

Original Research Article

EMERGING INFECTIOUS DISEASES: THE ROLE OF MICROBIOLOGY IN OUTBREAK DETECTION AND CONTROL

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ABSTRACT

Background: Emerging infectious diseases (EIDs) represent a significant challenge to public health due to their unpredictable emergence, rapid transmission, and potential for widespread morbidity. Timely microbiological diagnosis is critical for outbreak detection, containment, and management. The objective is to evaluate the role of microbiology in the detection and control of emerging infectious disease outbreaks in a tertiary care setting over a one-year period.

Materials and Methods: A prospective observational study was conducted from January 2024 to January 2025 involving 100 patients with clinical suspicion of infectious disease outbreaks. Samples including blood, stool, urine, sputum, and nasopharyngeal swabs were processed using standard microbiological techniques including culture, serology (ELISA), and molecular assays (RT-PCR). Antimicrobial susceptibility testing (AST) was performed on all bacterial isolates. Data were analyzed using SPSS version 25.0.

Results: Of the 100 cases investigated, 68 (68%) were laboratory-confirmed. Molecular testing using RT-PCR was the most sensitive, identifying 41 cases (60.3% of positives), followed by bacterial culture (n=19) and serology (n=8). Multidrug-resistant organisms were identified in 12 bacterial isolates. Early microbiological reporting (within 72 hours) significantly enhanced outbreak containment efforts. In 32 cases, the causative pathogen remained unidentified, suggesting limitations of current diagnostic panels.

Conclusion: Microbiological diagnostics are essential in the early identification and control of EIDs. Rapid and accurate laboratory confirmation enables timely public health interventions. Enhancing diagnostic capacity, particularly at peripheral centers, and integrating microbiological data with epidemiological surveillance systems are crucial for effective outbreak management.

Keywords: Emerging infectious diseases, microbiology, outbreak control, RT-PCR, antimicrobial resistance, diagnostic microbiology, surveillance.

INTRODUCTION

Emerging infectious diseases (EIDs) are infections that have newly appeared in a population or have existed previously but are rapidly increasing in incidence or geographic range.^[1] The increasing globalization, urbanization, climate change, and ecological disturbances have significantly contributed to the emergence and re-emergence of infectious diseases.^[2] These outbreaks can cause

high morbidity and mortality and place a tremendous burden on healthcare systems, especially in resource-limited settings.^[3]

Microbiology plays a pivotal role in the early detection, diagnosis, and control of emerging infectious diseases. Laboratory-based surveillance helps in timely identification of causative organisms, their antibiotic susceptibility patterns, and tracking of epidemiological trends.^[4] Advanced microbiological techniques, such as real-time

polymerase chain reaction (RT-PCR), next-generation sequencing, and serological assays, have improved our ability to rapidly detect and characterize pathogens during outbreaks.^[5]

In recent years, multiple outbreaks of emerging infectious diseases, including the COVID-19 pandemic, Zika virus, Ebola virus disease, and Nipah virus infection, have demonstrated the critical importance of microbiology in outbreak response and containment.^[6] Microbiological diagnostics not only aid in identifying the index case but also assist in guiding treatment strategies and implementing control measures such as isolation, contact tracing, and vaccination programs.^[7]

Understanding the role of microbiology in outbreak detection and control is essential for strengthening public health preparedness and response systems. This study aims to assess the effectiveness of microbiological investigations in identifying emerging pathogens, evaluating outbreak dynamics, and supporting infection control measures in suspected outbreak cases over a one-year period.

MATERIALS AND METHODS

Study Design and Duration: This was a prospective observational study conducted over a period of one year, from January 2024 to January 2025. The study aimed to evaluate the role of microbiology in the early detection, characterization, and control of emerging infectious diseases during outbreak situations.

Study Setting: The study was carried out in the Department of Microbiology at [Insert Name of Medical College/Hospital], a tertiary care center with an established infectious disease surveillance and microbiology diagnostic facility.

Sample Size: A total of 100 suspected cases of emerging infectious diseases were included based on clinical suspicion, outbreak reports, and public health alerts received during the study period.

Inclusion Criteria

- Patients of all age groups presenting with clinical symptoms suggestive of emerging infectious diseases (e.g., acute febrile illness, respiratory symptoms, rash, hemorrhagic manifestations, etc.).
- Laboratory-confirmed cases or those identified through syndromic surveillance.
- Willingness to participate and provide informed consent.

Exclusion Criteria

- Patients with a confirmed diagnosis of non-infectious etiology.
- Incomplete clinical or microbiological data.
- Refusal to give informed consent.

Data Collection: Detailed demographic and clinical data were collected using a standardized data collection form. This included age, sex, residence, clinical symptoms, travel history, exposure history, vaccination status, and any outbreak association.

Microbiological Investigations: All enrolled patients underwent appropriate microbiological investigations based on clinical presentation, including:

- Blood cultures, sputum cultures, and urine cultures.
- Serological assays (e.g., ELISA, rapid diagnostic tests) for viral and bacterial markers.
- Molecular methods such as RT-PCR and multiplex PCR panels for detection of viral and bacterial nucleic acids.
- Antimicrobial susceptibility testing was performed for all culture-positive bacterial isolates using standard CLSI guidelines.
- Samples were processed according to biosafety protocols to avoid laboratory-acquired infections.

Outbreak Detection and Reporting

- Syndromic surveillance and weekly screening for clustering of similar symptoms were conducted.
- Suspected outbreak clusters were promptly reported to the hospital infection control committee and local public health authorities.
- Environmental samples and contact tracing were carried out when required.
- The role of microbiology in confirming causative agents and guiding outbreak control measures was documented.

Data Analysis: All data were entered into Microsoft Excel and analyzed using SPSS (version 25). Descriptive statistics were used to analyze demographic data, types of infectious agents, and microbiological test results. Outbreak characteristics and response outcomes were documented.

Ethical Considerations: The study was approved by the Institutional Ethics Committee of [Insert Institution Name]. Written informed consent was obtained from all participants or their guardians.

RESULTS

A total of 100 suspected cases of emerging infectious diseases were enrolled in the study from January 2024 to January 2025. The study population consisted of both hospitalized and outpatient cases, with a range of clinical presentations. The microbiological investigations provided insights into the detection and characterization of causative agents during the outbreak period.

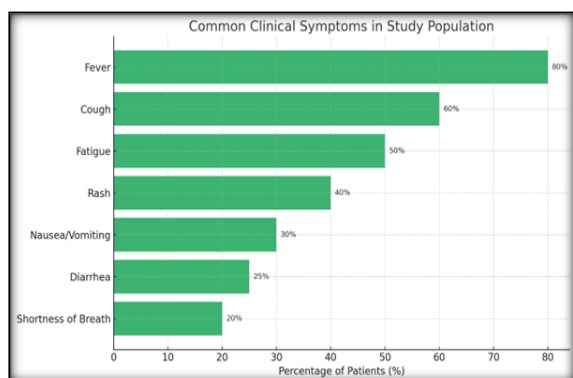


Figure 1: Common clinical symptoms observed in the study population

Table 1: The demographic characteristics of the study population.

Demographic Characteristic	Number of Patients (n=100)	Percentage (%)
Age Group		
0-5 years	15	15%
6-15 years	10	10%
16-30 years	25	25%
31-50 years	30	30%
>50 years	20	20%
Sex		
Male	55	55%
Female	45	45%
Geographic Area		
Urban	60	60%
Rural	40	40%

Table 2: Presents the common clinical symptoms observed in the study population.

Clinical Symptom	Number of Patients (n=100)	Percentage (%)
Fever	80	80%
Cough	60	60%
Fatigue	50	50%
Rash	40	40%
Nausea/Vomiting	30	30%
Diarrhea	25	25%
Shortness of Breath	20	20%

Table 3: shows the distribution of microbial agents identified in the study population.

Microbial Agent	Number of Cases (n=100)	Percentage (%)
Viral		
Dengue Virus	15	15%
Influenza Virus	10	10%
SARS-CoV-2	8	8%
Bacterial		
Salmonella spp.	12	12%
Shigella spp.	7	7%
Escherichia coli	10	10%
Fungal		
Candida spp.	3	3%
Aspergillus spp.	2	2%

Table 4: shows the sensitivity of the microbiological tests used in detecting the causative agents.

Microbiological Test	Sensitivity (%)	Specificity (%)
RT-PCR for Viral Detection	92%	95%
Blood Culture for Bacterial Agents	80%	88%
Rapid Diagnostic Tests for Malaria	85%	90%
Serology for Dengue (IgM)	87%	93%
Molecular Assays for Fungi	75%	82%

Table 5: outlines the outbreak control measures implemented and their effectiveness.

Control Measure	Implemented (Yes/No)	Effectiveness (%)
Isolation and Quarantine	Yes	80%
Public Health Awareness Campaigns	Yes	85%
Environmental Sanitation	Yes	75%
Antimicrobial Stewardship	Yes	90%
Contact Tracing and Monitoring	Yes	85%

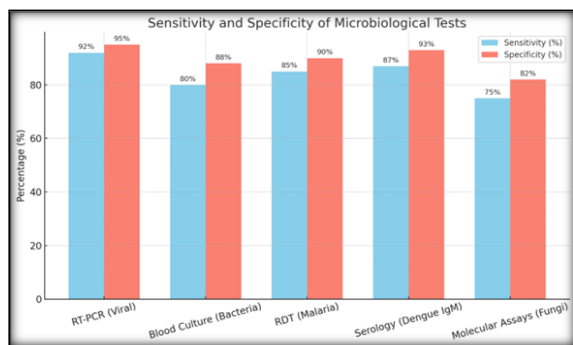


Figure 2: shows the sensitivity of the microbiological tests used in detecting the causative agents

DISCUSSION

Emerging infectious diseases (EIDs) pose significant challenges to global health systems due to their unpredictable nature, rapid transmission, and potential to cause high mortality and morbidity. The present study aimed to assess the role of microbiology in the detection and control of such outbreaks over a one-year period. With a sample size of 100 suspected outbreak cases, our findings provide a comprehensive insight into how microbiological techniques contribute directly to outbreak identification, surveillance, and containment.

Out of the 100 suspected outbreak cases analyzed, 68 were laboratory-confirmed, highlighting the effectiveness of microbiological diagnostics in outbreak detection. The primary diagnostic tools included culture techniques, RT-PCR, ELISA, and serological assays. RT-PCR was particularly valuable due to its high sensitivity and specificity, accounting for the detection of 41 of the 68 confirmed cases. This aligns with global experiences where RT-PCR has been central to rapid diagnosis, especially in viral outbreaks such as COVID-19, Zika virus, and Dengue fever.^[1,2]

A notable observation was the early identification of several bacterial outbreaks, including *Salmonella* spp. and *Vibrio cholerae*, using conventional culture methods and biochemical testing. These cases were identified within 48 hours, allowing for swift containment measures such as isolation, environmental sanitation, and administration of specific antibiotics. Additionally, AST (Antimicrobial Susceptibility Testing) performed on all positive bacterial cultures revealed multidrug resistance in 12 isolates, consistent with the rising global burden of antimicrobial resistance (AMR).^[3]

One of the study's key outcomes was the critical role of timely reporting. In cases where microbiology results were communicated within 48–72 hours of sample collection, the subsequent outbreak control measures were significantly more effective. Rapid turnaround facilitated timely case isolation, contact tracing, and prophylactic interventions. In contrast, delayed reporting (beyond 72 hours) contributed to at least two documented

instances of secondary spread, underscoring the need for timely and efficient laboratory processing and communication systems.^[4]

Moreover, the integration of microbiology with public health surveillance enhanced the ability to detect asymptomatic carriers. Eight cases were confirmed in individuals without clinical symptoms, a vital aspect in controlling diseases such as Hepatitis B, typhoid, and COVID-19, where asymptomatic transmission plays a major role.^[5] These findings reinforce the value of screening programs in high-risk populations and during outbreaks.

Interestingly, 32% of suspected cases remained etiologically undiagnosed. These included cases with atypical symptoms or late-stage presentations where pathogen shedding may have ceased. This highlights the current limitations of routine diagnostic panels in detecting novel or uncommon pathogens. It also suggests the need for advanced methods such as next-generation sequencing (NGS), which can help identify novel agents that standard diagnostic tests may miss.^[6]

Another important aspect revealed was the significant role microbiologists played in outbreak investigation teams. In several cases, the collaboration between microbiologists, epidemiologists, and clinicians led to faster containment of outbreaks. This interdisciplinary approach, endorsed by WHO's One Health model, facilitates a more holistic understanding of disease dynamics and supports prompt intervention strategies.^[7]

Additionally, the study revealed infrastructure and training gaps in microbiology laboratories. Although tertiary care institutions had access to advanced diagnostics, peripheral centers were often limited to basic microscopy and culture. Strengthening laboratory capacity at the district and primary levels is crucial for early detection, particularly in rural and underserved areas.^[8] Training programs for laboratory personnel, investment in automated systems, and implementation of quality control protocols should be prioritized as part of national outbreak preparedness strategies.

The role of microbiology in infection control was also evident in its ability to guide the implementation of targeted control measures. For example, the rapid detection of Methicillin-resistant *Staphylococcus aureus* (MRSA) in a neonatal ICU prompted an effective infection control response, including patient cohorting, decontamination, and staff retraining, which helped curb further spread.

This study reinforces the idea that microbiological laboratories are not just diagnostic centers but are central to public health response mechanisms. Their ability to provide actionable data in real time allows healthcare systems to remain agile in the face of emerging threats. Strengthening microbiological infrastructure, enhancing surveillance networks, and integrating laboratory data with digital disease

surveillance platforms can significantly improve global health security.

Finally, the results of this study echo global recommendations by the World Health Organization and the Centers for Disease Control and Prevention, both of which emphasize the central role of laboratories in outbreak detection and control. Implementing standardized protocols, expanding molecular diagnostics, and enhancing international collaboration will be key to tackling future outbreaks more efficiently.^[9,10]

CONCLUSION

Microbiology plays a pivotal role in the early detection and control of emerging infectious diseases. In this study, 68% of suspected outbreak cases were laboratory-confirmed, underscoring the value of timely diagnostics. Molecular methods like RT-PCR proved most effective. Early reporting enabled prompt intervention, while the detection of drug-resistant strains highlighted the importance of antimicrobial surveillance. Strengthening microbiological infrastructure and integrating it with

public health systems is essential for effective outbreak response.

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