Newer Viral Infection in Bangladesh--are We Prepared to Face Them?

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Despite an elaborate planning and measures to tackle microbes, emerging infectious diseases remain a crucial global challenge. Emerging infections can be defined as "infections that have newly appeared in a population or have existed previously but are rapidly increasing in incidence or geographic range"¹. Infectious diseases account for 20% of global mortality, with viral diseases causing about one third of these deaths². In addition to the rapidly changing social, environmental and behavioural patterns observed among the population of Bangladesh in the face of rapid urbanization and ever-increasing global travel and trade, there is also a dramatic rise in the spectrum of infectious diseases³. The unprecedented growth of the population sharing a fragile ecology, close interactions of humans and animals, large urban dwellers living in unhygienic conditions, highly vulnerable food security and safety along with the adversities of natural disasters and global warming, make Bangladesh increasingly vulnerable to the public health threats of emerging and re-emerging diseases (EIDs) and antibiotic resistance⁴. Emergence/re-emergence of several viral infections have been reported from Bangladesh in the past few decades⁵. They include infection due to Nipah, Severe acute respiratory syndrome (SARS), Influenza H5N1, Influenza H1N1, Influenza H7N9, Chikungunya, Human Enterovirus-71, Dengue, Japanese Encephalitis, Severe fever with thrombocytopenia syndrome (SFTS), Middle East Respiratory Syndrome Corona Virus (MERS CoV) and Zika virus infection. This increase is a result of both the increase in the rate of emerging zoonotic infections across the globe and our enhanced ability to detect and identify agents. When considering the threat posed by emerging viruses, it is useful to divide them into three categories: newly emerging viruses, reemerging viruses, and viruses deliberately spread by bioterrorists. Recognizing this urgent need to respond to the rising threats of highly infectious viral diseases and to comply with the International Health Regulations (2005) by WHO, Bangladesh has already taken several steps to address a number of issues for maintaining a sustainable animal, human and eco-health, including proper waste disposal in laboratories and industries, hygienic animal slaughtering practices in wet markets, and monitoring of antibacterial resistance and residual effect⁶-⁷. Genomic sequencing has facilitated identification of viruses, other advances are helping to diagnose viral illnesses. Specifically, polymerase chain reaction (PCR), the nucleic acid amplification technology initially developed in 1983, is now more rapid and less cumbersome, allowing for deployment in outbreaks, even in resource-poor settings⁸. New technologies have expanded the sensitivity and scope of our detection and diagnostic capability⁹. However, a pathogen may still go undetected if it does not cause a significant disease outbreak. Creating increased awareness and training of clinical microbiologists/virologists for identification of new/emerging pathogens, and prompt reporting and management of outbreaks is essential to tackle the threat posed by emerging/re-emerging infection.

References :

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