharmacologist and clinicians for the appropriate usage of the drugs on the basis of clinical condition, culture report and other investigations of the patient.

Result

Among the 272 prescriptions analyzed, 63.6% patients were males. Maximum number of reserve drugs were given to the patients belonging to the age group 61-80 years (41.5%). Audit report shows that most of the

reserve drugs were used in Medical wards & Colistin (124) and Tigecycline (91) were the most prescribed drugs. Single Restricted antimicrobial therapy was given in 66.5% patients, dual therapy in 25% and three drugs were given in 8.1%. Out of 272 Prescriptions of Restricted Antibiotics, 70.9% were found to be rational.

Conclusion

We can reduce the evolution of antibiotic resistance by its judicious use and education. So an audit is a mandatory exercise to provide quality care in the healthcare system.

Investigation of a Cluster of *Clostridioides Difficile* Diarrhea in a Medical Oncology Unit

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Background

The medical oncology unit of our tertiary care hospital experienced an increase in the number of *Clostridioides difficile* (*C. difficile*) infections from March 2024 to June 2024. This study aimed to investigate the rise in *C. difficile* cases, identify risk factors, and implement measures to contain and prevent future occurrences.

Methods

A comprehensive investigation involved collection from November 2022 to July 2024. The investigation included sample testing, analysis of infection trends across departments, and identification of risk factors such as outside hospitalization, multiple hospitalizations, prolonged hospital stays, antibiotic usage, infection control practices, and cleaning & disinfection protocols for environmental disinfection. Measures taken included strengthening hand hygiene, increasing the frequency of environmental cleaning, and introducing hydrogen peroxide fogging. Antibiotic stewardship programs were also reinforced, and active surveillance for new cases was implemented.

Results

The analysis revealed 34 positive cases out of *C. difficile* of 151 tested samples, with a 22.5% positivity rate. The majority of cases were concentrated in the Medical Oncology department (52%). Key risk factors identified included prior hospitalization (48%), multiple hospitalizations (72%), prolonged hospitalization (16%), and the use of high-risk antibiotics (76%). Environmental persistence of *C. difficile* spores and inadequate cleaning practices contributed to the outbreak. The implemented measures resulted in improved infection control, reduced antibiotic usage, and enhanced communication among staff.

Conclusion

The investigation highlighted critical risk factors and provided a basis for targeted interventions. Strengthening hand hygiene, optimizing environmental cleaning, and reinforcing antibiotic stewardship were effective in reducing the number of cases. Continued surveillance, staff education, and adherence to infection control protocols are essential to prevent future *C. difficile* in healthcare. The study underscores the importance of a multifaceted approach in managing healthcare-associated infections and enhancing patient safety.

Changing Trends of Antibiotic Susceptibility Pattern in *Escherichia coli* Isolates Obtained from Urinary Tract Infections in a Tertiary Care Hospital in Sub-Himalayan Belt

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Background

Escherichia coli is the most common organism causing both communities as well as hospital-acquired UTIs. The effectiveness of antibiotics used to treat these infections is threatened by growing bacterial resistance.

Aim & Objectives: This study aims to analyze the changing trends in the antibiotic susceptibility pattern of *Escherichia coli* isolates obtained from UTI cases in the last three years.

Methodology

Retrospective data analysis of antibiotic susceptibility pattern in *Escherichia coli* obtained between Jan 2021 -Dec 2023 in the Department of Microbiology.

Results

A total of 22,802 urine samples were tested over three years. In 2021, a total of 457 isolates of *E.coli* were obtained followed by 1072 in 2022 and 977 in 2023.. Maximum isolates were obtained in the age group of 18-45 years. Amongst the most commonly used urinary antibiotics, Nitrofurantoin showed a declining trend in susceptibility ranging from 93%in 2021 to 76% in 2022 and 51% in 2023. Fosfomycin showed steady susceptibility rates in 2021 (96%), 2022 (96%), and 2023 (94%). Ciprofloxacin showed a decrease in susceptibility rangingfrom40%in 2021 and 32% in 2023. Piperacillin showed a sharp decline in susceptibility rates from 84% in 2021,74% in 2022 and 20% in 2023. Ampicillin showed a decrease in susceptibility from 30% in 2021 to 25% in 2023.

Conclusion

An increasing trend in resistance to common urinary antibiotics like nitrofurantoin, Cephalosporins, and Fluoroquinolones emphasizes the need to strengthen hospital infection control practices and judicious use of antibiotics for empirical therapy according to standard guidelines. This can further help in the effective implementation of AMSP guidelines, guiding empirical treatment, and formulating hospital antibiotic policy according to local antibiograms.

Barriers and Facilitators of Outpatient Parenteral Antimicrobial Therapy Practice and its Role in Antimicrobial Stewardship – A Pilot Longitudinal Study

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Introduction

Outpatient parenteral antimicrobial therapy (OPAT) offers a crucial method for administering IV/IM antimicrobials outside hospitals, enabling patients to complete treatment safely outside and many hospital-acquired events. This pilot study evaluates real-world barriers and facilitators of OPAT's practice in resource-poor settings, emphasizing its role in antimicrobial stewardship.

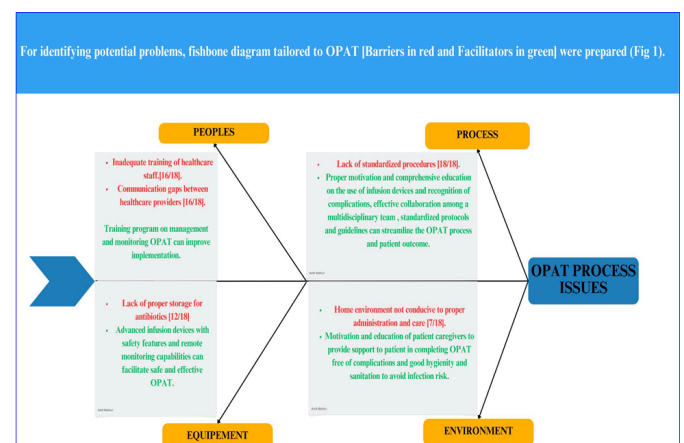
Methodology

This pilot longitudinal observational study included patients meeting OPAT checklist criteria and committed to post-discharge follow-up. Pre-discharge demographic data noted. Various barriers and facilitators were identified after an extensive literature review by preparing a fishbone diagram, data collection and analysis, and patient feedback.

Results

The mean age of patients was 38 years [Range=21-63], with Male-14 and Female-4, with all OPAT administered at the home by family members on 7/18, and by a local nurse on 11/18. Infections requiring OPAT were Typhoid-5, UTI-4, HAP-4, Meningitis-3, IE-1 and multiple visceral abscess-1. 17/18 patients became afebrile and one instance of prematurely discontinued OPAT regimens and one readmission was observed. Additionally, one patient developed thrombophlebitis. No education, counseling, or demonstrations were done before discharge in almost half of the patients. It saves 2 week's duration of hospitalization For identifying potential problems, a fishbone diagram

tailored to OPAT [Barriers (red) and Facilitators (green)] was prepared (Fig 1).



Conclusion

This pilot study provides a comprehensive understanding of barriers and facilitators influencing OPAT practices in a resource-poor setting. By addressing these barriers and leveraging facilitators, each healthcare facility can optimize OPAT delivery, improve patient outcomes, and ensure a more efficient and patient-centered approach to antimicrobial therapy. OPAT reduces hospital stays and hospital-acquired infections, is crucial in combating antimicrobial resistance (AMR), and thus directly contributes to integrated antimicrobial stewardship, aligning with a global action plan for AMR in infection prevention and antimicrobial optimal utilization.

Development and Characterization of *A. Annua* L. Whole Leaf Extract Loaded HSA Nanoparticles Using QbD Approach: A Novel and Potential Antimalarial Formulation

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Background

Malaria, caused by *Plasmodium* protozoa, remains a major global health issue. Significant efforts have been made to eradicate malaria, with chemotherapy being a key strategy in combating the disease. The most significant antimalarial drugs are artemisinin (ART) and its semi-synthetic derivatives, which have been used in combination therapy (ACTs) for improved efficacy. However, despite early success, ACTs have proven unsuccessful in many endemic nations due to drug resistance. The development of nano drug delivery systems (NDDSs) is critical to overcoming this challenge and improving treatment outcomes.

Methods

This study aimed to design, optimize, and develop *Artemisia annua* L. whole leaf extract-loaded human serum albumin (HSA) nanoparticles using a quality-based design (QbD) and risk assessment methodology. The nanoparticles were synthesized via the desolvation method and characterized by dynamic light scattering (DLS), transmission electron microscopy (TEM), scanning electron microscopy (SEM), and Fourier-transform infrared spectroscopy (FT-IR).

Results

The optimized nanoparticles exhibited a particle size of 84.73 nm, a polydispersity index (PDI) of 0.23, a zeta potential of -22.11 mV, and an entrapment efficiency exceeding 90%. Additionally, the ART release profile

demonstrated sustained release and stability studies indicated long-term stability.

Conclusion

The *A. annua* L. whole leaf extract-loaded HSA nanoparticles developed in this study present a promising antimalarial drug delivery system. The developed nanoparticles may have the potential to improve treatment outcomes and overcome drug resistance, thereby addressing a critical need in the fight against malaria.

Implementing Antimicrobial Prescription Checklist to Improve Antimicrobial Prescription Quality – A Pilot Project Towards Antimicrobial Stewardship

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Background

Inappropriate use of antimicrobials is a major cause of antimicrobial resistance (AMR). Antimicrobial Stewardship (AMS) interventions are required to curtail the AMR. The antimicrobial prescription of an indoor patient is an important target for AMS intervention. Enhancing the quality of antimicrobial prescriptions will help ensure that antimicrobials are used appropriately.

Objective: To assess the improvement in the quality of antimicrobial prescription against the 'Antimicrobial Prescription Checklist' and the effect of improved antimicrobial prescription practices on various clinical outcomes.

Methodology

Active surveillance was conducted in a randomly selected ward of the General Medicine department for two months. Prescription data was recorded daily using a pre-validated Data Abstraction form. Patients who received antimicrobial agents via systemic route were included for analysis. Patient days were counted daily at 6 pm for calculating days of therapy (DOT) per 1000 patient-days (PD). The 'Antimicrobial Prescription Checklist (APC)' was used to ensure compliance with antimicrobial prescriptions. Data was collected before and after the intervention. The educational intervention for antimicrobial stewardship included didactic lectures, small-group discussions, WhatsApp notifications, and posters in the ward. Data was analyzed for APC criteria indicators, clinical outcomes like DOT per 1000 PD, WHO prescribing indicators, antimicrobial therapy duration, and length of stay.

Results

The educational intervention received positive and immediate feedback from the prescribers. A slight improvement in a few APC criteria, including the notation of 'review after 48 hours', the day of antimicrobial prescribed, and the report of culture sensitivity was observed. The average number of antimicrobials prescribed per hospitalization (2.36 vs 1.9) and the average duration of antimicrobial treatment days (12.53 vs 8.1) were decreased. A decreased consumption (per 1000 PD) of 'Watch' group antimicrobials (Meropenem: 171 vs 41; Piperacillin-tazobactam: 264 vs 192; Vancomycin: 90 vs 12) was observed. The median length of antimicrobial therapy (629 vs 383 days) and hospital stays [10 (IQR: 7 – 18) vs 9 (IQR: 6 - 13) days] was decreased.

Conclusion

The APC Criteria can improve prescription quality and identify key antimicrobial stewardship areas. The educational intervention improved clinical outcomes. However, sustainable and long-term observations are needed to draw a meaningful conclusion.

Phenotypic and Genotypic Characterization of Carbapenem-Resistant *Enterobacterales* Obtained from Various Clinical Specimens

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Background

The rise of Carbapenem-resistant *Enterobacterales* (CRE) presents a significant threat to global health, particularly due to the production of carbapenemase enzymes, including New Delhi Metallo- β -lactamase (NDM-1).

Methodology

The study investigates the prevalence and phenotypic detection of carbapenemase-producing Enterobacteriaceae (CRE) in clinical specimens, emphasizing the distribution of the NDM-1 gene among carbapenem-resistant isolates.

Result

A total of 17,984 clinical specimens were analyzed, with 1,843 culture-positive samples revealing Gram-positive cocci (GPC) in 632 cases, Gram-negative bacilli (GNB) in 1,207 cases, and *Candida albicans* in 4 cases. Among Enterobacterales, *Escherichia coli* was the most common isolate (666), followed by *Klebsiella* species (172), *Citrobacter* species (93), and others. Phenotypic detection of carbapenemase production was done by employing modified Carba NP, Rapidec Carba NP, EDTA combined disc assay, modified Hodge test, and modified carbapenem inactivation assay. The study found significant variability in detection rates among these methods. The Modified Carba NP and Rapidec Carba NP tests showed a higher sensitivity for detecting carbapenemase production compared to other methods. Among the 150 CRE isolates, 68 were identified as NDM-1 gene carriers, predominantly in *E. coli* (33) and *Klebsiella* species (29).

Conclusion

The findings underscore the need for comprehensive phenotypic and genotypic testing to accurately identify and manage CRE infections, thereby aiding in the implementation of effective infection control measures.

Real-time Assessment of the Effectiveness of Alcohol-based Hand Rubs in Disinfecting Non-critical Patient Care Items in Tertiary Care Hospital Settings

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Introduction

Non-critical patient care items are frequently implicated in the transmission of Multi-drug resistant organisms (MDROs), resulting in hospital outbreaks. Alcohols are the most preferred agents for disinfection of these items due to their rapid action and compatibility with the majority of hard surfaces. Many hospitals use 'alcohol-based hand rubs' (ABHR) for this purpose due to their easy availability, even though they are designed for skin antisepsis and not hard surface disinfection. This study assessed the effectiveness of ABHR in disinfecting them, in real-time situations, as little is known about it.

Methodology

A cross-sectional study for two months was conducted in the inpatient facility of a 1,000-bed tertiary care hospital. Approximately 210 patient care items were sampled from various ICU and non-ICU locations. Two samples from each equipment were collected, one before and one after disinfection by ABHR, while the equipment was in real-time use. The collected samples were processed for aerobic colony count. Mean CFU/ml of bacteria grown from pre and post-disinfection sampling was compared to determine the efficacy of the disinfection procedure.

Results

The baseline contamination rate for the items was observed to be 72.4% (152/210) which was higher in non-ICU locations than in ICUs. Most of the items (96.1%) were contaminated with Coagulase Negative Staphylococcus. The effectiveness of ABHR in decontaminating patient care items was found to be

93.1% $\pm$ 6.3. ABHR was also found to be effective in decontaminating objects with an MDRO microbial profile; however, objects contaminated with Staphylococcus aureus were the least effectively removed by it.

Conclusion

ABHR was found effective for the disinfection of non-critical patient care items in real-time scenarios including its action on MDROs. Hence, it can be safely used to achieve the disinfection of hard surfaces in addition to its recommended use.

A Cross-sectional, Qualitative, and Observational Study was Conducted in the OPD Setting of a Rural and an Urban Setting Community Hospital

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Introduction

The overuse and misuse of antimicrobials in human health with other contributing factors like inappropriate prescriptions, prolonged use of antimicrobials, and sometimes the presence of bacteria in the environment and entry of these antimicrobials to the food chain through any means have increased the development and spread of AMR globally.

Methodology

This study is a cross-sectional, prospective, and observational study conducted in an outpatient department (OPD) of two community health settings (rural and urban). The main objective of the study is to evaluate the antibiotic prescription pattern in an OPD setting. Using a random sampling procedure, 1724 prescriptions were studied from an urban setting and 1151 from a rural setting.

Results

Some of the WHO core indicators were studied and it was seen that 98% of the prescriptions were from the essential drug list (EDL) in a rural setting and 87% were from an urban setting. 99% of the prescriptions were based on generic names in an urban setting and rural areas it was around 91%. 15.25% of prescriptions from urban settings were based on diagnosis. The most prescribed antibiotic from an urban setting was Cefixime, whereas, from a rural setting, it was Amoxycylav. Most of the cases reported from this setting were from general medicine OPD, whereas in rural settings it was from dental OPD where the antibiotics were prescribed just in case.

Conclusion

We see high use of antibiotics from the Watch group being prescribed in urban settings, whereas in the rural setting, it is from the Access group. But on the positive side, we saw most of the prescriptions in both settings were made from the essential drug list and most of them were written by generic names following the WHO guidelines. Making these prescription patterns available will help clinicians in studying the current practicing scenario.

A Study to Evaluate the Use of Antimicrobial Agents in Patients of Burns at a Tertiary Care Hospital.

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Introduction

Burns are the most painful of all injuries and a major global public health problem. Mortality due to sepsis is a major concern which can be prevented with proper antimicrobial agents according to culture sensitivity.

Methodology

This prospective, observational study was conducted on 104 adults who were having burns and were admitted at Civil Hospital, Ahmedabad. After approval from the Institutional Ethics Committee, each patient was followed up daily for the first seven days and then weekly till discharge. Demographic details, vitals, local wound examination, numeric pain rating scale, culture sensitivity report, drug treatment, and adverse events were recorded and analyzed using appropriate statistical tests.

Results

A total of 104 adult burn patients were enrolled. Male to female ratio was 1:1.89. At the time of admission the mean total body surface area (TBSA) burned was 24 + 16%. The most common type of burn was thermal burns(45.19%). Patients who suffered from severe full-thickness burns were 73(70.19%). The most commonly used topical antimicrobial was Silver sulfadiazine (100%) while Systemic antibiotics like amoxicillin + clavulanic acid (99%) was used empirically and prophylactically in 82 and 22 patients respectively. In addition to these Piperacillin tazobactam(18.27%), Linezolid(35.58%), Metronidazole(14.42%) were used as add on therapy. Culture sensitivity is done in 23(22.11%) patients among them 7 (30.43%) *Pseudomonas aeruginosa*, 4(17.39%) *Acinetobacter baumannii*, and 4(17.39%) staphylococcus aureus isolated. 64 patients were discharged with healing wounds. The total reported

adverse drug reactions were 31 among them 15(48.39%) were constipation. The mean length of hospital stay was 21 + 15 days. Overall mortality observed due to sepsis is 7(33%).

Conclusion

The extent of the burn area, days of hospital stay, and nature of the burn are the risk factors for the development of infection. Antimicrobials based on culture sensitivity reports can prevent infection-related mortality. Public awareness about burns prevention is also advisable.

A Point Prevalence Survey Study (PPS) of Antimicrobial Consumption in a Tertiary Care Super-speciality Hospital of West Bengal from 2021: Some Interesting Findings

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Background

The burden of antimicrobial resistance (AMR) is an emerging global health issue. Resistance occurs when bacteria, parasites, viruses, or fungi are exposed to antimicrobials but not killed by them. The study was conducted in the year 2021 to gather baseline information and to assess the antimicrobial consumption practices across six departments in a tertiary care super specialty hospital in West Bengal.

Methodology

The institutional ethics committee accorded the study protocol. A modified version of a patient data collection form proposed by Global PPS was developed on Epi Info software version 7 (CDC). Data of all patients admitted to a ward at 08.00 am data were studied. The use of antimicrobials was categorized as empiric, prophylactic, or lab-based. WHO-AWARE classification of antimicrobials was instituted for categorizing usage.

Results

A total of 85 patient-related data was collected in the designated survey form. The total number of beds covered was 340 and the number of patients on antimicrobials was found to be low at 21.27 %. The patients surveyed were predominantly female (78.8 %). The mean number of antimicrobials per patient was found to be 1.62 (range of 1.4 to 2.2). Relatively low number of patients were found to be on 2 or more antimicrobials. Double gram-negative and Double anaerobic coverage of AM used varied across departments covered in the survey.

Conclusion

Our point prevalence study was able to facilitate the conducting of a point prevalence survey in a high patient volume tertiary care hospital with a paper-based medical record system and depicted the baseline parameters of intervention for instituting future action and policy changes.

Normal Cytology Doesn't Preclude High-risk HPV Infection

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Abstract

Human papillomavirus infection is the major risk factor for cervical cancer. Infection with certain high-risk HPVs makes an individual more prone to develop cancer. Therefore, understanding the prevalence and distribution of HPV genotypes is necessary to implement targeted prevention strategies. Hence, in this study, we evaluated the burden and distribution of HPV genotypes among women across the cervical cytological changes. 125 symptomatic women were recruited in the following three groups; women with a) cervical cancer, b) squamous intraepithelial lesions, and c) normal cytology based on the Pap test. We evaluated the presence of HPV and its genotypes using RT-PCR with a 16/18 genotyping kit. The overall prevalence of HPV infection was observed to be (84/124) 67.2%. Prevalence of HPV infection in cervical cancer cases was highest with (48/50) 96% positivity followed by 56% and 44% among women with squamous intraepithelial lesions and normal cytology respectively. HPV 16 emerged as the predominant genotype, accounting for 59.60% of cases. Following closely, HPV 18 exhibited 8.33% positivity. HPV 8 GTS genotype demonstrated 17.85% positivity, suggesting the presence of any of the following genotypes: 31, 33, 35, 51, 52, 56, 58, or 66. Multiple genotypes coexisted in 4.76% of patients. Conclusively, our finding indicates HPV 16 to be the most common genotype followed by HPV 8 GTS. This study reflects the need to screen symptomatic women for HPV even if their pap test suggests normal cytology for early identification and monitoring of women infected with high-risk HPV.

Cognizance on Antimicrobial Stewardship Indicators in Intensive Care Unit

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Background

Antimicrobial resistance (AMR), is recognized as one of the top ten public health threats by the World Health Organization. About 4.95 million deaths were associated with bacterial AMR globally. Fighting antimicrobial resistance requires appropriate diagnosis and monitoring of the use of antibiotics through an effective Antimicrobial stewardship program which assists in monitoring antibiotic resistance trends, antimicrobial consumption, and clinical outcomes of patients in healthcare facilities.

Methodology

ICU WORKUP: Regular visits were made to the ICU and the patient's relevant data were collected including Dose, duration, blood culture practices, and course of antimicrobials given.

LAB WORKUP: Direct smear from the flagged blood culture was processed further along with Antibiotic susceptibility testing and the outcome and process indicators of AMS were calculated accordingly.

Results

The demographic parameters of the patients revealed that the number of males admitted to the ICU was almost equal to that of females and the median age of patients was around 46.5 years. The mortality rate in this study was found to be 36.5 %. In our study, we analyzed the utilization patterns of Antimicrobial agents in terms of DDD/100 patient days and DOT/100 patient days. The median DOT/100 patient days was 7 (0–29). The most commonly used antibiotics are Meropenem, Piperacillin-tazobactam, Ceftriaxone, and Azithromycin with DDDs 1.45, 3, 6, and 3 respectively. Appropriate timing of Blood culture collection was

followed in 70% of the cases, 30% of cases were already started on empirical therapy before sending samples for culture. The infection-specific mortality was found to be 12.1% while the infection-specific length of stay was increased in 3 patients by 10 days and antibiotic-associated diarrhea was not seen during the study period.

Conclusion

In conclusion, our study reveals that antimicrobials are widely prescribed in critically ill patients which forms a significant proportion of the total drugs consumed in the ICU. Decision-making by the clinicians can be optimized which will result in good treatment compliance since the patients will be exposed to fewer and more appropriate antibiotics.

Antimicrobial Stewardship: A Tertiary Care Experience

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Background

Antimicrobial resistance (AMR) is a global health concern. An antimicrobial stewardship program is the core strategy to tackle AMR. The goals of this program are to rationalize prescribing and avoid unnecessary use of antimicrobials. It plays a significant role in decreasing the incidence of resistance, reducing the development of multidrug-resistant organisms, and improving patient care with the long-term goal of preserving antimicrobials.

Methodology

The antimicrobial stewardship program was started in 2019 at our tertiary care center. This ongoing program includes microbiology, the infection control department, infectious disease physicians, and clinical pharmacists. The data is collected on data forms which are then collated on Microsoft Excel. The team provides stewardship advice based on the hospital antibiotic policy and the results of the culture.

Results

The total no. of patients admitted to the hospital for the period 2022-2023 was 85833. Of these, the percentage of patients administered antimicrobials varied from 8.1% to 25.5% per month with an average of 16%. Compliance with stewardship advice was around 78%. The percentage of compliance with de-escalation advice was 74.4%. The combined Duration of Therapy for all antimicrobials per 1000 patient days was 474. DOT per 1000 patient days for selected antimicrobials was as follows: Piperacillin-tazobactam (53.7), Carbapenems (49.5), Polymyxins (13.1), First-generation Cephalosporins (31.5), Third-generation Cephalosporins (78.5), Vancomycin (9.8) and

Fluoroquinolones (6.4). Consumption of selected antimicrobials in DDD per 1000 patient days was as follows: First-generation Cephalosporins (32.6), Third generation Cephalosporins (72.3), Piperacillin-tazobactam (40) Carbapenem (30) Polymyxin (3), Vancomycin (4.7). The Length of Therapy of antimicrobials per 1000 patient days has ranged from 146 to 322 days with an average of 220 days. The average length of stay of the patients on antimicrobials was 9.3 days. The average cost per patient per day was Rs. 485.

Conclusion

The implementation of the Antimicrobial Stewardship Program at our center has resulted in the rationalization of antibiotic prescriptions as compared to other institutions of a similar nature.

Drug Resistance Profile of TB Patients Attending a Tertiary Care Centre in the Northern Region of India

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Background

The emergence of multidrug-resistant MDR-TB and extensively drug-resistant XDR-TB are serious threats to global TB control. Molecular tests like GenoType MTBDRplus have revolutionized MDR-TB diagnosis by rapid detection of resistance, leading to early and appropriate management of DR-TB. Retrospective data was collected to determine the study drug-resistant pattern among TB patients.

reducing turnaround time and leading to early management of DR-TB cases.

Methodology

A retrospective analysis was done over 12 months from January 2023 to December 2023 to study the resistance pattern and mutations present in DR-TB in adults with suspected pulmonary or extrapulmonary tuberculosis using Hain's GenoType MTBDRplus VER 2.0.

Results

Over 12 months, samples from 6425 patients with suspected TB were received, 822 out of these 6,425 isolated were tested for drug resistance by Hain's GenoType MTBDRplus VER 2.0. 11.68% (751) were sensitive to isoniazid and rifampicin while 8.63% were DR-TB (71 out of 822). Out of these 71, 20 were MDR TB, 48 were isoniazid monoresistant (0.74%) and 3 were rifampicin monoresistant. Out of 3,726 males, 511 were tested for drug resistance; 9.00% (46 out of 511) were resistant, and out of 2,633 females; 314 were tested for drug resistance; 9.23% were resistant (29 out of 314).

Conclusion

We cannot solely rely on Xpert MTB/Rif Assay for detection of drug resistance due to the risk of missing

the isoniazid monoresistance. GenoType MTBDR plus has revolutionized MDR-TB diagnosis by substantially

Abstract 016

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Targeted Pharmacovigilance for Metronidazole-Induce Neuropathy in Patients of Amoebic Liver Abscess at A Tertiary Care Hospital: An Ambispective Study

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Introduction

Metronidazole has been the front-line choice for several anaerobic and protozoan infections worldwide (1). Metronidazole is widely prescribed and recognized for its effectiveness in treating various infections, including amoebic liver abscesses (ALA). However, its use comes with potential complications, particularly the development of neuropathy, which affects the nerves and can cause significant discomfort and impairment for patients (2). While existing knowledge suggests that neuropathy is a rare side effect, the exact incidence rates and risk factors remain areas of ongoing research. Our study addresses the neuropathy risk associated with metronidazole treatment for ALA. To assess the incidence and severity of metronidazole-induced neuropathy in patients with ALA managed with Metronidazole.

Methodology

It was an observational study conducted in the Department of General Medicine, General Surgery, Gastro-Enterology, Paediatric Surgery, and Pharmacology, All India Institute of Medical Sciences, Rishikesh, UK, India. The duration of the study was six months. All patients diagnosed with ALA managed with Metronidazole were included (Adult and pediatric above 5 years and patients from both genders). The suspected adverse drug reaction reporting form,

version 1.4, was used to collect data. All the suspected cases of adverse events were reported to the ADR Monitoring Centre, AIIMS, Rishikesh. The causal relationship was established by the standardized WHO UMC Causality Assessment. The analyzed data was then uploaded to Vigiflow software and sent to NCC. The severity and preventability of metronidazole-induced neuropathy were assessed using the Modified Hartwig severity assessment scale and Modified Schumock-Thornton Scale, respectively. Retrospectively, adverse drug reaction (ADR) data were collected from Vigiflow software at AMC, AIIMS, and Rishikesh and cross-checked with the respective department, treating physicians, and Patients (telephonically) for further information.

Results

A total of 165 patients were recruited. 23 developed neuropathies, primarily due to long-term medication use and risk factors like alcohol, tobacco, and diabetes. Most affected were males (69%) within the 20-50 age range. The occurrence of neuropathy symptoms was most frequently reported with the 800 mg dose. The most common duration of Metronidazole use was 30 days. 26% of patients suffered from severe symptoms of metronidazole-induced neuropathy.

Conclusion

This study identified the percentage of patients developing neuropathy after metronidazole use in ALA, including gender, age, severity, dose, and duration of treatment. Awareness of the drug's potential for neurotoxic effects is crucial, and its use should be

judiciously managed to prevent unnecessary exposure. These data will be helpful for the risk minimization and risk management plans.

Abstract 017

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A Comparative Study to Assess Sputum and Bronchoalveolar Lavage (BAL) Sample in Infective Lung Diseases

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Introduction

Establishing the causative pathogen early in infective lung diseases is cardinal for reducing diagnostic delay and ensuring the administration of effective antibiotics, especially in an era of growing antibiotic resistance, and will decrease mortality and morbidity.

Aims & Objectives: This study aims to analyze concordance or discordance among sputum and BAL samples.

Methodology

This retrospective study was conducted in a tertiary hospital in North India from January 2022 to December 2022 and a total of 3758 sputum samples and 253 BAL samples were received. Samples were directly inoculated on Blood agar, Chocolate agar, and MacConkey agar. Confirmation of isolates and antimicrobial susceptibility testing was done with the help of an automated system (VITEK 2@bioMerieux). These samples were analyzed for concordance amongst 56 samples for which both BAL and sputum were sent in close succession to rule out other infection incidence.

Results

Concordance among various microorganisms isolated from BAL and Sputum samples was observed in about 80 % of cases which indicates that sputum samples may be used in cases where BAL samples cannot be taken.

Conclusion

Sputum analysis is a more cost-effective and noninvasive sampling method. This study can be further used prospectively with clinicians to de-escalate antibiotics if started, and/or to change antibiotic therapy according to the sputum sample culture results.

Phage Selection Can Restore Antibiotic Sensitivity to Treat Urinary Tract Infections: Preclinical Study Findings

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Background

Phage-antibiotic synergy (PAS) has emerged in recent years to possibly replace or supplement traditional antibiotic therapy. Here, we report a unique mechanism involved in the effectiveness of phage PGN80A and antibiotics combination against multidrug-resistant uropathogenic *Escherichia coli* isolates.

Material & Methodology

To unravel the pharmacological dynamics, we employed whole genome sequencing for wild-type isolates and phage PGN80A-resistant variants (n=10) using NovaSeq6000. To determine the PAS in phage PGN80A (102 -109 PFU/mL) five different classes of antibiotics with varied concentrations (1–512 µg/ml) were co-cultured with bacterial culture. To examine the role of cell filamentation in synergism, we examined the morphology of wild and mutant bacterial isolates using scanning electron microscopy. A transurethral catheter mice model was used to test and compare the efficacy of phage and antibiotic synergy.

Results

In our pharmacogenomic analysis, we revealed that phage-resistant variants acquired multiple single nucleotides (n=3868) and frameshift mutations

(n=184). Cells were significantly shorter for bacteria grown in a Trypticase soy broth without imipenem compared to those grown in a medium with the sub-inhibitory concentration of imipenem and low titer of phage PGN80A. Phage 80A demonstrated a pharmacologically relevant synergistic effect in combination with imipenem at 1/100 of the MIC concentration. A significant reduction ($p<0.01$) in bacterial load was observed in mice treated with phage PGN80A (106 PFU/ml) and a low dose (25 mg/kg) of imipenem.

Conclusion

Our study undertakes a pharmacologically oriented comprehensive analysis of phage-antibiotic interactions for scrutinizing synergy, additivity, and antagonism. We observed morphological changes after antibiotic application, the occurrence of multiple mutations in phage-resistant variants indicating the involvement of multi-antibacterial strategies. We conclude that phages, such as PGN80A, represent a new approach to traditional therapeutics where phages exert selection for MDR bacteria to become increasingly sensitive to conventional antibiotics.

Evaluation of Antimicrobial Drug Promotional Literature: A Critical Analysis Using Diverse Standard Tools

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Background

Drug promotional literature (DPLs) are known to influence the drug prescribing practices of physicians. Previous studies in multiple therapeutic areas have often shown that the DPLs lack fair balance, manipulate data, and make unsubstantiated claims. There is a paucity of research on critical appraisal of antimicrobial DPLs in particular. This could lead to irrational prescribing patterns of antibiotics thereby exacerbating antimicrobial resistance. Therefore, the current study is planned to assess the compliance of antimicrobial DPLs to various standard guidelines.

Material & Methodology

A cross-sectional observational study was conducted over three months to analyze antimicrobial DPLs. These were collected from multiple sources including various clinical departments of our hospital, medical representatives, and nearby pharmacy outlets. Institute Ethics Committee approval was not sought as the utilized information was available in the public domain. DPLs were evaluated using multiple criteria such as WHO Ethical Criteria for Medicinal Drug Promotion, International Federation of Pharmaceutical Manufacturers & Associations (IFPMA) Code of Practice, Uniform Code for Pharmaceuticals Marketing Practices (UCPMP), Organization of Pharmaceutical Producers of India (OPPI) and Association of British

Pharmaceutical Industry (ABPI) code of practice. In addition, the “Fair balance” of DPL information was assessed using USFDA criteria. Data were entered in a Microsoft Excel sheet and analyzed using descriptive statistics.

Results

The majority of DPLs did not comply with the standard guidelines completely and lacked a fair balance of the information. Reference to scientific literature substantiating claims was not provided in most of the studied DPLs.

Conclusion

The study provided valuable insights into the quality of antimicrobial DPLs. Regulatory measures are needed to ensure healthcare professionals receive accurate, current, unbiased evidence-based information on antimicrobials for rational prescribing.

Abstract 020

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Linezolid Resistant Enterococci in Urinary Tract Infection

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Background

Enterococci are one of the most common isolates in urinary tract infections. It shows intrinsic as well as acquired resistance to a wide range of antibiotics. With the ascending trend of Vancomycin-resistant Enterococci in nosocomial infections, the use of Linezolid has proportionately increased because of its high oral bioavailability and good clinical cure rate. However, cases of Linezolid resistance have also come up in certain strains of Enterococcus species which can be attributed to nosocomial transmission and extensive usage. The study aimed to isolate and identify Linezolid-resistant Enterococcus species among urine samples and study the antimicrobial susceptibility pattern of Enterococcus species isolates at a tertiary care center.

Material & Methodology

An observational study was carried out for a period of 4 months from January 2024 to April 2024 at the Department of Microbiology, SMS Medical College, Jaipur. A total of 478 isolates of non-duplicated Enterococcus species from urine samples received at the Bacteriology Laboratory were processed. Isolation and identification were done using standard microbiological protocols. Antimicrobial susceptibility testing was done on Mueller-Hinton agar by the Kirby-Bauer disc diffusion method as per CLSI guidelines.

Results

Out of 802 urine samples showing positive aerobic growth, 478(59.6%) were Enterococcus species. Among these isolates, 48(10.04%) VRE (Vancomycin-Resistant Enterococci) and 8 (1.67%) Linezolid Resistant Enterococci were detected, all of which were inpatients. Among the Enterococcus species isolates, maximum resistance was seen towards Ciprofloxacin(85.3%) followed by Tetracycline(77%), High.Gentamycin(67%),..Ampicillin(58.5%), Fosfomycin(42%), Nitrofurantoin(25%), Teicoplanin(13.3%), Vancomycin(10.04%) and Linezolid(1.67%).

Conclusion

The emergence of Linezolid-resistant Enterococci poses an alarming threat to clinicians as these isolates have limited or no therapeutic options. Such a condition requires a reduction of inadvertent use of Linezolid by frequent monitoring of antimicrobial susceptibility patterns and strict infection control measures to contain the spread of these multidrug-resistant pathogens.

Abstract 021

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The Emergence of Carbapenem-Resistant Enterobacteriaceae (CRE) in Southern Haryana: A Hospital-Based Study

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Background

In the past two decades increasing number of bacteria in the family Enterobacteriaceae have become resistant to multiple antimicrobial classes leading clinicians to turn to agents from the broad-spectrum carbapenem class for the effective treatment of life-threatening infections caused by these organisms. Therefore present study planned to detect the presence, type, and estimate the burden thereof carbapenemase-producing Enterobacteriaceae.

Material & Methodology

An observational cross-sectional study was carried out from December 2022 to November 2023(1 year) in the department of microbiology of SHKM GMC Nalhar in Haryana. 94 consecutive carbapenem nonsusceptible (by Kirby Bauer Disc diffusion test) Enterobacterales isolated from various clinical samples were included in the study. Demographic details of patients and samples were included. Identification of isolates to the species level was done using conventional methods. Antimicrobial susceptibility testing of all the isolates was done by Kirby Bauer disc diffusion as per CLSI guidelines. All the carbapenem nonsusceptible isolates by Kirby Bauer disc diffusion method were further subjected to carbapenemase detection by Modified carbapenem inactivation method (mCIM) and further

to characterize the type of carbapenemase, EDTA carbapenem inactivation method(eCIM).

Results

570 Enterobacterales were isolated during the study period. 16.49% were carbapenem nonsusceptible by the Kirby Bauer disc diffusion method. These were subjected to mCIM in which 9.5% were found to be positive for carbapenemase production (CRE). 70.4% of Enterobacteriaceae were Metallo-beta-lactamase producers whereas 29.6 % were Serine-type beta-lactamase producers. The majority of isolates were from female and younger age groups. Further most common sample yielding CRE was pus.

Conclusion

Given that carbapenems are the last resort drug for GNBs, there is an urgent need for active antibiotic stewardship in hospitals and communities. Our study underlines the worrisome presence of these CREs in the hospital and hence calls for stringent infection control measures to curb the menace.

Abstract 022

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Trends in Antimicrobial Susceptibility Pattern of Salmonella Species Isolated from Bacteremia Patients at a Tertiary Care Center in Northern India

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Background

Incidence of Multidrug-resistant (MDR) *Salmonella enterica* serovars has been increasing globally over the past 20 years which in turn has led to limited options of treatment.

Aim: The study was done to assess the antimicrobial susceptibility pattern among *Salmonella enterica* serovars causing bacteremia in Northern India. In this observational study, blood samples positive for *Salmonella enterica* serovars.

Methodology

From January 2021 to April 2023 were studied. Species identification was done using MALDI-ToF MS. Serotyping was done using slide agglutination method using *Salmonella* polyvalent antisera O, monovalent O:1, O:2, O:9, O:12, and Vi for confirmation of different serovars. Antimicrobial susceptibility was interpreted as per the CLSI guidelines.

Results

During the study period, 32 *Salmonella enterica* serovars were isolated. *Salmonella enterica* serovar Typhi was the predominant serovar, followed by

Salmonella enterica serovar Paratyphi A. All isolates were susceptible to ceftriaxone, chloramphenicol, co-trimoxazole, and cefotaxime. Pefloxacin showed 100% resistance. Resistance to nalidixic acid was found in 81.2% of isolates. Of the isolates resistant to nalidixic acid, 19(73.08%) isolates were resistant to ciprofloxacin also. Of the 32 isolates, 2 of them were MDR *Salmonella* isolates. Only one patient died due to the infection. The patient was positive for *Salmonella enterica* serovar Paratyphi B.

Conclusion

This changing susceptibility pattern necessitates continuous surveillance of antibiogram of *Salmonella* isolates to rationalize the treatment protocols for invasive salmonellosis and prevent the emergence of resistant strains.

Abstract 023

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A Questionnaire-based Survey on the Knowledge, Attitude, and Practice of Antimicrobial Stewardship among the Faculties and Resident Doctors of a Teaching Tertiary Care Hospital in Punjab, India

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Background

Antimicrobial stewardship is a coordinated program that promotes the appropriate use of antimicrobials thereby improving patient outcomes and decreasing the spread of infections caused by multidrug-resistant organisms. Due to the rise of antimicrobial resistance (AMR), the efficiency of currently used antimicrobial drugs is declining, making it more difficult and expensive to treat illnesses and harder to manage outbreaks. This has emerged as one of the major public health issues globally. A KAP survey is meant to be a representative survey that aims to elicit what is known (knowledge), believed (attitude), and done (practiced) in the context of the topic of interest.

Material & Methodology

The study will be conducted among practitioners (faculty, senior residents, junior residents) of different departments of our hospital after obtaining ethics approval and informed consent. A pre-designed questionnaire will be used after content validation. A validated questionnaire consisting of 30 questions will be shared among potential participants through either

online (Google Forms) or offline modes. The respondents will be given one week to respond. In case there is no response one reminder will be given after 7 days. Existing gaps in the knowledge, attitude, and practice will be identified and presented using descriptive statistics.

Results

current status of knowledge, attitude, and practice among faculties and resident doctors.

Conclusion

These results will be used to design a tailored educational program for our healthcare setup.

The result of this survey will help in comprehending the

Abstract 024

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Comparative Analysis of Standard Treatment Guidelines for Common Infections in India: Identifying Discrepancies and Opportunities for Harmonization

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Background

Antimicrobial resistance (AMR) is a growing global concern. The availability of multiple standard treatment guidelines (STG) can cause ambiguity among prescribing physicians, resulting in either antibiotic underuse or overuse and thus can exacerbate AMR. In the absence of local or institutional antibiogram data, primary care physicians may face challenges in selecting the correct empiric antibiotic regimen, which may lead to potential errors. Thus, this study aims to compare and contrast various STGs and identify discrepancies that may perplex healthcare professionals.

Material & Methodology

A comparative analysis of various STGs of six common infections in India, i.e., UTI, CAP, Enteric Fever, Acute Otitis Media, Pharyngitis, and PID, was conducted. STG from national sources like ICMR (2019,2022), regional treatment guidelines, e.g., PGIMER, Indian Academy of Pediatrics (2022), Lung India, etc. and international guidelines from CDC 2021, Infectious Disease Society

of America (IDSA), CMDT 2024 were included. Ethical approval was not required as the data was available in the public domain. The recommended first- and second-line treatment regimens across these guidelines were compared and any discrepancies in the drug selection, dose, and duration of treatment were identified.

Results

Substantial discrepancies exist among national and

international STG, including variations in drug selection, with some guidelines designating a particular drug as a first-line treatment, while others classify it as second-line or fail to mention it altogether. Furthermore, additional discrepancies were observed in dosing regimens or treatment duration, with some guidelines providing a range of doses while others specified a single standard dose.

Conclusion

Wide variation exists amongst the studied national and international STG, which may cause confusion in

treatment decisions by prescribers and can potentially contribute to AMR. The findings highlight the need for harmonization and streamlining of treatment guidelines to ensure consistency and evidence-based practice in healthcare.

Abstract 025

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Prevalence and Antibiotic Susceptibility Patterns of *Haemophilus* Species Isolated from Lower Respiratory Samples and Blood Samples of Cancer Patients at a Tertiary Care Hospital

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Background

Haemophilus species are pleomorphic, oxidase-positive, fastidious, capsulated gram-negative coccobacilli that cause invasive diseases such as bacteremia and pneumonia. Cancer patients being immunocompromised are more prone to a wide range of infections including *Haemophilus*. CLSI-approved treatment for *H.influenza* includes Ampicillin, Carbapenems, Cephalosporins Fluoroquinolones, and Cotrimoxazole. So, determining their susceptibility pattern is also essential in understanding the antibiotic trend.

Methodology

One-year retrospective analysis of lower respiratory and blood samples of cancer patients was performed in the microbiology department at Tata Memorial Hospital, Mumbai. All samples were processed according to standard microbiological procedures. *Haemophilus* species identified on VITEK and confirmed with factors V and X were included in the study. Antibiotic susceptibility testing was performed

using the Kirby-Bauer disc diffusion method. Biotype determination was done using biochemical reactions.

Results

Haemophilus spp. were isolated mainly from respiratory samples (94.5%) whereas 5.4% were isolated from blood (bacteremia). The majority of the cases were males (64.5%), and 39% were in the age group 50-65 years. The prevalence of *Haemophilus* species isolated from lower respiratory samples and blood samples was 9.2 %. The distribution of *H. influenzae* was 81.81%, *H. parainfluenza* (13.6%), and others (4.5%). In *H. influenzae*, the biotype III was isolated most frequently (67.2%). All isolates were sensitive to Carbapenems and Cephalosporins, followed by Fluoroquinolones (87.7%), Ampicillin (74.6%), and Cotrimoxazole (30%).

Conclusion

Haemophilus spp. Cause significant respiratory tract infections in cancer patients. *H.influenzae* tends to be

the most common species isolated in biotype III and was found to be the predominant one in our population. Isolates were sensitive to carbapenems and cephalosporins.

Abstract 026

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Identification of Novel Inhibitors Against Multi-Drug Resistant Uropathogenic Escherichia coli Using Virtual Screening and Molecular Dynamics Simulation Study

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Background

Escherichia coli (E. coli) is a gram-negative bacterial pathogen that poses a significant clinical and epidemiologic challenge. The selection pressure brought by the insufficient use of antibiotics has resulted in the emergence of multi-drug-resistant E. coli in the past ten years.

Material & Methodology

Computational and bioinformatics methods for screening inhibitors have significantly contributed to discovering novel antibacterial agents. One possible target for novel anti-virulence drugs is motility. Motility inhibitors are generally effective at concentrations lower than those required for the antibacterial properties of traditional antibiotics, and they are likely to exert less selective pressure than current medicines. Motility may be essential for bacteria to survive, find nutrients, and escape unfavorable environments and biofilm formation. The FliN is a protein forming the

bulk of the C ring of the flagella and is present in multiple copies (more than 100) in bacteria. Its absence in mammals makes it an attractive drug target for drug discovery. Two-thousand seven hundred seventy-eight natural compounds from the ZINC library were screened against FliN (PDB ID: 4YXB) using PyRx AutoDock Vina, and the top compounds were selected for secondary screening after sorting the results based on their binding energy.

Results

kcal/mol), and inhibition constant (1.98 μ M), ZINC000000619481 was the best inhibitor. This compound binds exactly as per the defined active site residues of the receptor protein. Also, molecular dynamics was performed. The eigenvalue of the selected complex was 1.241657e-05. There were no ADME properties outside of the specified range for the identified hit; it fitted exactly to the binding site of the FliN receptor well and was found to be stable in MD simulation studies.

Conclusion

Further in vitro and in vivo studies are needed to confirm its antibacterial activity and use as a potential antimicrobial drug against urinary tract infections caused by *E. coli*.

Based on interactional analysis, binding energy (– 7.78

Abstract 027

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Comparative In Vitro Efficacy of Generic and Branded Levofloxacin, Ceftriaxone and Meropenem Injections

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Background

The broad acceptance of generic drugs encounters challenges related to their quality, effectiveness, and regulatory scrutiny, leading to significant bias against their substitution. Evaluating the therapeutic effectiveness of both branded and generic antibiotics becomes essential as antibiotic resistance grows. The WHO estimates that a considerable portion of circulating antibiotics, particularly in developing nations, is sub-standard. These substandard antibiotics can result in treatment failure and prolonged illness, significantly contributing to the global antimicrobial resistance (AMR) issue.

Aim & Objective: To assess and compare the In Vitro efficacy of Generic and Branded Injections of Levofloxacin, Ceftriaxone, and Meropenem

Methodology

One generic and one branded formulation of Levofloxacin, Ceftriaxone, and Meropenem were selected for microbiological analysis. Standard (ATCC) bacterial strains of *Staphylococcus aureus* and *Escherichia Coli* were used. Standard blank sterile disks were impregnated with 25µl of the Generic and

Branded formulations of Levofloxacin, Ceftriaxone, and Meropenem after serial dilutions, achieving a final concentration equivalent to commercially available discs. The antimicrobial activity of the formulations was evaluated using the Kirby Bauer disk diffusion method on Mueller Hinton Agar plates, and zones of inhibition were measured using an Antibiotic zone scale as per CLSI guidelines. Statistical analysis was done.

Results

and Branded formulations of all 3 Antibiotics in both bacterial strains was in the Standard range as per CLSI guidelines.

Conclusion

The study concludes that generic formulations of Levofloxacin, Ceftriaxone, and Meropenem are as effective as their branded counterparts In Vitro. These findings support the use of generic antibiotics as cost-effective alternatives to branded formulations, without compromising therapeutic efficacy. The above study will be further supported with a larger group of samples and variable parameters.

The mean value of the Zone of inhibition for Generic

Abstract 028

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Myocarditis - A Punch From Gut ”– A Complicated Enteric Fever Case Report

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Background

Enteric fever is a common infectious disease occurring worldwide, especially in developing countries. The mortality of enteric fever used to be 10-20% in the pre-antibiotic era. The fatality has significantly decreased to < 1% in the endemic areas with antibiotic treatment. Enteric fever can have diverse extra-intestinal complications including encephalitis, Guillain-Barré syndrome, endocarditis, osteomyelitis, renal abscess, and splenic abscesses. Myocarditis is a rare complication of enteric fever accounting for about 1-5% of affected cases.

Summary of Case

A case 13-year-old girl, presented to pediatric OPD with complaints of fever associated with chills, haematuria, pain abdomen, vomiting, and bi-frontal headache for 4 days with the rest of the vitals being stable. CBP, CUE, Blood culture, Urine culture, CRP, LFT, RFT dengue serology, and Widal were sent. USG abdomen was done which showed no abnormal findings. Urine culture did not show any growth. CRP was found to be in the normal range and Serology for dengue was negative.

Important Findings

Blood culture was flagged positive after 2 days, which was identified to be *Salmonella Typhi*. 2D Echo and ECG were taken on the background of chest pain which showed LV dysfunction and signs of myocarditis respectively. Based on these, the diagnosis was made as complicated enteric fever for which she received IVIG – 1 dose and Ceftriaxone for 14 days. She recovered fully with reversed 2D echo and ECG changes.

Conclusion

of enteric fever like myocarditis which can occur even in immune-competent cases, pose a diagnostic challenge for paediatricians and can be fatal if not treated properly and timely. Hence ECG and Echocardiography should be part of the regular assessment of enteric fever. These patients should also be under close follow-up till ventricular function is fully recovered.

In developing countries like India, rare complications

Abstract 029

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Infectious Diseases and the Status of Women in India with a Specific Emphasis on the Caste Aspect

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Background

Biological differences, social determinants, and cultural norms contribute to increased susceptibility and vulnerability to infections among women. Women are more prone to reproductive tract infections and urinary tract infections, fungal infections, and sexually transmitted infections (STIs/RTIs) such as HPV, syphilis, gonorrhea, herpes, and Chlamydia due to biological and behavioural factors. In the patriarchal structure of Indian society, where women often hold subordinate positions, their health is frequently overlooked or deprioritized. This neglect is further compounded by various socio-cultural, educational, economic, and geographical factors that negatively influence women's health. Indian women are disproportionately affected by a range of infectious diseases, facing a higher likelihood of infection and more severe disease outcomes than men. Due to cultural and economic constraints, underprivileged women face a higher risk of HIV/AIDS infection. Alarming, 24% of Indian women are undernourished, while 54% suffer from anemia.

Methods

In-depth interviews of 54 women were recorded using stratified random sampling from different castes (SC, ST, OBC, and general). Women from rural and urban areas with varying education levels and socioeconomic status were included. Secondary data analysis of existing literature and reports were made. Informed consent and Confidentiality and anonymity with Ethical approval from the Institutional Review Board were taken.

Results

contracting infectious diseases. Dalit women belonging to lower castes endure the compounded disadvantages of caste and gender discrimination, leading to a life expectancy that is, on average, 15 years shorter than that of upper-caste women. The root causes include poor nutrition, strenuous labor, and inadequate healthcare access. Lower caste women's access to healthcare is severely restricted, heightening their susceptibility to infectious diseases. WHO reports indicate a higher prevalence of maternal mortality, anemia, and other health issues among lower caste women, exacerbated by poor healthcare access, sanitation, and nutrition. Moreover, social and cultural

practices like child marriage and restricted mobility further elevate their disease risk. Addressing the health inequities faced by lower caste women necessitates a comprehensive strategy, including policy reforms, education, and community initiatives. The National Family Health Survey shows a marked disparity in life expectancy between general and lower castes (SC/ST), with a significant risk of HIV-AIDS and prevalent skin conditions among Dalits.

Conclusion

Throughout their lives, women are at continuous risk of

Key to improving women's health status in India are education, awareness of personal health, economic independence, and autonomy in health-related decisions, particularly reproductive health. We emphasize the need for gender-sensitive public health strategies, improved access to healthcare, and increased research to address the unique needs of women in preventing, diagnosing, and treating infectious diseases. These are crucial measures for enhancing the overall health and wellness of Indian women.

Abstract 030

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Analysis of Changing Trends of Antimicrobial Resistance in *Enterobacterales* Isolates to Commonly Used β -Lactam Antibiotics in a Tertiary Care Hospital: Sub-Himalayan Region

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Background

Multidrug-resistance in gram negative bacteria is a global threat, requiring urgent attention. Third-generation cephalosporin-resistant and other broad spectrum β -lactam antibiotic-resistant *Enterobacterales* are emerging worldwide and are classified as high-priority pathogens by WHO. Carbapenem-resistant *Enterobacterales* were uncommon in the past and served as treatment options for these resistant organisms, but the trend is shifting in recent years.

Methods

Study includes retrospective analysis of resistance trend to commonly used parenteral β -lactam drugs in *Enterobacteriaceae* isolates obtained from culture positive clinical samples, between Jan 2022- May 2024 in Department of Microbiology, DRPGMC Tanda.

Results

Total of 8,185 *Enterobacterales* isolates were obtained from 2022 to 2024. In 2022, a total of 3262 of *Enterobacterales* were obtained, followed by 3602 in 2023 and 1321 in 2024 (till date). Ceftriaxone showed steadily increasing resistance rates of 60.94% in 2022, 63.32% in 2023 and 65.44% in 2024. Piperacillin-tazobactam showed a drastic increase in rate of resistance from 24.81% in 2022 to 64.50% in 2023 and 72.34% in 2024. Meropenem showed steady resistance rates of 17.29% in 2022, 16.14% in 2023 and 13.85% in 2024.

Conclusion

There is an increase in overall resistance rates during the study period. This could be due to over-the-counter use of broad-spectrum antibiotics, unnecessary antibiotic use, poor adherence to hospital antibiograms by clinicians, poor implementation of HIC and AMSP.

Abstract 031

ASPICON 2024 | August 22 -25 | 2024

Trends in Antimicrobial Resistance Patterns Among Gram Negative Non-fermenters in Intensive Care Units of a Tertiary Care Hospital in Eastern India.

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Background

Antimicrobial resistance (AMR) is increasing at an alarming rate among Gram negative bacteria (GNB) in intensive care units in recent years and has become a matter of concern around the world.¹ Limited availability of evidence-based data and empirical antibiotics, non-fermenting Gram-negative bacilli (NF-GNB) have emerged as noteworthy pathogens in health care setting.² This study aimed to find out the prevalence of different non fermenter GNB in ICUs and to analyse the trend of antimicrobial resistance of the isolates over a period of 5 years.

Materials & Methods

It is a cross-sectional retrospective study from January 2019 to December 2023 including 993 isolates identified by VITEK 2 from clinical samples of various ICUs in KIMS hospital, Odisha. Data collected was entered into Microsoft Excel spreadsheet, analysed using Epi Info software, presented using descriptive

statistics. MH-chi-square was used as a test of significance with a p-value of 0.05 considered statistically significant.

Results

Over a 5 year period the most commonly isolated non fermenter GNB was *Acinetobacter* spp. (44.6%) followed by *Pseudomonas* spp. (29.7%) and *Burkholderia* spp. (13.7%) and other non-fermenter GNB. With advancing age of the patients, a statistically significant increasing trend in presentation of GNB seen in *Acinetobacter* spp. (p=0.042) followed by

Burkholderia spp. (p=0.048). Decreasing trend of resistance to antibiotics in *Acinetobacter* spp. were seen for carbapenems (88 % to 81.5%) and ciprofloxacin (86% to 79.6%). Increasing trend of resistance seen for piperacillin/tazobactam (8.8 % to 85.2%) and amikacin (55.6% to 72.2%). In *Pseudomonas* spp. aztreonam (74.3% to 54.2%) and ceftriaxone (2.9% to 0) showed decreasing trend of resistance.

Conclusion

Acinetobacter spp. is the most common isolated bacteria showing increased sensitivity to carbapenems and Pseudomonas spp. showed increased sensitivity to aztreonam. The trend of antimicrobial resistance highlights for thorough follow up of antibiograms in hospitals and their judicious use by implementation of antimicrobial stewardship.

Abstract 032

ASPICON 2024 | August 22 -25 | 2024

Integrating Aerobic and Anaerobic Cultures in Wound Bio-Burden for Surgical Decision Support

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Background

Necrotising soft tissue infections (NSTIs) are rare but severe conditions characterised by rapid necrosis extending into deep fascia. Diagnosis is challenging and delays in treatment can be fatal. Microbiological cultures are essential to identify the causative organisms and understand the infection.

Methods

This study investigated the wound bioburden in 169 tissue biopsies from patients with NSTIs using both aerobic and anaerobic cultures. Tissue samples were cultured on blood agar and MacConkey agar plates, incubated under both aerobic and anaerobic conditions, and colonies were identified using MALDI-TOF MS.

Results

Results showed a wide range of bacterial counts, with heavy or overburdened plates reaching 1017 CFU/ml/g, while positive necrotic tissue had a minimum count of

109 CFU/ml/g. Most cases required amputation due to the severity of the infection. A semi-quantitative ranking system (sterile, scanty, mild, heavy, overburdened) based on aerobic blood plates was developed to aid in the rapid assessment of active infection at the point of care, facilitating surgical decision-making.

Conclusion

Integrating aerobic and anaerobic cultures, along with standardized quantitative and semiquantitative assessments, provides a comprehensive understanding of wound bio-burden in NSTI biopsies. This point-of-care knowledge enables timely and informed decisions regarding surgical intervention in cases of active infection.

Abstract 033

ASPICON 2024 | August 22 -25 | 2024

Unravelling the Microbiological Silhouette Causing Orthopaedic Infections in a Tertiary Care Centre

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Background

Orthopaedic infection of the bone, joint and associated structures are commonly caused by bacteria, which is acquired either by endogenous or exogenous routes (surgical site, trauma, and implant related infections), posing a global challenge with alarming consequences in resource constrained settings leading to increased medical cost, drug resistance and difficult to manage infections resulting in significant mortality. This study aimed to determine the bacteriological profiles and antimicrobial susceptibility patterns associated with orthopaedic infections.

Methods

An institution-based retrospective study was conducted from Jan 2023 to April 2024 at VMMC and Safdarjung Hospital, New Delhi. About 2467 patients with clinically suspected orthopaedic infections were enrolled. Aseptically collected Wound swabs or pus aspirates were processed aerobically for culture and sensitivity and antibiotic sensitivity testing as per standard microbiological techniques and CLSI 2023. SPSS version 25 was used for analysis.

Results

Most patients were 18-30 years old (34.5%), with a male preponderance (77.4%). The infection was more in winter season (35.6%). Pathogenic bacteria were grown in 54.3% of samples. *Staphylococcus aureus* (22.5%) and *Klebsiella pneumoniae* (7.1%) were the most common organisms. Gram positives were mostly resistant to Penicillin, Ciprofloxacin, and Erythromycin, while Gram negatives were mostly resistant to Amoxicillin-clavulanic acid, Cefepime, Cefotaxime, Ciprofloxacin, Piperacillin-tazobactam and Aminoglycosides.

Conclusion

The most isolated bacteria were *S. aureus* and *Klebsiella* spp. To mitigate the problem, aseptic surgical practice, conventional wound management, appropriate infection control practices and constant observation of antimicrobial resistant patterns, should be followed.

Abstract 034

ASPICON 2024 | August 22 -25 | 2024

Patterns of Antibiotic Prescription in Outpatients at a Public Health Care Facility in a Northern State of India

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Objectives

Conducting Outpatient survey (OPS) is especially challenging in the absence of electronic medical records, lack of dedicated resources and a high patient load in resource poor settings. The study aimed to analyze the patterns of antimicrobial prescriptions among patients visiting outpatient departments of a secondary care facility in an industrial town in northern India. This survey was conducted to provide background data for planning and strengthening the antimicrobial stewardship program (AMSP) in a public health care facility in Baddi, Himachal Pradesh.

Methods

A prospective, single-center outpatient survey (OPS) on antimicrobial prescriptions was carried out at Civil Hospital, Baddi, Himachal Pradesh, India. Information on patient demographics, reasons for antimicrobial use, and details of each prescribed antimicrobial, including supportive investigation reports, was collected using predesigned forms. The IEC approval was taken from Postgraduate Institute of Medical Education and Research (PGIMER), Chandigarh (IEC No. IEC-10/2022-2575).

Results

From September 2023 to June 2024, a total of 2098 patients were screened for potential eligibility visiting

the outpatient facility at Civil Hospital, Baddi. 1400 (66.73%) were prescribed antibiotic(s). The commonest indication for which antibiotic was prescribed was Urinary tract infections (UTI) (24.2%). Majority of prescriptions included single antibiotic (85.7%) and third generation oral Cephalosporin, Cefixime accounted for 42.2% of these prescriptions. Though the majority of prescriptions were oral, 2.6% patients received injectables. Majority of prescriptions were from the 'Watch' category of antibiotics. Importantly, the culture facility at the site was used for guiding a treatment in a vast majority, *E. Coli* was the commonest isolated organism.

Conclusion

The present study gives us a background of prevalent conditions presenting at the setting, profile of organisms and their susceptibility. Secondly, an evidence based policy will be generated for the facility.

Abstract 035

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Antibiotic prescription patterns among Admitted Patients in a tertiary Healthcare Facility in India: findings from a Point Prevalence Survey

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Background

In India, data on antibiotic prescription patterns at the patient level is scarce, and intervention measures to optimize antibiotic prescription patterns and reduce antimicrobial resistance are rarely performed. This study aims to determine the antibiotic prescribing pattern in admitted patients in the tertiary care hospital, in North India.

Methods

A Point Prevalence Survey (PPS) on antibiotic prescription patterns according to the World Health Organization (WHO) Methodology for PPS on antibiotic prescription in hospitals, was carried out in tertiary care hospitals. PPS was conducted over 2 weeks across five different wards in this hospital. Eligible inpatients

were those who were hospitalized in the ward at 8.00 a.m. on the day of the survey.

Results

In our findings, the overall prevalence of antibiotic usage among inpatients was 62%. Empirical treatment of antibiotic prescriptions was found in 95.2% of the patients. In 10% of patient's bacterial culture and AST were performed. In most of the patients, an antibiotic course was started before the sample was sent for culture, and cultures were sent on the 2nd or 3rd day of admission. The antibiotic ceftriaxone, which belongs to the WHO "watch" category, was the most prescribed antibiotic (49%), followed by azithromycin (16.40%) and metronidazole (16%), both belonging to the WHO "access" group of antibiotics. Overall, parenteral routes

of antibiotic administration were common in 84% of patients, whereas oral drugs were given only in 19% of the patients. We observed that 30% of patients received more than one antibiotic and broad-spectrum antibiotics were prescribed in more than 50% of cases. Antimicrobial prescriptions for both medical and surgical prophylaxis were 23% and 12% respectively.

Conclusion

In conclusion, the findings suggest the need for strengthening diagnostic and antimicrobial stewardship programs given the high proportion of antibiotic use, underutilization of bacterial culture, and complete reliance on empirical antibiotic treatment. Furthermore, this study has identified areas for quality

improvement, including education programs focusing on prescription practice in admitted patients and the urgent need for antimicrobial stewardship policy in hospital settings.

Abstract 036

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Evaluating the consumption pattern of antibiotics in the In-patient clinical department of a tertiary care hospital by using WHO AWaRe and DDD/100 bed-days

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Background

Antibiotic resistance is on rise and has become a worldwide problem, because of higher utilization and irrational uses of antibiotics. Consumption patterns of drugs can be classified by WHO AWaRe and measured by ACT/DDD/100 bed-days (Define daily dose) methodology. Surveillance data of antibiotics consumption can be used for developing hospital antibiotic policy as well as stewardship programs.

Methods

Detailed data of antibiotic dispensed to various clinical Inpatient (IPD) from 01/04/ 2023 to 31/03/ 2024 was collected from Central hospital pharmacy and pattern of use and consumption was calculated in terms of ACT/DDD and AwaRe Classification.

Results

Total 19 antimicrobials were used during the study period. Out of which 6 (Access), 7 (Watch), 1 (Reserve), 5 (Non-recommended) and 13 (E.M.L.), 1 (Non-EML) and 5 (Non-classified). Majority Cephalosporin (54.60%, {Ceftriaxone 31.12 DDD/100 bed-days and Cefotaxime 5.1 DDD/100 bed-days) class of drug were most commonly used followed by imidazole derivatives (16.97%, 8.2 1 DDD/100 bed-days) and lastly penicillin including beta-lactamase inhibitor (6.11%, 2.55).

Conclusion

As the data shows in comparison to antibiotics of the Access category, antibiotics of the Watch group were used more frequently and should pay attention toward the non recommended category because consumption of this class is also a considerable amount. 3-4 antibiotics are used more than others that will lead to resistance. Institutions must go for an antibiotic policy

and stewardship program that will help to procure and dispense the antibiotics as per requirements.

Abstract 037

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An Evaluation of Antibiotic Prescription Pattern and Drug Rationality Analysis Among Outpatients at public health setting, India

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Background

Antimicrobial resistance (AMR) poses a significant threat to global public health, increasingly prevalent due to inappropriate antibiotic prescribing patterns. Rational prescribing is essential to combat AMR, as rational analysis can identify irrational practices and build physician consensus. This study evaluates antibiotic prescription patterns and assesses the rationality of drug use among outpatients in a public health setting in India.

Methods

A cross-sectional study was conducted at a 30-bed public community health centre serving a population of

12,900, including urban and rural residents, between August 2021 and August 2022. Data were collected from 1,219 outpatient antibiotic prescription slips, randomly selected, and analyzed for drug type, dosage, duration, and adherence to clinical guidelines. For rationality assessment, each prescription was analyzed by a team of ID specialists (HS) and infectious disease clinical pharmacologists (NS) based on standard principles of rational prescriptions of antimicrobials. When required, NCDC treatment guidelines for infections, guidelines for the management of infections by the Postgraduate Institute of Medical Education and Research, and standard reference guidance documents were referred. Data was analyzed to evaluate demographics, the most prescribed antibiotics their

class and AWaRe category, diagnoses among these patients, drug rationality, and the drugs from the Essential Drug List (EDL).

Results

A total of 1,219 antibiotic prescription slips were analyzed. Among these, 45.9% were for males and 54% for females, with most patients (30.9%) being aged 20-40 years. Oral infections (21.4%) were the most common diagnosis followed by Skin and soft tissue infection (20%). The most frequently prescribed antibiotic was amoxycylav (27.2%), followed by metronidazole (13.4%) and azithromycin (10.3%). According to the WHOAWaReclassification, 49.7% of the prescribed antibiotics were in the "Access" group, while 27.3% were in the "Watch" group. No antibiotics were prescribed from the "Reserve" group. Regarding rationality, 57% of prescriptions for urinary tract

infections were rational, whereas only 29% of respiratory tract infection (RTI) prescriptions were rational. Diarrhea cases showed the highest rate of irrational antibiotic use (95%) followed by RTI (71%).

Conclusion

The study highlights significant issues in antibiotic prescription practices, with a considerable proportion not adhering to recommended guidelines. These findings underscore the necessity for optimized antibiotic prescribing practices to combat AMR in outpatient settings in India. Implementing targeted education programs for health care providers, enhancing prescription monitoring systems, and adhering to evidence-based guidelines are critical steps to promote rational antibiotic use and improve public health outcomes.

Abstract 038

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Descriptive Analysis of Antimicrobial Resistance Patterns in WHO-Critical Priority Pathogens from Clinical Samples in ICU at a Tertiary Care Hospital, India

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Background

Antimicrobial resistance (AMR) is a serious concern to world health, especially in intensive care units (ICUs) where patients are more susceptible to infections. The World Health Organization (WHO) has identified certain pathogens as a critical priority due to their high resistance rates and limited treatment options. It includes *A. baumannii* (carbapenem-resistant), *P. aeruginosa* (carbapenem-resistant), and *Enterobacteriaceae* (carbapenem-resistant and third-generation cephalosporins resistant). This study aims to provide a comprehensive analysis of AMR patterns in WHO-critical priority pathogens isolated

from clinical samples in the ICU of a tertiary care hospital in India.

Methods

Clinical samples were received from the ICU between January 2021 to December 2023. These clinical samples were processed by using conventional culture methods in the microbiology laboratory. Antibiotic susceptibility testing (AST) was performed by using disc diffusion methods as per CLSI guidelines. The pathogens were categorized according to the WHO priority list, and their resistance patterns were analyzed.

Results

A total of 4,339 clinical samples were received from the ICU. From clinical samples, 850 WHO-critical priority pathogens were identified. The most common isolates were *A. baumannii* 365 (43%) followed by *K. pneumoniae* 192 (23%), *P. aeruginosa* 178 (21%), and *E. coli* 115 (13%). The resistance to carbapenems was observed to be 88% in *A. baumannii* and 36% in *P. aeruginosa*. In *Enterobacteriaceae*, carbapenem resistance was found to be 54%, while resistance to third-generation cephalosporins was 81%.

Conclusion

The antibiotic resistance among critical priority pathogens observed in this study reflects a significant challenge in treating ICU infections in a tertiary care hospital in India. The high rates of resistance necessitate urgent action in implementing robust antibiotic stewardship programs and enhancing

infection control practices to mitigate the spread of resistant pathogens. The data generated can aid in developing targeted treatment guidelines and policy decisions to combat AMR effectively.

Abstract 039

ASPICON 2024 | August 22 -25 | 2024

Resistance Profile of Candida Isolates in Women with Vulvovaginal Candidiasis Attending Gynecology OPD in a Tertiary Care Hospital

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Background

Vulvovaginal Candidiasis (VVC) is a substantially common mucosal infection of the female genital tract caused by a polymorphic opportunistic fungus *Candida* spp. Over the years, a marked rise in vaginal yeast infections has been observed. Over the counter availability and rampant use of antifungals have contributed massively in development of antifungal resistance, thus have been conducive to increased morbidity among females with VVC.

Methods

197 (HVS) samples were collected from symptomatic females having at least one of the symptoms of itching, vaginal discharge and burning sensation. The samples were processed and subjected to antifungal

susceptibility (AFS) by both Kirby Bauer Disc Diffusion(DD) method and by using Octofungi 1 HiMIC plate kit. Three drugs were used for both the methods- fluconazole, voriconazole and caspofungin. The data was analysed statistically and comparison of the resistance pattern by both the methods was done and depicted in the form of a comparative table and graph.

Results

Of 197 samples, 52 were positive for VVC. The species isolated were *C. albicans*(26), *C.parapsilosis* (10), *C. tropicalis*(9), *C. krusei*(5) and *C. glabrata*(2). For fluconazole, *C. albicans*, *C. tropicalis* showed no resistance by DD method and MIC testing. However, (1/2) of *C. glabrata* showed resistance by disc diffusion whereas both(2/2) were resistant by MIC. Regarding voriconazole, 3.8% resistance was observed for *C.*

albicans, 50% for *C. glabrata*, 10% for *C. parapsilosis* and 11.1% for *C. tropicalis* by MIC testing. Similarly, for caspofungin, all species were susceptible to DD. Whereas, *C. krusei*, and *C. tropicalis* showed no resistance by MIC but *C. albicans* (3.8%), 50% *C. glabrata* and 10% *C. parapsilosis* showed resistance by MIC testing.

Conclusion

Testing by MIC method is a better method for antifungal susceptibility testing, as described by CLSI. Since resistance in *Candida* isolates is on a rise, it is important to incorporate routine AFS testing which would improve patient outcomes and ultimately reduce morbidity and mortality.

Abstract 040

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Prevalence and Antimicrobial Susceptibility Profile of Multidrug Resistant *S. maltophilia* Causing Infections in a Tertiary Healthcare Setting in North India

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Background

Stenotrophomonas maltophilia is an emerging opportunistic pathogen in healthcare settings. It is notoriously resistant to many antimicrobials intrinsically and has shown acquired resistance increasingly to many antimicrobials due to multiple drug resistant mechanisms, making therapeutic management of critical cases difficult. This study aimed to determine the prevalence and antimicrobial susceptibility profile of *S. maltophilia* causing infections at a tertiary hospital setting.

Methods

This observational study was conducted for a one-year duration from 1st May 2023 to 31st May 2024 in the

department of microbiology after taking ethical approval from the institute's ethical committee. All clinical samples received were subjected to aerobic culture at 37°C. Identification and susceptibility testing was performed using VITEK 2 COMPACT automated systems. Antimicrobial susceptibility profile was interpreted in accordance with CLSI breakpoints.

Results

Out of 1870 non lactose fermenters isolated after aerobic culture from various clinical specimens, 66 (3.52%) were identified as *S. maltophilia*. The isolates were predominantly recovered from 41-60 (34.84%) age group. A male predominance of 1.5:1 was seen. The isolates were identified as multi drug resistance organisms (MDROs) and were found sensitive to

minocycline (80.73%), levofloxacin (71.21%), cotrimoxazole (63.36%), ticarcillin and clavulanic acid (62.12%) and chromaphenicol (59%). Resistance was seen predominantly for tetracycline (100%) and ceftazidime (61%).

Conclusion

Although a low prevalence of *S. maltophilia* was reported at our setup, the antimicrobial susceptibility profile determined would be useful in selection of appropriate antimicrobials to manage infections caused considering the paucity of data on the same.

Abstract 041

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Isolation of *Cedecea lapagei* in a Case of Hemothorax: A Rare Emerging XDR Organism

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Background

Cedecea is a gram negative bacteria belonging to the *Enterobacteriaceae* family with six known species of *Cedecea*. Among them three of these species are said to be human pathogens: *Cedecea davisae*, *Cedecea lapagei* and *Cedecea neteri*. It was first described in 2006 as a potential pathogen in a peritoneal fluid of a 55 years old man having peritonitis. *Cedecea lapagei* was further isolated in pneumonia, bacteremia, soft tissue infection, peritonitis, sepsis, hemoptysis and urosepsis patients. Till now, only thirteen cases of *Cedecea lapagei* have been described in the literature and none of them was extensively drug resistant. *Cedecea* causing infections at a tertiary hospital setting.

Case Summary

A 60 year old female patient of a road traffic accident was brought to the emergency department at AIIMS Bathinda with a history of difficulty in breathing for 1 day. She was diagnosed with hemothorax & subsequently an intercostal drain tube was placed. On day 7th, the patient developed a fever, so ICD content were sent for microbiological bacterial culture and

susceptibility. Growth of *Cedecea lapagei* was obtained on the culture which showed resistance to piperacillin tazobactam, cefuroxime, ceftazidime, ceftriaxone, cefepime, ertapenem, imipenem, meropenem, doxycycline, tetracycline, amikacin, gentamicin, ciprofloxacin, cotrimoxazole and intermediate susceptible to cefaperazone- sulbactam and susceptible to minocycline only. Repeat sample was taken to exclude the contamination and for confirmation of

suspected bacteria. Same organism with a similar susceptibility pattern was obtained on repeated culture. Patient was put on minocycline and after 3 days a sample was sent which was sterile after 48 hours of incubation. Patient became afebrile and the ICD tube was removed. Patient was discharged under stable conditions.

Conclusion

We report a rare case of hemothorax having infection of *Cedecea lapagei* which was multi-drug resistant. Earlier it was rarely described in the literature to be extensively drug resistant. Hence, early identification of this bacteria and correct antibiotic treatment is important for good outcome.

Abstract 042

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Surgical Site Infection in Pediatric Population: Incidence, Etiology and Risk Factors

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Background

Surgical site infection (SSI) is one of the most common healthcare-associated infections (HAIs). However, there is a lack of data available about SSI in children, especially from low-income and middle-income countries. In children, SSI has important social and economic consequences, leading to loss of school days and lost working days by parents. The burden of SSI in terms of healthcare costs and the requirements for antibiotic therapy have important consequences for antibiotic resistance and health system capacity.

Methodology

A retrospective study was conducted over a period of 5 years from January 2019 to December 2023 to study the incidence, etiology and risk factors of SSI in children less than 12 years of age undergoing gastrointestinal, urological and orthopaedic surgery in a government paediatric tertiary care centre.

Result

Over a period of 5 years, samples from 12628 surgeries were conducted, out of which, 365 patients developed SSI, SSI rate being 2.89 SSI per 100 operative procedures. Out of these 365 patients, 55.9% of patients were males and 44.1% were female patients. 14.8% of these patients were neonates. In 10.4% of the patients,

infection was present at the time of surgery (PATOS). The most common causative agent of SSI was found to be *E.coli* (49%), followed by polymicrobial growth (15.9%), and *Acinetobacter baumannii* (7.7%). The most common risk factors associated were low birth weight, malnutrition, lack of traffic control in OT, patients not taking pre operative showers, longer duration of surgeries, anastomosis leak, inadequate staffing for post op care in ward, and contamination of incision post op.

Conclusion

SSI represents a significant burden of postoperative morbidity in children who receive surgery. Focused initiatives and research, aiming to reduce SSIs in children should be a key priority for surgery agendas.

Abstract 043

3-Step Model- An Explorative Novel Approach to Classify Sepsis: A Longitudinal Observational Study

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Introduction

Sepsis remains a critical healthcare challenge worldwide, demanding prompt identification and treatment to improve patient outcomes. Given the absence of a definitive gold standard diagnostic test, there is an imperative need for adjunct diagnostic tools to aid in early sepsis detection and guide effective treatment strategies. This study introduces a novel 3-step model to identify and classify sepsis, integrating current knowledge and clinical guidelines to enhance diagnostic precision.

Methodology

This longitudinal observational study was conducted at a tertiary care teaching hospital in northern India. Adult

patients admitted with suspected sepsis underwent screening using predefined criteria. The 3-step model consisted of Step 1, assessing dysregulated host response using a National Early Warning Score-2 (NEWS-2) score of ≥ 6 ; Step 2, evaluating risk factors for infection; and Step 3, confirming infection presence through clinical, supportive, or confirmatory evidence. Patients were categorized into Asepsis, Possible sepsis, Probable sepsis, or Confirmed sepsis at various intervals during hospitalization.

Result

A total of 230 patients were included. Initial categorization on Day 1 showed 13.0% in Asepsis, 35.2% in Possible sepsis, 51.3% in Probable sepsis, and 0.4% in confirmed sepsis. By Day 7, shifts were observed with

49.7% in Asepsis, 9.5% in Possible sepsis, 25.4% in Probable sepsis, and 15.4% in confirmed sepsis. At discharge or death, categories were 60.4% Asepsis, 5.2% Possible sepsis, 21.7% Probable sepsis, and 12.6% Confirmed sepsis. Transitions between categories were noted throughout hospitalisation, demonstrating the dynamic nature of sepsis progression and response to treatment.

Conclusion

The 3-step model effectively stratifies sepsis status over hospitalization, facilitating early identification and classification of septic patients. This approach holds promise for enhancing diagnostic accuracy, guiding clinical decision-making, and optimizing antibiotic stewardship practices. Further validation across diverse patient cohorts and healthcare settings is essential to confirm its utility and generalizability.

Abstract 044

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Bacteriological Profile and Antimicrobial Susceptibility Pattern in Blood Culture Specimen in a Tertiary Care Hospital of Uttar Pradesh: Six Years Retrospective Study

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Background

Globally, bloodstream infections (BSI) are major contributors to morbidity and mortality. Antimicrobial surveillance is crucial for spotting emerging resistance and developing empirical treatment guidelines. The goal of this study is to examine trends in BSI antimicrobial susceptibility from 2018 to 2023 in a tertiary care centre.

Methodology

This was a retrospective study conducted in the department of microbiology, Sarojini Naidu Medical College, Agra. Blood samples for culture were received from various clinical departments which were processed by conventional methods for identification.

Result

4195 samples were tested positive from a total of 11882 samples. Most of the samples that tested positive were from the pediatric age group. Klebsiella species was the

most frequent organism to be isolated (45%) amongst the gram-negative organisms followed by *E. coli* species (30.9%). Other gram-negative bacilli that were isolated included *Burkholderia* species, *Acinetobacter* species, *Pseudomonas* species, and *Citrobacter* species. While *klebsiella* species demonstrated a decline in resistance to doxycycline and ceftriaxone sulbactam antibiotics, they demonstrated a significant rise in resistance to imipenem and piperacillin tazobactam antibiotics. However, there was no change in the resistance pattern of cotrimoxazole and ceftriaxone during the years. Antibiotic resistance to amikacin, piperacillin, tazobactam, and cotrimoxazole increased significantly in the *E. coli* isolate, while antibiotic resistance to levofloxacin, doxycycline, and ceftriaxone decreased in these 6 years' time. Among the various antibiotic groups, *Pseudomonas* species exhibited the highest emergence of resistance, with the exception of piperacillin and carbapenem antibiotics.

Conclusion

Rising resistance to last-resort antibiotics such as beta-lactam and carbapenem antibiotics, especially in Gram-negative bacteria, indicates that there is urgent

need for exploring new antibiotics such as Fosfomycin, tigecycline and other antibiotics against these carbapenem resistant isolates and stronger antimicrobial stewardship efforts to combat antimicrobial resistance and stop the spread of multidrug resistant bacteria amongst population.

Abstract 045

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Analysis of Antibiotic Consumption Patterns, Practices and Trends Based on Procurement Data of Public Health Institutions in the Hilly State of Himachal Pradesh, India. (2018-2022)

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Introduction

India, despite being the largest consumer of antibiotics in absolute volume, lacks a formal system for antibiotic use unlike the US and Europe. Consequently, much antibiotic use data remains "invisible." This paper aims to make antibiotic use data within the public health sector of a northern Indian state "visible".

Methodology

This study analyzes antibiotic use patterns and trends in five Public Health Facilities in Himachal Pradesh from 2018 to 2022. The analysis employs the Anatomical Therapeutic Chemical (ATC) classification system and uses Defined Daily Dose (DDD) and Drug Intensity Dosage (DID) as metrics.

$$Y_i = \sum_{i=1}^5 M_i / \frac{ni}{Ni}$$

Using the formula where:

Yi: The adjusted DDDs in a given year in a certain type of healthcare institution i;

Mi: DDDs used in a given year in a certain type of healthcare institution i;

ni: Number of a certain type of healthcare institution i from where data was collected; PHC, CHC, DH, Tertiary and Super Speciality Hospitals

Ni: The total number of a certain type of healthcare institution i in the Health Statistics Yearbook

Result

This study examines antibiotic procurement in Himachal Pradesh from 2018 to 2022. During this period, total antibiotic consumption initially increased, peaking in 2019, a sharp decline in 2020, due to COVID 19 followed by a steady rise and then a decline in 2022. The most commonly consumed drug classes were Macrolides followed by Fluoroquinolones and Penicillins whereas Cephalosporins & Carbapenems were the least consumed.

Conclusion

This study provides empirical evidence on trends in antibiotic procurement and expenditure from

2018-2022. The introduction of an essential drug list by the state government improved medicine availability at hospital pharmacies. Concurrently, interventions like prescription audits, digitization of procurement and consumption data, and antimicrobial stewardship have arguably contributed to the decrease of antibiotic consumption. However, the patterns of antibiotic use do not provide strong evidence to show rational use of antibiotics.

Abstract 046

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Identification of Factors Driving Decision Making in Prescribing Empirical Antibiotics-An Observational, Cross-sectional Pilot Study

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Introduction

Antimicrobial Stewardship requires multiple stakeholders who are responsible for patient care. Hence it is prudent to understand the factors which are responsible for decision making for prescribing empirical antibiotic therapy. The pilot study aims to identify the attributes and its levels affect decision making while prescribing empirical antibiotic therapy.

Methodology

An observational, cross-sectional study with study Population of 200 Registered medical practitioners will be done from May 2024 to July 2024 online through Google forms. An e-Informed consent will be taken before attempting the questionnaire in which the purpose of the study will be explained. Demographic profile of the doctor including the place of his practice (urban, rural) and type of practice (private, government) and his/her qualification along with the years of experience will be taken. A total 15 questions will be asked which shall not take more than 10 minutes to answer.

Result

Although data collection is ongoing, preliminary projections suggest that factors such as antibiotic spectrum, adherence to clinical guidelines, and patient-specific characteristics (e.g., age, comorbidities) are likely to emerge as significant drivers in the decision-making process. Cost is anticipated to have a moderate impact.

Conclusion

After completion of data analysis, the study will highlight the factors or attributes affecting the decision for prescribing empirical antibiotics. These insights will help in tailoring solutions for enhanced acceptance and implementation of antimicrobial stewardship by the clinicians. The results will help us conduct a robust and validated discrete choice experiment study.

Abstract 047

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Efficacy of Conventional and Molecular Methods in Clinically Suspected Cases of Musculoskeletal Tuberculosis in a Tertiary Care Hospital

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Objective

Lab diagnosis of clinically suspected cases of musculoskeletal tuberculosis by ZN staining, MGIT, and CBNAAT.

Material & Methodology

The samples were collected in the Department of Microbiology in association with the Department of Orthopaedics between August 2022 to March 2024. Eighty clinically suspected musculoskeletal

tuberculosis patients were included and the samples were processed for ZN staining, CBNAAT, and MGIT.

Result

The mean age of the study population ranged from 1.5 to 69 years with a maximum number of cases aged between 21 and 30 years (23.98%). Knee joints (n=33; 41.3%) followed by spine (n=31; 38.8%) were the most commonly involved sites. In half the cases (n=40; 50%), the specimen was fluid aspirate from the affected site, 33 (41.3%) were pus specimens and 7 (8.8%) were

biopsy specimens. The positivity rate for Ziehl-Neelsen staining and MGIT was 11.3% and 12.5% respectively. A total of 18 (22.5%) cases were confirmed as tuberculosis. ZN stains had the lowest sensitivity, negative predictive value, and accuracy (50%, 87.3%, and 88.8% respectively) followed by MGIT (55.6%, 88.6%, and 90% respectively). However, CBNAAT was not only highly sensitive (94.4%) but also had high negative predictive value (98.4%) and accuracy (98.8%).

Conclusion

The present study showed that CBNAAT is a useful, sensitive and specific molecular diagnostic method compared to MGIT and ZN stain for evaluation of extrapulmonary tuberculosis so the CBNAAT could be recommended as the first line diagnostic method for the detection of tuberculosis in clinically suspected musculoskeletal tuberculosis cases.

Abstract 048

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Achromobacter Species: An Underestimated Contributor to Post-operative Infections in Immunocompromised Patients in a Tertiary Care Cancer Center

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Background

Achromobacter is a genus of non-fermenting Gram-negative bacteria under order Burkholderiales. They are mostly associated with the respiratory tract of patients with cystic fibrosis. They are often isolated from severe healthcare-associated infections in immunocompromised patients.

Case Summary

A total of 8 cases of infections due to *Achromobacter* spp. were identified in the year 2023 at our centre; 4 cases each of *A. denitrificans*, and *A. xylosoxidans*. Most of the patients were males from 40 to 80 years of age. All the patients were diagnosed with advanced stage solid organ cancers and had undergone surgery in

the recent past, with the patients belonging to head and neck tumours accounting for the maximum cases. None of the above patients were lost during treatment and follow-up.

Important findings

Heavy growth of *A. denitrificans* was isolated from chemo port catheter and pus swab from pus/discharge oozing from surgical wounds. Significant growth of *A. xylosoxidans* was also isolated from urine, bronchoalveolar lavage and pus swab of post-operative admitted patients. The isolates were identified by manual biochemical reactions and the Vitek-2 automated ID and susceptibility testing system. *A. xylosoxidans* isolates showed high susceptibility to cotrimoxazole, cefoperazone-sulbactam and ceftazidime, and resistance to fluoroquinolones. *A. denitrificans* isolates showed high susceptibility to piperacillin-tazobactam and resistance to fluoroquinolones. Although the infections were serious, requiring ICU admission for some patients and long-term therapy, all the above patients recovered successfully.

Conclusion

These cases illustrate the important role of this bacterium in serious infections in patients with advanced-stage cancers of solid organ systems. Although infections reported by these bacterial isolates are less frequent, they can be a serious threat to immunocompromised patients. Differences in biochemical reactions and antibiotic susceptibility patterns from other commonly isolated Gram-negative bacteria should be borne in mind for rapid diagnosis and prompt institution of appropriate therapy.

Abstract 049

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Mycetoma in North India: A Clinico-epidemiological and Investigative Study- New Patterns of a Neglected Tropical Disease

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Background

Mycetoma is a chronic, suppurative granulomatous infection affecting skin and subcutaneous tissues and is characterized clinically by a triad of painless soft tissue swelling, underlying sinus tracts, and extrusion of grains. It is endemic in India and other tropical and subtropical countries between latitude 15° S and 30° N all around the globe.

Methodology

This study involved a cross-sectional design in which cases of mycetoma presenting to a tertiary care hospital in north India were included. Clinicoepidemiological parameters including age, gender, occupation, geographical distribution and site of involvement were noted. Diagnostic investigations included microbiological and histopathological analyses of the clinical samples. Radiological studies including X-ray/ultrasound and MRI were done to evaluate the extent of tissue involvement.

Result

A total of seven cases were studied. Female outnumbered males (71%) and age of the patients varied from 13-71 years. Site of involvement was right foot in 57.1% cases and left foot in 42.8% cases. 85% of the patients had occupational exposure in the form of working in fields. There was a history of passage of granules in 4 patients, however pale white granules were retrieved only in 2 patients. Etiologically, 71.4% cases consisted of actinomycotic mycetoma and 28.6% of cases were caused by fungi (eumycetoma). Confirmation of causative agent was done on gram staining, ZN staining, KOH examination and histopathology. However, species identification was not done due to non-availability in our center. Radiological studies were normal in 28% cases, 2 patient demonstrated soft tissue swelling, lytic lesions with focal erosions/scalloping were seen in 2 patients; and soft tissue radiopacity with sclerosis, conglomerate of hypoechoic nodules and dot in a circle appearance was seen in one patient. All cases of actinomycetoma were started on a modified Welsch regimen and cases of eumycotic mycetoma were started on itraconazole. All cases demonstrated significant improvement at 3 months follow up, assessed clinically and radiologically.

Conclusion

Traditionally eumycetoma is considered to be more common in north India and actinomycetoma is predominant in south India. However, by reporting the contrasting findings, elucidating disease characteristics, associated risk factors and causative agents in this specific geographical context, our study may shed light on changing pathogen profile, to facilitate better management, prevention and control strategies for this deforming granulomatous infection.

Abstract 050

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Performance Evaluation of Disk Diffusion Test Directly from Positive Blood Culture Broth for Gram negative Bacteremia - a Step Towards Antimicrobial Stewardship

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Background

Performing & reporting Direct Susceptibility Testing (DST) from positive blood culture broth will help in the management of critically ill patients with sepsis. However, published studies mention that categorical agreement (CA) between direct susceptibility testing from-blood culture by disk diffusion & disk diffusion

from colonies may vary between different laboratories. Therefore, a present study was conducted to determine categorical agreement (CA) and very major error (VME), major error (ME) & minor error (minE) of DST in comparison to the reference method.

Methodology

A retrospective observational study from February 2023 - May 2024 was conducted after Institutional Ethics Committee approval. DST (as per CLSI guidelines) results of Enterobacterales and non-fermenters (*Pseudomonas aeruginosa* & *Acinetobacter* spp.) were compared with reference AST method (Vitek-2 Compact & manual disk diffusion) done from colonies. CA and various types of errors of DST were evaluated. As per FDA, acceptable cut off for CA is $\geq 90\%$ and ME $< 3\%$. For VME, the upper limit of 95% confidence interval should be $< 7.5\%$.

Result

A total of 225 isolates were tested for 1186 antimicrobial agents. Overall CA was 94.10% (1116/1186). Non-fermenters showed better CA of 96.38% (133/138) compared to enterobacterales which showed CA of 93.80% (983/1048). A total of 2.53 % minE (30/1186), 6.43 % ME (29/451), and 1.51 %VME (11/727) were observed. Enterobacterales showed less numbers of minE as compared to nonfermenters (2.39% vs 3.62%). ME and VME were not observed in non-fermenters, however, enterobacterales showed 2.77% ME, and 1.05% VME. In Enterobacteriaceae, maximum ME & VME errors were seen with aztreonam (9/151, 16.39%) and ceftazidime (5/148, 4.35%) respectively.

Conclusion

DST from positive blood culture broth showed a good categorical agreement and range bound errors with reference method. DST can be confidently implemented for patient's reporting at our institute. In *Enterobacterales*, DST results of ceftazidime should be reported with caution.

Abstract 051

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Antimicrobial and Diagnostic Stewardship of a Novel Beta Lactam/Beta- Lactamase Inhibitor in Extensively Drug-Resistant Gram-Negative Pathogens: Establishing the Realm of Promise

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Background

Carbapenem resistant Enterobacterales and *Pseudomonas aeruginosa*, producing metallo-beta-lactamases, are a potential nosocomial threat. The combination of aztreonam (ATM) with ceftazidime-avibactam (CZA) is a possible therapeutic approach. However, rising resistance and lack of an approved and feasible testing method of synergy is worrisome. The study aimed to evaluate the synergy and rapid detection of carbapenemases.

Material & Methodology

50 extensively drug-resistant (XDR) isolates of Enterobacterales and *Pseudomonas aeruginosa* were selected. Synergy between ATM and CZA was assessed using the disk stacking method, by placing one disk each of ceftazidime, CZA, ATM and stacked disks of CZA over ATM onto a cultured Muller Hinton agar plate. Positive synergy was defined as individual resistance to both CZA and ATM but susceptible to the combination. Carbapenemases were determined in isolates exhibiting positive synergy by immunochromatographic based methods.

Result

Klebsiella pneumoniae, *Escherichia coli*, *Pseudomonas aeruginosa* and *Klebsiella aerogenes* accounted for 60% (30/50), 22% (11/50), 16% (8/50), and one isolate respectively. Overall, positive synergy was observed in 40% (20/50) of the isolates, maximum in *Escherichia*

coli (54.5%; 6/11), followed by *Klebsiella pneumoniae* (40%; 12/30), *Pseudomonas aeruginosa* (12.5%; 1/8), and the lone isolate of *Klebsiella aerogenes*. NDM was the predominant carbapenemase (50%; 10/20), followed by its co-expression with OXA-48 (25%; 5/20), OXA-48 alone (15%; 3/20), and no carbapenemase in two isolates. Clinical response to the synergy along with microbiological clearance was observed in 75% (15/20) of the patients

Conclusion

Albeit limitations, Disc stacking method, being economic, can be employed in clinical microbiology laboratories in resource limited settings, lacking rapid molecular diagnostic systems. The rapid reporting of synergy results enabled clinicians to tailor treatments. This further empowered in improvement of patient outcome, reduced duration of hospital stay, optimal usage of antibiotics and reduction in antimicrobial resistance. This helped in improvement of existing antimicrobial and diagnostic stewardship practices.

Abstract 052

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A Longitudinal Study of Antimicrobial Resistance Dynamics in *Pseudomonas Aeruginosa* - Implications in Treatment Strategies

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Background

Pseudomonas aeruginosa is a significant pathogen known for its resistance to multiple antibiotics, posing a challenge in both adult and pediatric patient populations. This study aims to analyze the antimicrobial susceptibility testing (AST) patterns of P.

aeruginosa isolates, focusing on differences between adult and pediatric populations, temporal trends and burden of multidrug resistant (MDR) isolates.

Methodology

The present study was conducted from 2021 to 2024 at a super-speciality government teaching hospital with medical and surgical departments. Samples were collected and processed as per clinical requests, and those showing growth were analyzed using an automated identification and antibiotic susceptibility test analyzer, following CLSI M-100 guidelines. Samples growing *Pseudomonas aeruginosa* were included, excluding repeat samples from the same patient. Colistin resistance was confirmed using colistin broth disc elution (CBDE) and compared with automated analyzer results. Suitable antibiotics for treatment were evaluated based on CLSI M39 guidelines.

Result

In the current study, *Pseudomonas aeruginosa* showed varying susceptibility rates to different antibiotics ranging from 43.85% to 54.68% in adults and 41.3% to 49.32% in pediatric populations. MDR isolates were 32.8% in total with 43.2%, 12.5%, 22.5% and 22.1% from urine, blood, respiratory and exudate samples. MDR rates increased from 2021 to 2023, reaching 43.6%. Colistin susceptibility rates were relatively high with 84.45% in adults, 90.41% in paediatric population and ranged from 79% to 93% across samples. There was a discrepancy of 14.9% between automated and CBDE methods for detection of colistin resistance. All antibiotics except colistin show less than 60% susceptibility, making higher antibiotics or combinations necessary for treatment..

Conclusion

This study highlights the increasing multidrug resistance in *Pseudomonas aeruginosa*, necessitating tailored treatment and enhanced surveillance. While colistin remains effective, its resistance requires ongoing monitoring. Effective antibiotic stewardship is crucial to managing these infections, especially in pediatric patients.

Abstract 053

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Randomized Comparative Study of Novel Teaching-Learning Methods Versus Conventional Pedagogy in Educating Phase II Mbbs Students on Different Aspects of Antimicrobial Use

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Background

Medical undergraduates are introduced to/(taught) the field of antimicrobials as a part of Pharmacology

curriculum in 2nd year of MBBS. Implementation of effective teaching learning methods during this phase can better equip the future healthcare professionals to acquire necessary knowledge and skills to practice rational use of antimicrobials as standalone physicians. The present work aims to compare the novel teaching-learning methods against the conventional pedagogy on various aspects of antimicrobial use.

The knowledge gain will be determined by comparing pre-test and post-test mean scores(+SD) obtained by both the groups after the teaching sessions. Further, two groups will be compared to assess if the experimental group outperforms the control group. The feedback of students will be utilized for improving the teaching strategies in future.

Material & Methodology

A randomised comparative study will be conducted among the phase II MBBS students after obtaining approval from the Institute Ethics Committee. After receiving written informed consent, the students will be randomly allocated to one of two groups: control group (using conventional pedagogy i.e. didactic lecture-based learning) or experimental group (using novel teaching-learning methods). A pretest will be administered to both the groups to have baseline assessment of the knowledge of the topic. After completion of respective teaching-learning sessions, a post-test (immediately after the session) will be administered and scores will be calculated. In addition, students' feedback regarding novel teaching-learning methods will be obtained with the help of a post intervention survey. Data will be analysed using descriptive statistics and student's t-test wherever appropriate.

Result & Conclusion

Abstract 054

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Comparison of Bacteriological Profile and Resistance Pattern in Two Consecutive Years in a Tertiary Care Hospital

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Background

Medical undergraduates are introduced to/(taught) the field of antimicrobials as a part of Pharmacology curriculum in 2nd year of MBBS. Implementation of

effective teaching learning methods during this phase can better equip the future healthcare professionals to acquire necessary knowledge and skills to practice rational use of antimicrobials as standalone physicians. The present work aims to compare the novel teaching-learning methods against the conventional pedagogy on various aspects of antimicrobial use.

Methodology

The bacteriology data of the past two years was collected, analysed and compared.

Result

Although the profile of bacteria isolated was largely the same in 2022 and 2023, there was a slight decrease in the number of *Klebsiella* species and *Acinetobacter* species isolated. There was a twofold rise in sample positivity from ICUs. The percentage of *Escherichia coli* isolated from urine specimens increased from 49% to 59% and *Staphylococcus aureus* from soft tissue and body fluid specimens increased from 34% to 40%. However, the percentage of *Pseudomonas aeruginosa* from respiratory samples decreased from 38% to 29%. There was an increase in the percentage of multi-drug resistance (MDR) isolates from 46.5% in 2022 to 57.9% in 2023. These include increased MDR strains of *E. coli*, *Klebsiella* species, *P. aeruginosa* and *Acinetobacter* species. Percentage of MRSA increased from 50% to 65.9% in the consecutive year.

The practices of infection control and awareness of prudent use of antibiotics needs to be made more stringent to curb the upward trend in frequency of MDR bacteria isolated.

Conclusion

Abstract 055

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Understanding Diabetic Foot Infections: Microbiological Insights and Clinical Implications

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Background

Diabetic foot infection denotes a serious complication of diabetes, characterized by microbial invasion into

foot tissues. Symptoms range from localized inflammation to severe ulceration and tissue necrosis, potentially leading to limb amputation if left untreated.

Most moderate-to-severe diabetic foot infections (DFIs) require hospitalization with urgent surgical approach and administration of empiric antibiotherapy. To ensure optimal antibiotic coverage, regular microbiological background updates are imperative.

Aim of this study is to characterize the microbiological profile and the antibiotic sensitivity pattern of the DFI causative pathogens per antimicrobial agents according to culture sensitivity.

Methodology

Our study is a prospective study in which 101 patients of moderate to severe Diabetic foot infection were admitted in Orthopaedic deptt of a tertiary care hospital of North India over a period of 3 years. Microbiological cultures from deep tissues were taken and sent to the microbiology deptt of the centre where Gram staining, Culture and corresponding antibiotic sensitivities were performed. Specific antibiotics were administered subsequently along with debridement of the diabetic ulcers as per required.

Result

A total of 101 patients were included in the study over the period of 3 years. The mean age of patients was 57.1 ± 9.1 years. There were more males (64.4%), mostly with type 2 diabetes (99%), with a median duration of 9 years. Their median blood sugar on admission was 246 mg/dL. Five percent of patients died and 23% had a major amputation. Twenty eight germs were isolated in

different cultures. A total of 60.0% of the isolates were Gram-positive bacteria and 40 % were Gram Negative bacteria. *Staphylococcus aureus* was the most common Gram-positive aerobe bacteria, followed by

Enterococcus spp and *Streptococcus* spp. Gram Negative organisms were *Pseudomonas aeruginosa*, *Proteus* spp, *Escherichia coli*, *Enterobacter* spp, and *Klebsiella* spp. Considering the AST data and the comparison established between recommended empiric antibiotics, there are some highlights to address. Excluding MRSA, Gram-positive bacteria represented 45% of the pathogens recorded, and almost all of them were sensitive to amoxicillin/clavulanic acid and piperacillin/tazobactam. Gram-negative bacteria showed a high resistance to amoxicillin-clavulanic acid, fluoroquinolones, and gentamicin. They were susceptible to imipenem, amikacin, and showed moderate susceptibility to third generation cephalosporins.

Conclusion

Gram-positive bacteria are still the main causative pathogens of Diabetic foot infections. Our study highlights piperacillin/tazobactam as the best first-line empirical antibiotic option to treat severe DFI. Along with the antibiotic administration, debridement of the infected diabetic wound is also to be emphasized for better antibiotic action.

Abstract 056

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In Vitro Activity of Aztreonam in Combination with Ceftazidime - Avibactam, Piperacillin - Tazobactam, Cefoperazone – Sulbactam and Amoxicillin – Clavulanate in NDM producing *Escherichia coli*, *Klebsiella pneumoniae* and *Pseudomonas aeruginosa* from Bloodstream Infections

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Background

Multidrug-resistant (MDR) Gram-negative organisms are a major health concern due to lack of effective therapy. Emergence of NDM producing organisms further magnifies the problem. In this context, combination therapy of Aztreonam with other antimicrobials may have potential in treating these pathogens. Unfortunately, there are limited data regarding these combinations.

Methodology

51 NDM producing clinical isolates of *Escherichia coli*, *Klebsiella pneumoniae* and *Pseudomonas aeruginosa* being the potential pathogen in bacterial septicemia were included in this study. The potential for synergy was evaluated via the broth disc elution method. The disk content/CA-MHB ratio (4/8 µg/mL ATM) was purposefully chosen as the ATM concentration would be around the CLSI susceptibility breakpoint for both the *Enterobacteriales* (4 µg/mL) and *P. aeruginosa* (8 µg/mL) strains. After overnight incubation, the tubes were assessed for no growth (susceptible) or growth (not susceptible) based on the breakpoints

Result

According to CLSI breakpoints, Aztreonam susceptibility was fully restored for 50%, 22.7%, 13.6% and 4.5% of the NDM producing *E. coli* isolates when combined with Ceftazidime - Avibactam, Amoxicillin –

Clavulanate, Piperacillin – Tazobactam and Cefoperazone – Sulbactam respectively. In *K. pneumoniae* and *P. aeruginosa*, the Aztreonam + Ceftazidime - Avibactam combination was the most potent in 74% and 100% isolates respectively.

Conclusion

Our findings show that overall, Aztreonam + Ceftazidime - Avibactam was the most potent combination to treat infections caused by NDM producers. Aztreonam in combination with other BL/BLI has potential for therapeutic options in difficult to treat pathogens but further evaluation of is warranted.

Abstract 057

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A Serial Point Prevalence Survey (PPS) of Antimicrobial Use Amongst Admitted Patients in Critical Care Settings of a Tertiary Care Teaching Hospital in Raipur District

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Background

Antimicrobial resistance (AMR) poses a significant global health threat exacerbated by inappropriate antimicrobial use. The WHO predicts AMR could cause up to 10 million deaths annually by 2050, primarily due to misuse. In India, antimicrobial consumption surged by 103% from 2000 to 2015, emphasizing the urgent need for effective antimicrobial stewardship programs (AMSPs). Conducting point prevalence surveys (PPSs) in India's resource-limited settings is challenging, resulting in sparse data on antimicrobial use. This study aimed to establish baseline data on antimicrobial utilization in critical care settings through serial PPSs at AIIMS Raipur, a tertiary care teaching hospital.

Methodology

Over 12 months, data was collected monthly from June 2023 to May 2024 across nine critical care settings. The study included all patients admitted to critical care settings by 8:00 am on survey days, excluding emergency department patients and those on topical or antitubercular antimicrobials. Data on patient demographics, reasons for antimicrobial use, and therapy specifics were collected using a Google Form for Data Abstraction.

Result

A total of 908 admitted patients in nine critical care settings were surveyed in 12 months. Of these, 505 (55.61%) patients were on antimicrobials out of which 320 (63.37%) patients were on two or more antimicrobials. Out of the total antimicrobials prescribed, 79.84% of the antimicrobials were

prescribed for hospital-acquired infections followed by community acquired infections (15.02%). Antibiotic usage was highest in Critical care units (15.7%), Trauma and Emergency ICU (15.05%), and paediatric ICU (14.94%). The antimicrobials used were commonly from WHO "WATCH" category of which Piperacillin-Tazobactam was prescribed most (14.61%) followed by Meropenem and Vancomycin (11.78% each).

Conclusion

This serial PPS demonstrated crucial insights into antimicrobial prescribing patterns, essential for developing targeted AMSP strategies and reducing AMR, thereby improving patient outcomes in critical care facilities.

Abstract 058

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Leveraging Antimicrobial Stewardship in Paediatric Pneumonia: An appraisal of prescription practices at a tertiary care hospital

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Background

Community-acquired pneumonia (CAP) is a leading cause of mortality among children under five years old, particularly in developing countries such as India. Despite the predominance of viral causes for CAP, the overprescription of antibiotics remains prevalent, exacerbating antimicrobial resistance (AMR) and undermining patient care.

Objective: This study aims to evaluate the antimicrobial prescription practices for paediatric CAP in a tertiary care hospital in India. The primary objectives include assessing adherence to hospital antibiotic policies, optimizing antibiotic use, and improving patient outcomes through an antimicrobial stewardship program (AMSP).

Methodology

A prospective observational study was conducted over three months, enrolling 100 patients aged 1 month to 12 years with CAP. Participants were recruited from the paediatric emergency and wards at Chacha Nehru Bal Chikitsalya, New Delhi. Antimicrobial prescriptions were evaluated for compliance with established guidelines. Biomarkers, radiographic patterns, and microbiological tests were utilized to guide treatment and monitor adherence to AMSP protocols. The study also compared clinical outcomes between children receiving stewardship intervention versus those who did not.

Result

The study observed 51% of prescriptions non-adherent

to hospital policy. Implementation of AMSP led to a noticeable reduction in inappropriate antibiotic use,

with increased compliance to first-line treatments and decreased reliance on broad-spectrum antibiotics. The stewardship intervention group showed improved clinical outcomes, including reduced hospital stay duration without an increase in morbidity or mortality.

Conclusion

The findings underscore the critical need for AMSP in paediatric settings to mitigate the adverse impacts of AMR. Adherence to antibiotic guidelines can significantly enhance the quality of care for children with CAP, fostering responsible antimicrobial use and better health outcomes. This study highlights the importance of continuous education and monitoring to ensure the success of stewardship programs in improving paediatric health care.

Abstract 059

Real World Evidence on the Impact of Implementation of Antimicrobial Stewardship (AMS) Practices on Antimicrobial Resistance and Therapeutic Outcomes on a Tertiary Care Hospital of West Bengal

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Background

The study was conducted to gather information to evaluate the extent of impact of antimicrobial stewardship (AMSP) practices on antimicrobial prescribing patterns and bacterial resistance profiles among indoor patients across ten ICU and non-ICU departments at the Institute of Postgraduate Medical Education and Research, Kolkata.

Methodology

The study protocol was submitted to the Institutional Ethics Committee of the institute and it was approved unanimously. Data from indoor patients were collected prospectively from June, 2022 till October, 2023. The use of antimicrobials was categorized as empiric, prophylactic or targeted/culture driven. The WHO-AWARE classification of antimicrobials was used to classify systemic antimicrobials being prescribed to patients. Study data was archived and analyzed using software such as SPSS 20.0 (IBM), Jamovi and Med Calc 8 (Belgium).

Result

Total number of beds covered annually was 2972 and the number of patients on antimicrobials was found to be as high as 98.57 %. Relatively higher number of patients were found to be on two or more antimicrobials predominantly from WATCH and RESERVE categories. Interestingly, among nosocomial CLABSI and CAUTI rates were 22.05 % and 22.11 % respectively with a relatively lower proportion of SSIs 0.88%. Mortality relatedness to infection was found to be 48.62 %. The overall compliance rate to hospital antibiotic policy was found to be 72. 51% while culture

was sent in nearly 59% of the patients surveyed. Rationality assessment and prospective audit and feedback of antimicrobials was performed which is under analysis.

Conclusion

This preliminary study resulted in acquisition of data not only related to resistance profiles of microbial pathogens to antimicrobials but also yielded data about determinants of hospital acquired infection including surgical site infections which may aid policy makers to formulate newer antimicrobial guidelines in the state and country based on such definitive findings.

Abstract 060

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Trends of Antimicrobial Resistance among Blood Isolates at a Tertiary Care Centre in North India: A Retrospective Six-Year Study

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Introduction

Bloodstream infections are important causes of morbidity and mortality worldwide. Candidemia is also being frequently reported and is mostly associated with immune suppression, prolonged antibiotic usage, invasive surgical procedures, intravenous devices and parenteral nutrition. The rising rates of antimicrobial resistance among pathogens in India are alarming, because of the high infection rates and poor control of antibiotic use thus antimicrobial surveillance is essential for identifying emerging resistance.

Aim: To study trends of antimicrobial resistance among blood isolates.

Materials & Methodology

This retrospective, observational study was performed in the Microbiology laboratory of a tertiary care hospital from 2018 to 2023. Blood samples from patients with suspected BSI were cultured on automated systems (BD Bactec Fx/ BactAlert 3D). Identification and antimicrobial susceptibility testing were performed on the Vitek2 system. The data was retrieved from the laboratory records and antimicrobial resistance profiles were analysed.

Result

Overall blood culture positivity of pathogenic bacteria and *Candida* spp was 12% (varying from 11.8% in 2018 to 11.5% in 2023), while year-wise positivity rates of candidemia was 0.68%, 0.73%, 0.82%, 0.71%, 0.50% and 0.55% from 2018 to 2023 respectively. *Escherichia coli* (21.5%) and *Klebsiella pneumoniae* (18.2%) were the predominant isolates, followed by *Acinetobacter baumannii* (8.9%), *Staphylococcus aureus* (8.3%) and

Candida spp (5.8%). Among *Candida* isolates, *C. tropicalis* was the most common followed by *C. albicans* and *C. parapsilosis*. The Gram-negative isolates showed high rates of resistance to third generation cephalosporins, β -lactam- β -lactamase inhibitor combinations and carbapenems. Methicillin-resistant *S. aureus* isolates increased from 67% to 83% over the six-year period, while vancomycin-resistant enterococci also escalated from 31% to 40% in 2022, with a fall to 22% in 2023. In comparison, *Candida* isolates showed good susceptibility to amphotericin B & echinocandins whereas increased resistance to azoles (10-25%) was observed in *C. tropicalis* and *C. parapsilosis*.

Conclusion

The study revealed that the majority of bloodstream infections were caused by Gram-negative pathogens with high rates of resistance to most antimicrobials and predominance of non-*albicans* *Candida* species. Regular monitoring of local trends of antibiotic resistance detects early shifts in susceptibility and also serves as a basis for empirical therapy, formulary decisions, and infection control practices.

Abstract 061

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Emergence of Multidrug-Resistant *Klebsiella pneumoniae*: Antimicrobial Susceptibility Trends (2019-2023) in an Indian Tertiary Care Hospital

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Background

The rise of multidrug-resistant (MDR) *Klebsiella pneumoniae*, causing nosocomial infections and significant mortality, has made it a WHO “priority pathogen.” This study examines the antimicrobial susceptibility of *K. pneumoniae* from pus specimens in an Indian tertiary care hospital, identifying trends in MDR spread.

Materials & Methodology

Clinically relevant isolates from culture-positive pus samples were identified using MALDI-TOF, and antimicrobial susceptibility testing (AST) was performed via Vitek2 systems. Colistin resistance was assessed using microbroth dilution as per CLSI/EUCAST guidelines. Data management and statistical analysis were conducted using WHONET and GraphPad Prism, respectively. Categorical variables were analyzed using χ^2 or Fisher exact test, with Odds Ratios (ORs) and 95% Confidence Intervals (CIs) calculated.

Result

From 2019-2023, 5801 *K. pneumoniae* isolates (13.1%) were identified from 44,414 positive pus specimens. The prevalence of *K. pneumoniae* increased by 14% in 2023 compared to 2019 ($p < 0.05$). AST showed a significant rise in carbapenem resistance, with 9.2% of isolates resistant to both carbapenems and colistin ($p < 0.0001$). MDR *K. pneumoniae* was more prevalent among inpatients compared to outpatients ($p < 0.0001$, OR: 1.817, CI: 0.7945 to 2.079). A moderate positive association was found between *K. pneumoniae* isolates from Medicine wards and MDR cases ($p < 0.0001$, OR: 1.583, CI: 0.7014 to 1.786). Resistance increased significantly post-COVID-19.

Conclusion

The study revealed concerning trends in *Klebsiella pneumoniae* antimicrobial resistance at an Indian tertiary care hospital, with a 14% surge in prevalence by 2023. Notably, 9.2% of isolates were resistant to carbapenems and colistin. Inpatients had significantly higher rates of multidrug-resistant *K. pneumoniae*, highlighting the urgent need for enhanced infection control measures.

Figure 1: Increase in %R for carbapenem and colistin post COVID-19.

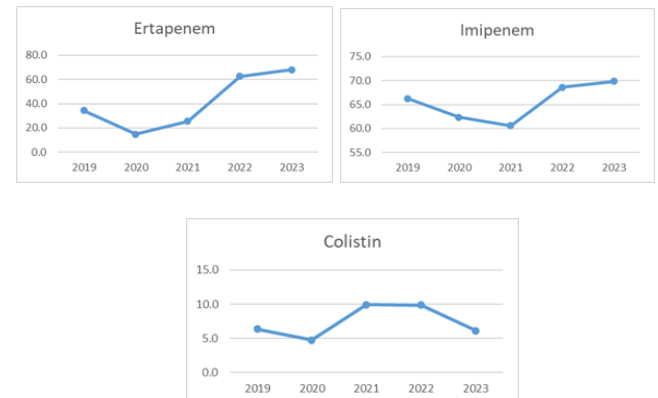
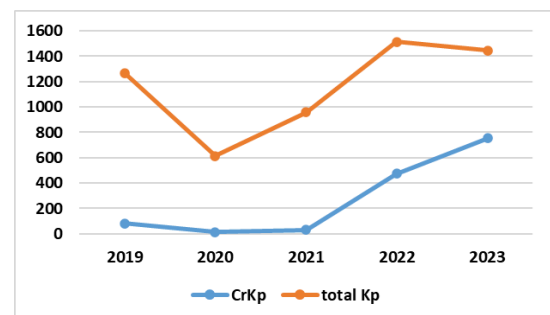


Figure 2: Increase in Carbapenem resistance *K. pneumoniae* isolates.



Abstract 062

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To Assess Symptomatic Improvement with Prescribed Antibiotics in Uncomplicated Urinary Tract Infections (UTIs)

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Introduction

Antibiotics are among the most frequently prescribed medications for outpatients suffering from uncomplicated urinary tract infections.

Aims and Objectives: To assess symptomatic improvement with prescribed antibiotics in uncomplicated urinary tract infections.

Materials & Methodology

This study was conducted on 60 male patients diagnosed with uncomplicated UTI. It was an observational, follow up study. Inclusion criteria was male patients age >16 years, diagnosed as case of uncomplicated UTI based on symptoms suggestive of UTI including burning micturition, increased frequency of urination, urgency, suprapubic pain, blood in urine, cloudy urine, fever with or without chills. Exclusion Criteria was catheter associated UTI, complicated UTI cases due to stones, strictures, neurogenic bladder, recurrent UTI and previous history of surgery involving genitourinary tract.

In this study, we found that Levofloxacin and Nitrofurantoin are the two most commonly prescribed antibiotics for uncomplicated UTI patients. Nitrofurantoin 100 mg BD provides symptomatic relief earlier than other antibiotics which make it a better choice for such patients.

Result

Abstract 063

Antibiotic	Dose	Duration of therapy in days	No of Prescriptions (%)	Symptomatic Relief in no of patients (%)	Symptomatic Relief in number of days (Mean±S.D.)
Levofloxacin	500 mg OD	7	19 (31.66%)	5 (26.31%)	5±0.63
Levofloxacin	750 mg OD	5	6 (10%)	1 (16.66%)	4.5±1
Ofloxacin	100 mg OD	5	4 (6.66%)	4 (100%)	5
Ciprofloxacin	500 mg OD	5	4 (6.66%)	2 (50%)	5±0.63
Norfloxacin	400 mg OD	7	9 (15%)	9 (100%)	3
Nitrofurantoin	100 mg OD	7	18 (30%)	14 (77.77%)	4.2±2.11

Interpretation

The Novel Automated Multiplex Pcr Respiratory Panel a Boon or Bane? For Stewardship Practices, An Experience from Dedicated Liver Hospital in North India

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Background

Ventilator associated pneumonia [VAP] is second most common HAI in ICU settings. The diagnosis of VAP is complex and requires correlation between Clinical, radiological and Microbiological criteria. The CDC supports semi quantitative reporting for respiratory samples sent for culture to rule out colonization. The novel multiplex PCR pneumonia panel has a TAT of 1 hour which when compared to Culture of respiratory specimens require minimum 48 hrs. The role of respiratory sample culture in stewardship practices is well known because of CDC semi quantitative threshold values but role of This Novel multiplex Respiratory PCR is not well described in studies and need further evaluation, secondly the Automated PCR gives its final result in unit Bin copies/ml which is totally different from culture result unit CFU/ml leading to confusion for treating physician in significance of pathogen as a coloniser or as a true pathogen

Materials & Methodology

- **Study Type :** Retrospective observational study
- **Study period :** September 2020 – Dec2020
- **Study population :** Liver disease patients of all age groups with clinically suspected lower respiratory tract bacterial infection admitted to Liver CICU
- **Sample size :** 157

- All respiratory samples received in microbiology lab were subjected to culture as well as PCR and results were compared at last

Result

- Majority of 10⁵ culture positive isolates were identified by PNP as 10⁷ and 10⁶ bins
- Both PNP and culture were +ve in 19.1% cases
- Both PNP and culture were –ve in 51% cases
- Genotypic resistance detected by PNP matched Phenotypic resistance detected by VITEK in 100% cases in case of gram neg bacteria
- MeC-A/C gene were detected by PNP in all cases of MRSA detection by VITEK [PHENOTYPIC]
- Additional 28.6% gain was seen in diagnosis of bacterial isolates which were negative on culture

Conclusion

Novel multiplex respiratory PCR is a great tool for early detection of MDR pathogen causing pneumonia and aids in stewardship practices

Abstract 064

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Evaluating the Efficacy of Ceftazidime-Avibactam and Aztreonam synergistic combination among Carbapenem Resistant *E. coli* and *Klebsiella spp.*

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Introduction

Carbapenems are preferred over other types of antimicrobial agents in treating invasive or severe infections due to their concentration-independent killing effect on the infecting bacteria. Carbapenem resistance in Gram-negative bacteria has been increasingly encountered in healthcare-associated infections in India. Ceftazidime-avibactam (CZA) and Aztreonam (ATM) combination therapy is used specifically to target infections caused by Metallo-beta-lactamase (MBL) producing Carbapenem-resistant Enterobacteriaceae (CRE). When combined, Ceftazidime-avibactam protects Aztreonam from hydrolysis by non-MBL beta lactamases. Aztreonam remains active against the MBLs, providing a synergistic effect. This study aimed to evaluate the synergistic effect of Ceftazidime-avibactam and Aztreonam combination in Carbapenem resistant Enterobacteriaceae.

Materials & Methodology

Carbapenem – resistant Enterobacterales were included in a study conducted in the Department of Microbiology, S.N. Medical College, Agra. The isolates were collected from different sources such as urine, blood, pus, respiratory specimens like sputum, bronchoalveolar lavage fluid and other body fluids. Samples received were processed by conventional methods for identification. Gram-negative isolates obtained were subjected to antimicrobial susceptibility testing by the Kirby–Bauer disc diffusion method using the following discs amikacin, ceftazidime, cefepime, ciprofloxacin, imipenem, cefoperazone/sulbactam, piperacillin/tazobactam, co-trimoxazole, ceftazidime/avibactam and aztreonam.

Result

Out of 293 Carbapenem-resistant isolates, 71.33% are *Klebsiella spp* and 28.66% are *E. coli*. Synergy test among *E. coli* and *Klebsiella spp* is under process. Out of 293 Carbapenem resistant Enterobacteriaceae studied, 78% isolates showed resistance to piperacillin-tazobactam, 47% isolates are resistant to amikacin, 55% isolates are resistant to cotrimoxazole.

Conclusion

The use of Ceftazidime-avibactam together with Aztreonam is a potentially suitable combination for the treatment of Carbapenemase-producing microorganisms and is available in clinical laboratories. As the combination has less side effects so the in vivo efficacy of this combination needs to be evaluated.

Abstract 065

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Emergence of Multidrug-Resistant Non-fermenting Gram Negative Bacilli and Carbapenem Resistant *Acinetobacter baumannii* (CRAB) in a Tertiary Care Hospital of Northern India

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Background and Objective

Nonfermenting gram negative bacilli (NFGNB) are now emerging as important nosocomial pathogens causing hospital-acquired infections and opportunistic infections. This study aimed to determine the antibiotic susceptibility pattern of NF-GNB isolated from all clinical samples to estimate the prevalence of Multidrug resistant NF-GNB and to screen for Carbapenem resistant *Acinetobacter baumannii* (CRAB) strains.

Materials & Methodology

An observational study with a cross-sectional design was conducted between September 2023 to May 2024 in the Department of Microbiology, S.N. Medical College Agra. We identified MDR NF-GNB from clinical samples using standard procedures and antimicrobial susceptibility testing done by Kirby Bauer disc diffusion method and MBL detection by EDTA double disc synergy test by using commercially available discs (Hi-media) i.e. Amoxicillin/clavulanic acid (20/10mcg), Piperacillin, (100mcg), Piperacillin/tazobactam (100/10mcg), Ceftazidime (30mcg), Cefepime(30mcg), Trimethoprim/sulfamethoxazole (1.25/23.75mcg), Ciprofloxacin(5mcg), Levofloxacin (5mcg), Amikacin (30mcg), Meropenem (10mcg), Imipenem (10mcg), Polymyxin B (300 mcg), Doxycycline (30mcg), Co-trimoxazole (25mcg), Tobramycin (10mcg), Minocycline (30mcg), Colistin (10mcg), Tigecycline (15mcg).

Result

A total of 270 MDR-NFGNB were isolated in 700 non fermenters gram negative bacilli and most common

samples were blood (n= 90, 33.3%) followed by pus (n=74 ,27.4%). Among MDR-NFGNB the most common organisms were *Pseudomonas aeruginosa* (n=126,46.6%), followed by *Burkholderia cepacia* complex (n=68,25.1%), *Acinetobacter baumannii* (n=67,24.8%) and others. The susceptibility pattern of these isolates were piperacillin-tazobactam (45.1%), imipenem (39.2%), aztreonam (18.5%), amikacin (17.4%), ceftazidime (14.0%), ciprofloxacin (10.7%), doxycycline (8.8%), cefepime (7.4%), cotrimoxazole (5.5%).Maximum MBL production was obtained in *Acinetobacter* spp. followed by *Pseudomonas aeruginosa* and the Carbapenem resistant *Acinetobacter baumannii* were (n=42,55%).

Conclusion

Since these organisms have great potential to survive in the hospital environment, improved antibiotic stewardship and infection control measures are required to prevent the emergence and spread of drug-resistant NFGNB in healthcare settings.

Abstract 066

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From Blueprints to Reality: Journey of a Clinical Pharmacologist Post-3rd ASPICON

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Submit a Manuscript: <https://journal.saspi.in/>
SASPI: <https://saspi.in/>

ISSN: 3048-4510 (Online)
DOI: 10.62541/jaspi2024

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Introduction

Antimicrobial resistance (AMR) poses a significant threat to global public health, largely driven by the misuse and overuse of antibiotics. Antimicrobial Stewardship (AMS) programs aim to optimize antibiotic use, enhance patient outcomes, and reduce AMR. This study details the experiences of a clinical pharmacologist in implementing AMS interventions in two tertiary care hospitals, focusing on the strategies employed and their impact on antibiotic use and patient outcomes.

Methodology

The AMS program, led by a clinical pharmacologist, was implemented over a one-year period at two different tertiary care hospitals. Key interventions included: a) PostPrescription Review and Feedback: Reviewing ongoing antibiotic therapies and providing recommendations based on patient progress and microbiological data. b) Educational Initiatives: Conducting workshops, case discussions, and disseminating guidelines to promote judicious antibiotic use. c) Multidisciplinary Collaboration: Participating in rounds and collaborating with infectious disease specialists, microbiologists, and nursing staff. Data on antibiotic usage, compliance with guidelines, incidence of *C. difficile* infections, and patient outcomes were collected and analyzed.

Result

AMS interventions led to a 20% decrease in continued use of broad-spectrum antibiotics and an increase in de-escalation practices. Compliance with evidence-based guidelines increased from 60% to 85%.

Additionally, the average hospital stay was reduced by 2 days, with a corresponding decrease in antibiotic-related adverse events.

Conclusion

AMS interventions, including post-prescription review, educational initiatives, and multidisciplinary collaboration, significantly optimized antibiotic use and improved patient outcomes. Continuous monitoring and collaboration are essential for sustaining AMS benefits and addressing emerging challenges.

Abstract 067

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Prevalence of Dengue and Leptospirosis Co-infection at Tertiary Care Hospital, Mumbai

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Background

Dengue and Leptospirosis are both infectious diseases and thus pose a threat to public health, especially in tropical regions like Mumbai. During monsoon, the incidence of the infections is widespread. Due to concurrent transmission, the possibility of co-infection may increase. The severity of the infection is heightened because of the concomitant infection. Despite this, there are not many studies available to address the severity of co-infection.

Aim: To determine the prevalence of Dengue and Leptospirosis co-infection, by Dengue NS1 antigen and IgM, Leptospirosis IgM ELISA.

Methodology

Retrospective observational laboratory-based study has been conducted over a period of 6 months in tertiary care hospital, Mumbai. A total of 3372 serum samples were received from patients with acute febrile illness and examined for dengue NS1 and IgM. Positive Dengue samples were additionally tested for IgM antibody against Leptospirosis by ELISA and co-infection was diagnosed.

Result

Out of 3372 received serum samples, 418 (12.39%) found positive for Dengue and among these 86 were co-infected with prevalence of 2.55 %. 19 cases of co-infection reported in the month of October and November.

Conclusion

These findings added a new dimension in the management of Dengue and Leptospirosis. The possibility of these concurrent infections should be considered especially in endemic areas for early diagnosis and proper treatment, thereby decreasing the mortality.

Abstract 068

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A Quality Improvement Initiative to Improve the Implementation Fidelity to Institutional Surgical Antibiotic Prophylaxis Policy

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Background

A Quality Improvement (QI) initiative aimed at enhancing infection control and reducing Surgical Site Infections (SSI) was undertaken at our institute. A significant component of this initiative focused on improving compliance with Surgical Antibiotic Prophylaxis (SAP) guidelines due to observed non-uniform practices across surgical departments.

Methodology

The QI project employed the Plan-Do-Study-Act (PDSA) methodology to systematically address SAP compliance. Initial steps included a gap analysis conducted by the Infection Prevention and Control (IPC) and Antimicrobial Stewardship Program (AMSP) teams, leading to the development and dissemination of institutional SAP guidelines. Data was collected on SAP practices, specifically focusing on the choice, dose, and timing of antibiotic administration, and compliance was audited monthly. The project aimed to increase SAP compliance to over 80% by April 2024, using targeted education, training, and continuous feedback mechanisms.

Result

The initial audit in July 2023 revealed varied compliance with SAP guidelines, with significant deviations in antibiotic choice, dose, and timing. Following the implementation of the QI interventions, compliance rates improved notably by April 2024. The proportion of surgeries with the correct antibiotic choice increased from 75% to 100%, the correct dose from 75% to 100%, and the correct timing from 50% to 100%. Correspondingly, the rate of SSIs decreased from 7.8% in November 2022 to 3.9% by April 2024, demonstrating the effectiveness of the QI initiative.

Conclusion

The QI project successfully improved compliance with SAP guidelines through structured interventions, significantly reducing SSI rates. Continuous monitoring, feedback, and iterative adjustments to the implementation strategies were crucial to achieving these outcomes. The initiative underscores the importance of a collaborative & multidisciplinary approach in driving quality improvements in surgical practices and infection control. Future steps will focus on sustaining high compliance levels and further reducing postoperative antibiotic use in non justified cases.

Abstract 069

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Case of Multifocal Abscesses in Uncontrolled Diabetes

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Abstract

Burkholderia cepacia is a pathogen usually causing infection in immunocompromised, uncontrolled Diabetes. It is also associated with infections in patients with underlying lung disease, such as cystic fibrosis and chronic granulomatous disease. *Burkholderia* infection mimics tuberculosis closely due to a similar clinical profile.

We present a case of disseminated *B. cepacia* infection in a 42 years old male patient with newly detected Uncontrolled type 2 diabetes mellitus and ischemic heart disease.

Patient was symptomatic for 1 month with intermittent low grade fever, chest discomfort and intermittent episodes of diaphoresis and palpitations. He also complained of burning micturition for 15 days, associated with incomplete voiding. He had a well defined, fleshy ulcerated lesion with purulent discharge over his left tibial tuberosity. Investigations were suggestive of neutrophilic leukocytosis anemia of chronic disease, mild transaminitis, HbA1c of 12.1% and elevated inflammatory markers – ESR, CRP, Ferritin and Procalcitonin was also elevated. Urine routine examination had 25-30 pus cells/hpf, albeit the cultures were sterile on serial 3 samples. USG revealed splenic abscess with prostatomegaly, hence CT Thorax and Abdomen-pelvis was done and was suggestive of Splenic, Prostatic abscess and loculated, septated collections in anterior and middle mediastinum encasing the ascending aorta, arch of aorta, the SVC and right atrium. There were also multiple subcentimeter mesenteric, retroperitoneal, inguinal and mediastinal reactive Lymph nodes.

CT guided Pus was drained from prostatic abscess, and pus from the ulcer revealed the growth of *Burkholderia cepacia* sensitive to Meropenem and cotrimoxazole which were started as per the indicated preferred regimen for 4 weeks to which the patient dramatically

responded. Cotrimoxazole was further continued for 3 months and the patient has clinicoradiological resolution on follow up.

Burkholderia is a great mimic of tuberculosis especially in endemic areas. It affects the immunocompromised individuals, and also specifically in poorly controlled Diabetes Mellitus and can cause multitude of clinical manifestations including disseminated abscesses as was in our case. Clinical suspicion of Melioidosis can prompt early diagnosis and treatment.

Abstract 070

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The Crucial but Unrecognized Role of Nurses in Antimicrobial Stewardship: A Systematic Review on the Challenges and Opportunities

Asheema Khosla*

Introduction

The widespread misuse of antimicrobials has led to the development of resistance in bacteria, viruses, fungi and parasites. Antimicrobial resistance is a worldwide concern which has contributed to national and global health crises, and with the global threat of AMR emergence there should be wider involvement between the multidisciplinary healthcare team. This article proposes possible ways of engagement of Nurses as Antimicrobial Stewards.

Objective: The aim of this evaluation is to determine the challenges and opportunities for the nurse role in the antimicrobial stewardship.

Methodology

PubMed/MEDLINE, EMBASE and Web of Science were searched for articles in English. The search strategy included indexing (MeSH) terms and text words for nurses, antimicrobial stewardship, antimicrobial resistance, antimicrobial stewardship programme and training and curriculum. Internet search engines were used to search the grey literature for reports, qualitative studies, cross-sectional studies and Randomized controlled trials. Rayyan® was used to organize and extract data.

Result

Out of 618 initial hits, 61 articles were included for data extraction. 219 duplicates were removed. Majority of the studies were Surveys (44%) followed by Qualitative studies (21%), Cross-Sectional studies (5%) and RCTs (1%). Studies were included from both LIMCs (46%) and HICs (53%) specifically from the Europe region. AMS challenges should be addressed by collaborative and methodical approach, aiming to implement AMSP in an effective manner.

The studies have attempted to evaluate the impact of educational initiatives on AMS practices for nursing students. It is evident from various studies that their curriculum lacks Education and proper training on AMS. Numerous surveys, interviews were conducted to assess the knowledge, attitude and perceptions of nursing students, clinical nurses and staff nurses. They were aware of their deficiency, the majority affirmed that the current curriculum of nursing degree should have more training on antibiotics and infection control because nursing staff plays a critical role in the rational use of antibiotics, so their training can be a key in fighting the antibiotic resistance. Furthermore, Inter-professional hierarchy and lack of cooperation by the prescribers was a major area of concern in the implementation of AMS. A number of potential interventions are identified that may prevent or mitigate these problems. Aside from preparing, administering and monitoring the effects of antimicrobials, nurses are involved in coordinating care, clinical assessment, taking specimens, monitoring patient progress and reporting adverse events, all of which offer opportunities to influence antimicrobial prescribing.

Conclusion

The inclusion of Nurses in Antimicrobial Stewardship is pivotal and through proper education and training they can have significant impact on the antimicrobial resistance patterns and can contribute in reducing the AMR burden.

Abstract 071

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When a Bad Hand-rub Choice Led to an Outbreak of *Candida Pelliculosa* in the ICU

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Background

The importance of hand hygiene cannot be over-emphasized. Standard alcohol-based handrubs (ABH) contain approx. 2% chlorhexidine and 60-80% alcohol. Some brands also offer emollient/ moisturizer-blended handrubs, which are softer on hands but may compromise the recommended % of constituents.

Methodology

Automated blood cultures with budding yeasts (BYCs) were identified on MALDI-TOF-MS (Biomerieux) and antifungal susceptibility was performed. A team of Microbiologists and infection control nurses conducted surveillance cultures and IPC audits, including evaluation of ABHs used in the ICU.

Result

The study was conducted from February to April 2022, when an unexpected 77 isolates of *Candida pelliculosa* were cultured from febrile HAI patients, susceptible to Caspofungin (84.41%), Amphotericin B (80.51%), Flucytosine (76.62%), Voriconazole (74.02%), Fluconazole (48.05%) and Micafungin (5.19%). Our investigation found the common link of all cases as the adult ICU. While surveillance cultures did not yield any *Candida pelliculosa*, an audit of ABHs showed a non-approved emollient ABH product with <60% alcohol being used for skin preparation, despite the availability of the standard ABH product in the store. There had been a routine shuffling of junior nursing staff between the wards and the ICU, around two months ago.

The new staff had switched to the emollient-based product for hand hygiene and had inadvertently used it for skin preparation as well. The use of the new product was immediately discontinued. A proper training was held and all the staff were re-trained. The

Candida pelliculosa outbreak resolved within 10-12 days of these active interventions. Routine surveillance was strengthened.

Conclusion

We report an outbreak of *Candida pelliculosa* in the adult ICU of a tertiary care hospital in Western Rajasthan, which was probably caused by the use of a new inappropriate ABH product. Prompt identification, proper training and re-introduction of the appropriate ABH led to an immediate cessation of the outbreak. The use of hand rubs with unregulated constituents can compromise infection prevention and control (IPC) in the hospital.

Abstract 072

ASPICON 2024 | August 22 -25 | 2024

Comparison of Population Pharmacokinetics of Two Different Dosing Regimens of Vancomycin in Critically ill Patients with Sepsis

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Background

Sepsis management requires urgent and optimized antimicrobial use. In sepsis, there is alteration of the pharmacokinetics of the drugs on account of several pathophysiological changes. So, pharmacokinetic studies to optimize antimicrobial dose and improve probability of target attainment need to be conducted. Vancomycin is a key component of empiric and/or lab-based use in patients with sepsis. The aim of present study was to compare two intravenous dosing regimens 1000 mg bid and 500 mg qid in critically ill patients with sepsis and to validate nomogram generated from pop-pk-pd modeling in a sample cohort.

Methodology

The study was conducted by the Department of Pharmacology in collaboration with critical care units in a tertiary care center of North India after obtaining approval from IEC (No. INT/IEC/SPL-480) and registration with CTRI (CTRI/2023/02/049473). Eligible patients were randomly assigned to receive either the vancomycin dose routinely administered 1000 mg iv twice daily or 500 mg iv four times daily. Relevant covariate data were collected. Sparse sampling methodology with capture of trough concentrations was carried out. Structural model followed by covariate model was built using PUMAS software. Model diagnostics conducted include goodness-of-fit, visual predictive check and 1000 bootstraps to validate the model. Data simulation for 1000 mg bid and 500 mg qid were carried out for different weight and creatinine clearance combinations ranging from 45-80 kg and 50-100 ml/min. Key primary endpoint was an increased probability of target attainment.

Result

A total of 26 patients were randomized to the two treatment arms. A 2-compartment combined error model with the covariates creatinine clearance and body weight explained the data adequately. AUC₀₋₂₄/MIC estimation from simulated data showed that the number of individuals attaining the target was significantly higher in the 500 mg qid group as compared to 1000 mg bid group. Nomogram based dosing and evaluation is ongoing. The trough concentrations in the therapeutic range of 15-20 mg/L was observed for a higher proportion of patients in 500 mg qid as compared to 1000 mg bid across all creatinine clearance and weight ranges tested.

Conclusion

The results of the study indicate that a higher proportion of patients achieve the required target in 500 mg iv qid group and it may be used in therapeutics to increase clinical efficacy.

Abstract 073

ASPICON 2024 | August 22 -25 | 2024

Clinical Efficacy and Pharmacokinetics of Colistimethate Sodium in Critically ill Paediatric

Patients Admitted in a Tertiary Care Hospital in India : A Prospective Longitudinal Observational Pilot Study

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Background

The emergence of multidrug-resistant Gram-negative infections in India's paediatric intensive care units (PICU) has necessitated the renewed use of colistin. Current evidence on colistin's use in Indian paediatric patients is limited, underscoring the need for further research. This study uniquely evaluates the clinical efficacy and safety of colistimethate sodium (CMS) in critically ill paediatric patients, correlating it with pharmacokinetics.

Methodology

: Children meeting the inclusion criteria were enrolled from the PICU of a tertiary care centre after obtaining parental consent. Demographic details were recorded, and blood samples were analysed for kinetic parameters using HPLC. Clinical efficacy indicators included clinical improvement or treatment failure at 72 hours, clinical cure at 14 days, all cause mortality within 28 days, ventilator days, length of PICU stay, and any adverse reactions before discharge. The microbiological outcome was culture response rate at the end of 14 days of CMS therapy. Data were managed with MS Excel (2019) and analysed using SPSS.

Result

The study included 23 patients. The main indication for colistin was sepsis (52.2%). At 72 hours, 52.2% of patients showed clinical improvement, while 47.8% had

treatment failure. Clinical cure at 14 days was observed in 43.5% of patients. All-cause mortality within 28 days was 43.5%. The median PICU stay was 21 days (IQR: 13-30), and the median ventilator days were 15 (IQR: 8-21). Nephrotoxicity occurred in 6 patients (23.1%), with no cases of neurotoxicity. Culture response was observed in 61.5% of patients. Kinetic parameter results are awaited.

Conclusion

Colistin therapy, administered as loading and maintenance doses, appears to be effective and safe in the paediatric population based on clinical and microbiological outcomes.

Abstract 074

ASPICON 2024 | August 22 -25 | 2024

Prospective Assessment of Antimicrobial Prescribing Pattern in Medicine Intensive Care Unit of a Teaching Hospital

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Background

Antimicrobial agents (AMAs) are one of the greatest contributions of the 20th century, used for the treatment and prevention of infections. However, their widespread use has led to the development of resistance. Understanding the pattern of their use is crucial for optimizing treatment efficacy and curbing the emergence of antibiotic-resistant organisms. Hence, it is important to evaluate the prescription pattern of AMAs to avoid irrational use of drugs.

Objectives:

1. To study the prescribing patterns of antimicrobial agents.
2. To Analyse the rational use of antimicrobial agents in the ICU

Methodology

The study was a retrospective observational study conducted over a period of three months, from January 2024 to March 2024. A total of 216 case records of patients admitted to medical intensive care unit, receiving at least one dose of antimicrobial drugs, were reviewed for the pattern and rationality of antimicrobial drug usage. Rationality was assessed by analysing adherence of antimicrobial prescriptions to Institutional Antimicrobial policy. Data were summarised and analysed using mean and percentage values.

Result

Of the total 216 patients evaluated, 65.7% were male and 34.2% were female. A common indication of AMAs use was Acute Febrile illness (AFI) at 46.3%, followed by COPD at 23.1%. The most commonly prescribed

AMAs were cephalosporins (55.56%), followed by tetracyclines (27.78%), and combination of antimicrobials from different groups (16.67 %). In 62.5% of patients, antimicrobial therapy was considered rational.

Conclusion

Prescribers did not adhere to the Institutional Antimicrobial policy in more than one - quarter of the prescriptions. This non-adherence may increase the risk of treatment failure, adverse drug reactions and the development of antibiotic resistance.

Comparative Evaluation of Agar Dilution, Broth Microdilution, and Vitek-2 Methods for Detection of Colistin Resistance in *Escherichia Coli* Urine Isolates

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Introduction

Polymyxin (Colistin) is a novel class of antimicrobial that has a surface-active detergent-like action against most Gram-negative bacteria like *Escherichia coli*. The emergence of colistin-resistant bacteria, although reported infrequently to date, However, there is still no consensus regarding the breakpoints for outlining resistance to colistin.

Aim: To compare the agar dilution, broth microdilution method, and Vitek-2 for the detection of colistin resistance in *E. coli* urine isolates.

Materials & Methodology

A one-year cross-sectional observational study was conducted in the Department of Microbiology at Maulana Azad Medical College, New Delhi. A total of 200 *E. coli* isolates were subjected to Agar dilution, Broth microdilution (BMD), and Vitek-2 methods for colistin. Using BMD as the gold standard, a comparative analysis between different methods was carried out.

Result

Out of 200 *E. coli* isolates, Vitek-2 detected colistin resistance in six isolates. By Broth microdilution, two isolates were resistant to colistin. No isolates were resistant to colistin by the agar dilution method. Among the 198 isolates that were colistin-sensitive by the broth microdilution method, 4 were colistin-resistant by Vitek-2. All the isolates were subjected to PCR using primers targeting the *mcr* gene to compare the resistance pattern of colistin in *E. coli* isolates. Out of 200 *E. coli* isolates, 20 had the *mcr* gene.

Conclusion

The overall concordance rate was 1%, and the discordance rate was 99% between Colistin resistance from BMD and Vitek-2. Therefore, the use of molecular methods (PCR) needs to be increased for better detection of resistance marker genes in routine laboratory settings.

Incidences of Missed Mucormycosis in Immunocompetent Individuals: A Study of Diagnostic Challenges and Treatment Patterns in Tbilisi, Georgia

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Background

Mucormycosis, typically seen in immunocompromised individuals, is a rare and diagnostically challenging fungal infection. However, recent reports suggest an increasing incidence among healthy, immunocompetent individuals. This research paper aims to highlight cases of missed mucormycosis in healthy patients, emphasizing the diagnostic challenges and efficacy of treatments post-misdiagnosis. We conducted a primary research study involving 15 respondents in Tbilisi, Georgia, and reviewed past cases of pulmonary and renal mucormycosis to understand evolving incidence patterns and diagnostic approaches. Two major hospitals in Tbilisi, where these patients were treated, were also included in the study.

Methodology

This study employed a mixed-methods approach. We conducted primary research with 15 respondents residing in Tbilisi, Georgia, to gather data on missed mucormycosis diagnoses and treatment patterns. The survey included questions about their health status, medical history, and any instances of mucormycosis diagnosis. Additionally, we reviewed case studies focusing on missed mucormycosis infections, misdiagnoses, and treatment outcomes. Diagnostic and differential diagnostic criteria were analyzed alongside novel treatment guidelines for the infection. Data from two major hospitals in Tbilisi, Georgia – Tbilisi State Medical University Clinic and Neo Hospitals – were included to provide clinical insights and case management details.

Result

Our primary research indicated that among the respondents, a significant proportion reported

generalized symptoms that were initially misdiagnosed, leading to delayed treatment. Radiological findings often revealed signs indicative of mucormycosis, such as pleural effusion and multiple nodules, which were overlooked in initial assessments. The review of past cases corroborated these findings, highlighting the need for improved diagnostic criteria and awareness among healthcare providers. Both Tbilisi State Medical University Clinic and Neo Hospitals documented several cases where initial misdiagnosis led to complications, emphasizing the necessity for comprehensive diagnostic protocols.

Conclusion

Although mucormycosis presentation in healthy, immunocompetent individuals is rare, the incidence rate is increasing, necessitating vigilant differential diagnosis and prophylactic testing. Early detection and treatment initiation are crucial, as delays in such cases can prove fatal. The study underscores the importance of heightened awareness and prompt action to mitigate the risks associated with missed mucormycosis diagnosis.

Diagnostic Accuracy of Point-of-Care Tests for Urinary Tract Infections: A Systematic Review and Meta-Analysis Protocol

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Background

Urinary tract infections (UTIs) are a common health problem that leads to a considerable number of consultations and antibiotic prescriptions. The conventional diagnosis is still based on empirical methods dependent on symptoms reported by the patient and urinalysis performed using test strips, with urine culture as the gold standard despite its result turnaround of 48 hours.

Objectives: It is particularly designed to juxtapose the existing point-of-care tests for UTI in terms of diagnostic accuracy in primary and secondary care. This systematic review will help describe more effective POCTs at the point-of-care level about identification (ID) and antimicrobial susceptibility testing (AST) and their respective specificity and sensitivity.

Methods and Analysis:

This review will include studies that evaluate POCT accuracy for UTI diagnosis. Detailed literature searches in electronic databases will be conducted in PubMed, Web of Science, Embase, and the Cochrane Database of Systematic Reviews from their inception to July 2024. The studies to be included are random control trials (RCTs), cluster RCTs, observational studies, and regulatory reports reporting on direct comparisons of POCTs to standard urine culture. Data extraction will be done by two reviewers who work independently. Discrepancies will be resolved through discussion. The Quality Assessment of Diagnostic Accuracy Studies risk of bias tool, QUADAS-2, will be used to assess the risk of bias in the studies included. A metaanalysis will

be undertaken when sufficient data are available. It will use bivariate random effects models for estimating summary sensitivity and specificity, accounting for heterogeneity among studies

Ethics and Dissemination

Formal ethical approval will not be sought because no identifiable personal data will be collected. This protocol for the systematic review will be registered with PROSPERO. The findings of review will be disseminated to stakeholders, including patient groups and clinicians and also presented at conferences. This review is expected to go on and make a contribution to the understanding of the available evidence about the effectiveness of POCTs for diagnosing UTIs, which may have an impact on clinical policy.

Analysis of Aetiology & Prevalence of Catheter Associated Urinary Tract Infection in a Tertiary Care Hospital of Uttarakhand

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Background

Catheter associated urinary tract infection (CAUTI) continues to be among the most common healthcare-associated infections in India. Despite advances in prevention guidelines, there remains a lack of knowledge concerning the risk factors for CAUTI. Hence this study was undertaken for better understanding of CAUTI.

Methods

The study was conducted in a tertiary care hospital attached to a teaching institute. Retrospective analysis of CAUTI bundle data was done from March 2023 to March 2024.

Result

Out of 15,842 catheterised patients, 61 patients (0.8%) developed CAUTI. Average duration of catheterisation among CAUTI patients was 15 days. Majority of the patients belonged to the age group 40yr to 69yr, followed by above 69 yr. There was no gender disparity. *Klebsiella pneumoniae* (36%), *Enterococcus spp* (31%) & *Escherichia coli* (25%) were the most common pathogen identified among 61 CAUTI patients, followed by *Pseudomonas aeruginosa* (6.6%) and *Acinetobacter baumannii complex* (4.9%). *Pseudomonas aeruginosa* & *Klebsiella pneumoniae* were the most resistant pathogen among all. Total deaths among CAUTI patients were seen in 11(18%) cases probably due to MDR organisms and multiple comorbidities.

Conclusion

This study underscores the importance of a comprehensive approach to CAUTI, encompassing

timely analysis, prevention & surveillance to reduce its prevalence & impact on patient care. Behavioural changes, including empowerment of nurses and adherence to all elements of the care bundle, can lead to significant and sustained improvement in reducing the CAUTI rate in hospitals.

Prevalence of Aeromonas Septicemia in Cancer Patients in Tertiary Care Hospital

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Background

Aeromonas Species (*A. hydrophila*, *A. caviae* and *A. veronii*) are emerging and potential human pathogens causing both intestinal and extra intestinal infections. *A. hydrophila* is the most prevalent Aeromonas species. It is an opportunistic pathogen and causes infections in immunosuppressed and anemic patients. The main clinical expressions are gastrointestinal tract disorders, wound, soft tissue infection and septicemia. We investigated the prevalence of Aeromonas species in cancer patients and studied their sensitivity pattern against CLSI approved antibiotics, which will help in management of clinical cases.

Methods

A 2 year retrospective study was conducted in the department of Microbiology at Tata Memorial Hospital. 179 isolates of Aeromonas species were obtained from clinical samples (blood, bile, pus, wound, urine, sputum, BAL, tracheal secretion). Identification and susceptibility testing was performed using standard guidelines by Vitek 2 system. VP Test was used to differentiate between *A. hydrophila* and *A. caviae*.

Result

Males (59.22%) were more commonly infected with Aeromonas sp. as compared to female (40.78%) patients. Patients in the age group of 31- 65 years were most infected (76.5%). *Aeromonas hydrophila* was the single most commonly isolated species. 60.9 % of the Aeromonas species were isolated from bile and PTBD followed by 17.3 % from pus/ swab and 15.7% from blood. 97.21% of the isolates were susceptible to Amikacin, followed by Gentamicin (95.53%) Imipenem (93.86%), Meropenem (92.19%), Ciprofloxacin (83.80%),

Ceftazidime (80.45%), and Piperacillin-Tazobactam (51.96%).

Conclusion

Aeromonas species mainly infect the GI tract in cancer patients. Amikacin, Gentamicin, Imipenem, Meropenem shows better susceptibility towards Aeromonas species when compared with Ciprofloxacin and Ceftazidime.

Microbacterium aurum Bacteremia in Immunosuppressed Patients- An Emerging Threat - Case Series

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Background

Bloodstream infection with *Microbacterium* species in humans is rare and frequently linked to the presence of immunosuppressed conditions such as patients on chemotherapy or corticosteroids. Presence of indwelling catheters is also a potential risk factor for *M. aurum* infection. No case report has been documented in the literature regarding the pathogenic potential of *M. aurum* in causing bacteremia. This is the first case series reporting bacteremia by *M. aurum* describing the risk factors and sensitivity pattern of this pathogen. In this case series, we have described bacteremia caused by *M.aurum* The risk factors and sensitivity pattern of this pathogen have also been evaluated.

Cases

We have described the clinical course and presentation of three patients whose blood culture showed growth of *M.aurum*. Indwelling venous catheter for hemodialysis or for chemotherapy for the treatment of acute lymphoblastic leukemia was found to be a risk factor in two patients. Rheumatoid arthritis was the underlying condition in the second patient and was started on immunosuppressants. Blood samples were collected during the febrile period. The blood culture samples of all these patients had pure isolates of *M. aurum*, identified by matrix-assisted laser desorption ionization–time of flight mass spectrometry. All three isolates were susceptible to ceftazidime and doxycycline. The patients were managed according to the sensitivity reports and were discharged in stable condition.

Conclusion

Microbacterium aurum can cause bacteremia in immunocompromised patients but respond well to susceptibility guided antibiotic therapy.

Unravelling the Trends of Sexually Transmitted Infective Syndromes Reported in an Apex Regional Centre

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Background

Sexually transmitted infections (STIs) pose a significant global public health challenge. The significance of addressing STI syndromes cannot be overstated. Publicly, they contribute significantly to the global disease burden, necessitating targeted interventions. Timely identification and treatment are pivotal at an individual level, preventing complications and enhancing overall health. Managing STI syndromes also aids in preventing transmission within communities and mitigating threats of antibiotic resistance. This study examines the demographic profiles of STI healthcare seekers, focusing on GUD Herpetic, Non-Herpetic, Vaginal, Cervical, and urethral discharge syndromes and analyses monthly trends to contribute insights for adaptive public health strategies.

Methods

A retrospective analysis of patient records from July 2022 to June 2023 was conducted. Demographic data, syndrome classification, and monthly trends were systematically extracted and analysed.

Results

The demographic analysis revealed distinct gender patterns among individuals seeking STI-related healthcare. Males were predominant in GUD Non-Herpetic (87.88%) and GUD Herpetic syndromes (66.89%), while females dominated VDRL testing, comprising 79.14% of cases. Syndromic classification further highlighted variations, emphasizing the importance of gender-specific factors in STI dynamics.

Monthly trend analysis depicted the dynamic nature of STIs, with fluctuations in reported cases. GUD non-herpetic syndrome peaked in April 2023 (6 cases), while GUD Herpetic and vaginal discharge peaked in December 2022 (19 and 96 cases respectively), indicating diverse seasonal trends in different syndromes. Amongst the GUD non-herpetic cases Syphilis was more common compared to Chancroid and Donovanosis.

Conclusion

This study provides significant insights into STI epidemiology, demographic profiles, syndrome prevalence, and temporal trends. Continuous monitoring and adaptive public health measures are crucial for addressing emerging patterns. Acknowledging limitations, prospective studies on socio-cultural factors' impact on STI dynamics are warranted

Seroprevalence of Leptospirosis in a Tertiary Care Hospital in Mumbai, India

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Background

Leptospirosis, a widespread re-emerging zoonotic disease especially in developing countries like India. According to WHO more than 5 lakh cases of leptospirosis are reported worldwide per year with mortality more than equal to 10% and its outbreak has been increased for the past 3 decades.

Aim: This study aims to determine the seroprevalence of leptospirosis among febrile patients at tertiary care hospital Mumbai.

Methods

It is a retrospective observational laboratory-based study over a period of one year in tertiary care hospital, Mumbai. A total of 5566 blood samples of febrile patients received for diagnosis of leptospirosis IgM antibodies using ELISA.

Results

A total of 1173 of 5566 blood samples that were received during the study period tested positive for leptospirosis IgM antibody. Seroprevalence of leptospirosis was found to be 21.07 % with maximum incidence observed during monsoon.

Conclusion

Leptospirosis has emerged as a health challenge among patients with acute febrile illness especially during monsoon, hence early diagnosis & proper antibiotic treatment is necessary to decrease mortality and morbidity.

Factors Influencing Patient Adherence to Prescribed Antifungals in Dermatophytosis and its Impact on Treatment Outcomes

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Background

Adherence to treatment protocol is of paramount importance in all areas of clinical medicine, including dermatology. Unfortunately, various reports in literature suggest that adherence may be the exception rather than the rule. There has been an unprecedented epidemic-like scenario of dermatophytosis in India in the past few years and nonadherence to treatment protocol is postulated as one of the most important factors contributing to it. This study aimed to investigate the factors that contribute to patient adherence to prescribed antifungal medications in treatment of dermatophytosis and examines how adherence influences treatment outcomes.

Methods

This was a cross-sectional descriptive study carried out over a period of one year in which clinically diagnosed patients of tinea infection were enrolled. The study employed a mixed-methods approach, combining quantitative surveys regarding demographic information, followed by clinical examination regarding type of dermatophyte infection; and qualitative interviews regarding understanding of treatment, medication-related beliefs, socioeconomic factors and other perceived barriers to adherence, to comprehensively explore the factors influencing patient adherence to prescribed antifungal medications. Finally the treatment outcomes in terms of mean duration of therapy for disease control and side effects were observed and compared

Results

A total of 500 patients, comprising 233 men and 267 women aged 18-60 years were included in the study. *Tinea corporis* was the most common variant seen in 260 (52%) patients followed by *Tinea cruris* (31.6%), *Tinea faciei* (7%), *Tinea pedis* (5.2%), *Tinea manuum* (3.4%) and *Tinea barbae* (0.8%). Only 8.8% of patients were treatment naïve with the majority of the patients having taken topical or oral antifungal treatment in the past. Antifungals were prescribed by dermatologists in 19.5% cases, general practitioner (GP) in 27.4% cases, local chemist (38.1%) and self-medication by 14.9% patients through TV advertisements.

Majority of the patients treated by GP, local chemist and self-medication; used topical steroid containing cream (92.6%) and oral fluconazole (56%). While itraconazole (88.2%) and topical luliconazole (78.6%) was most commonly prescribed by dermatologists. Factors affecting adherence were poor instructions given to patient by prescriber (34.2%), cost issues (29.3%), patients belief regarding safety of medication (17.8%), lack of patient education regarding disease (10.7%), poor memory on part of patients (7%) and side effects (3%). The average duration of antifungals treatment in the non-adherence group was 13.4 weeks.

Conclusion

These findings will provide valuable insights into the existing knowledge gaps and aid in designing targeted educational programs and interventions to enhance treatment adherence which might improve treatment outcomes.

Note: Few presented abstracts were removed in this publication considering lack of data or incompleteness.

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