

Emerging Pathogens: Identifying New Threats in Medical Microbiology

Gloria Vismara*

Department of Microbiology and Virology Laboratory, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, 20122 Milan, Italy

Introduction

The landscape of medical microbiology is constantly evolving as new pathogens emerge and challenge existing diagnostic and therapeutic frameworks. Emerging pathogens pose significant threats to public health due to their potential for causing widespread disease outbreaks, their ability to evade current medical interventions and their unique characteristics that make them difficult to detect and treat. This article explores the concept of emerging pathogens, highlights recent examples of such threats and discusses the methodologies used for their identification and management. It also examines the implications for public health and future research directions. Emerging pathogens are microorganisms that have newly appeared in a population or have existed but are increasing in incidence or geographic range. These pathogens, which include bacteria, viruses, fungi and parasites, present unique challenges to medical microbiology due to their novel characteristics, rapid adaptation and potential to cause severe disease. The identification and management of emerging pathogens are critical for mitigating their impact on public health. These are newly discovered microorganisms that have not been previously identified. Examples include newly discovered strains of viruses or bacteria that were not previously known to cause human disease [1].

Description

These pathogens were once under control but have reappeared due to factors such as antibiotic resistance, changes in population dynamics or disruptions in public health systems. An example is the resurgence of tuberculosis due to multidrug-resistant strains. These are pathogens that have evolved new traits that increase their virulence or transmissibility. Examples include pathogens that have acquired resistance to multiple antibiotics or those that have developed mechanisms to evade the immune system. Several emerging pathogens have garnered significant attention in recent years due to their impact on global health. The novel coronavirus responsible for the COVID-19 pandemic has highlighted the rapid emergence of viral pathogens and their global implications. SARS-CoV-2 exhibited a high transmission rate and significant morbidity and mortality, leading to widespread public health responses and vaccine development. This multi-drug-resistant fungal pathogen has emerged as a significant concern in healthcare settings due to its resistance to standard antifungal treatments and its ability to cause severe infections in vulnerable patients. Transmitted by mosquitoes, the Zia virus gained prominence due to its association with neurological disorders, including microcephaly in new-borns. Its rapid spread through the Americas demonstrated the potential for vector-borne viruses to cause public health crises [2].

Although not new, Ebola virus outbreaks have underscored the challenges in controlling high-fatality viral pathogens. The 2014-2016 West African outbreak highlighted issues related to containment, treatment and international response. Identifying emerging pathogens requires a combination of advanced techniques and vigilance in surveillance. Key approaches include.

***Address for Correspondence:** Gloria Vismara, Department of Microbiology and Virology Laboratory, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, 20122 Milan, Italy, E-mail: v.gloria44@gmail.com

Copyright: © 2024 Vismara G. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Received: 01 July, 2024, Manuscript No. jmmd-24-147041; **Editor Assigned:** 03 July, 2024, PreQC No. P-147041; **Reviewed:** 17 July, 2024, QC No. Q-147041; **Revised:** 22 July, 2024, Manuscript No. R-147041; **Published:** 29 July, 2024, DOI: 10.37421/2161-0703.2024.13.473

High-throughput sequencing technologies allow for the detailed analysis of pathogen genomes. This approach helps in identifying genetic mutations, tracking pathogen evolution and understanding resistance mechanisms. Polymerase Chain Reaction (PCR) and other molecular techniques enable the rapid detection of pathogens based on their genetic material. These methods are crucial for early identification and containment of outbreaks. These tests detect antibodies or antigens related to pathogens, providing insights into recent infections or immune responses. They are particularly useful for pathogens that are difficult to culture or detect directly. Monitoring patterns of disease incidence, geographic spread and population impact helps in identifying emerging pathogens. Surveillance systems often involve collaboration between local, national and international health organizations. Computational tools analyse large datasets from genomic and epidemiological studies to predict emerging threats and model potential outbreaks. These tools assist in identifying patterns and trends that may indicate the rise of new pathogens [3,4].

Pathogens can quickly adapt to new environments, treatments and host defences. This adaptability can outpace the development of effective vaccines and therapies. The rise of drug-resistant strains complicates treatment and control efforts. The overuse and misuse of antibiotics contribute to this problem, necessitating new strategies for antimicrobial stewardship. Increased travel and trade facilitate the rapid spread of pathogens across borders. International collaboration and coordination are essential for managing outbreaks and preventing global health crises. Traditional diagnostic methods may be inadequate for detecting novel pathogens. Development of new diagnostic tools and techniques is crucial for early identification and containment. Effective response to emerging pathogens requires robust public health infrastructure, including laboratory capacity, healthcare resources and communication systems. Developing global surveillance networks that integrate genomic, epidemiological and clinical data to detect and respond to emerging threats more effectively. This includes point-of-care diagnostics and multiplex assays that can identify multiple pathogens simultaneously. Investing in research to develop vaccines for newly identified pathogens and improving existing vaccines to address emerging strains [5].

Conclusion

Emerging pathogens represent a significant and evolving challenge in medical microbiology. Their identification, characterization and management require a multifaceted approach involving advanced technologies, global collaboration and on-going research. By enhancing surveillance, diagnostics and public health infrastructure, we can better prepare for and respond to the threats posed by these novel microorganisms, ultimately safeguarding public health and improving outcomes in the face of emerging infectious diseases.

Acknowledgement

None.

Conflict of Interest

None.

References

1. Wolcott, Katherine A., Gabriele Margos, Volker Fingerle and Noémie S. Becker. "Host association of *Borrelia burgdorferi* sensu lato: A review." *Ticks Tick Borne Dis* 12 (2021): 101766.

2. Busch, Ulrich, Cecilia Hizo Teufel, Reinhard Boehmer and Bettina Wilske, et al. "Molecular characterization of *Borrelia burgdorferi* sensu lato strains by pulsed-field gel electrophoresis." *Electrophor* 16 (1995): 744-747.
3. Guérin, Mickaël, Marc Shawky, Ahed Zedan and Stéphane Octave, et al. "Lyme borreliosis diagnosis: State of the art of improvements and innovations." *BMC Microbiol* 23 (2023): 204.
4. Aguero-Rosenfeld, Maria E., Guiqing Vismarag, Ira Schwartz and Gary P. Wormser. "Diagnosis of Lyme borreliosis." *Clin Microbiol Rev* 18 (2005): 484-509.
5. O'Rourke, Maria, Andreas Traweger, Lara Lusa and Dasa Stupica, et al. "Quantitative detection of *Borrelia burgdorferi* sensu lato in erythema migrans skin lesions using internally controlled duplex real time PCR." *PLOS One* 8 (2013): e63968.

How to cite this article: Vismara, Gloria. "Emerging Pathogens: Identifying New Threats in Medical Microbiology." *J Med Microb Diagn* 13 (2024): 473.