A Short Review on Nipah Virus

P. Swathi*, K.Shruthi*, G.saiprasana*

1 Department of Microbiology, Siddhartha Women's Degree and pg college Kukatpally, Hyderabad **Corresponding author:** K shruthi, Department of Microbiology, Siddhartha Women's Degree and pg college Kukatpally, Hyderabad

Abstract:

Nipah viral complaint is a zoonotic infection caused by Nipahcontagion (NIV), a paramyxo contagion. Belonging to the rubric Henipavirus of the family Paramyxoviridae. The scientific name of the Nipah contagion is Nipah henipavirus'. The Nipah genome is a Single (non-segmented) negative-sense, single-stranded RNA of over 18kb, which is mainly longer than of other paramyxoviruses. This complaint was. first linked in 1998 by a platoon of experimenters at the Faculty of Medicine, University of Malaya. during an outbreak in Malaysia. numerous of people in Malaysia are diagnosed with the Nipah contagion & and treated at the University of Malaya Medical Centre. The contagion was insulated and linked in 1999. The complaint is named after a vill in Malaysia, (Sungai Nipah) It was also honored in Bangladesh in 2001, and nearly periodic outbreaks have passed. in that country since. this complaint has also been linked periodically in Eastern India. The contagion has been set up in the given natural force and several other club species in several countries.

Introduction:

Human contagious range from asymptomatic infection to acute respiratory infection (mild, severe) and fatal encephalitis. Infected people originally develop symptoms including fever, headaches, myalgia (muscle pain), Vomiting and sore throat. This can be followed by dizziness, internal confusion neurological signs, altered knowledge and disorientation severe respiratory problems. Including acute respiratory torture. Encephalitis and Seizures do in Severe Cases and can lead to coma within 24 to 48 hours. Encephalitis, inflammation of the brain, is a potentially fatal complication of Nipah virus. The people who have breathing difficulty are more likely than those of normal people. The complaint is suspected in characteristic individualities in the environment of an epidemic outbreak still the incubation period is believed to range from 4 to 14 days, an incubation period as long as 45 days has been reported. Although the Nipah contagion has caused only a many known outbreaks in Asia. It infects a wide range of creatures and causes several conditions and death in people, making, it a public health concern. Roughly 20 of cases are left with residual neurological Consequences like Seizure complaints and personality changes. Only a small number of people who recovered later fall (or) developed delayed onset encephalitis Transmission During the first honored outbreak in Malaysia & and in Singapore, utmost mortal or infections redounded from direct contact with sick gormandizers (or) their defiled napkins. Transmission is allowed to have occurred via vulnerable exposure to concealment from the gormandizers, (or) vulnerable contact with the towel of a sick" beast. Consumption of fruits or fruit products" defiled with urine on slaver from infected" fruit club was the most likely source of infection. Mortalto-mortal transmission of Nipah contagion has also been reported among family and Caregivers of infected cases. Close contact with a person infected with Niv or their b body fluids (includes blood, nasal or respiratory driblets). During the after outbreaks in Bangladesh and India Nipah contagion spread directly from person to person through close contact with People's stashing and excretions. In India and Siliguri in 2001, the transmission of the contagion was reported within a healthcare setting Around half of the reported cases in Bangladesh were due to mortal transmission, through furnishing course to infected cases.

Opinion original Signs or Symptoms of Nipah contagion are nonspecific and the opinion is frequently not suspected at the time of donation. In addition, the quality, volume type, and timing of clinical sample

collection & and the time demanded to transfer samples to the laboratory can affect the delicacy of laboratory results. This can also hamper accurate opinion and Create challenges in outbreak, and discovery. The main tests used are real-time polymer chain response (RT- PCR) from body fluids and antibody discovery via enzyme-linked immunosorbent assay (ELISA). Other tests used include PCR assay and contagion insulation by cell Culture RT- RT-PCR is taken from throathearties, Cerebrospinal fluid, urine, and blood analysis. After recovery, IGG and IGM antibody discovery can confirm a previous Nipah contagion infection." Immunohistochemistry on apkins collected during necropsy also confirms the complaint. There are presently no medicines (or) vaccines specifically for Nipah contagion infection although WHO has linked Nipah as a precedence complaint for the WHO Research and Development of Blueprint. Treatment is limited. Probative care, including rest, hydration, and treatment of symptoms as they do. Still, immunotherapeutic treatments are presently under development and evaluation. for treatment of NIV infections. One similar. monofloral Antibody" m102.4', has completed phase 1 clinical trials. The medicine ribavirin was used to treat a small number of cases in the Malaysian NIV outbreak, but its efficacity in people is unclear. The Natural host of NIV is fruit club; it outbreaks in domestic creatures similar as gormandizers (whi ch are substantially 4 seen), nags, scapegoats, lamb, pussycats, and tykes. Educating people about the measures they can take to reduce" exposure or threat of NIV infection to them. WHO is supporting affected and threat countries with specialized guidance on how to manage NIV and how to help their occurrence. Fruit with signs club mouthfuls should be discarded.

1. Origin and Reservoir Hosts:

Nipah virus is believed to originate from fruit bats of the Pteropus genus, which serve as natural reservoirs. These bats shed the virus in their saliva, urine, and excreta, and their proximity to humans and domestic animals can facilitate transmission. NiV has been detected in various animal species, including pigs, which can act as intermediate hosts in transmitting the virus to humans.

2. Transmission:

Human-to-human transmission of Nipah virus primarily occurs through close contact with infected individuals, particularly in healthcare settings. Transmission can also occur through the consumption of contaminated date palm sap or fruits that have been partially eaten by infected bats. The virus can spread through respiratory droplets, making it a significant concern during outbreaks.

3. Clinical Presentation:

Nipah virus infection can lead to a wide range of clinical manifestations, from asymptomatic or mild flu-like symptoms to severe respiratory illness and encephalitis. The latter can result in altered consciousness, seizures, and death. Nipah virus is notorious for its high mortality rate, which can exceed 70% in some outbreaks.

4. Diagnosis:

Diagnosing Nipah virus infection is challenging, as early symptoms often resemble those of other respiratory illnesses. Laboratory tests, such as reverse-transcription polymerase chain reaction (RT-PCR) and antibody-based assays, are crucial for confirmation. Timely diagnosis is critical for implementing control measures during outbreaks.

5. Prevention and Control:

There is no specific antiviral treatment for Nipah virus, making prevention a paramount concern. Preventive measures include avoiding consumption of raw date palm sap and fruits, practicing proper hygiene, and implementing infection control practices in healthcare settings. Vaccines and experimental treatments are being researched to combat this deadly virus.

6. Ongoing Research:

Research on Nipah virus continues to advance our understanding of its molecular biology, pathogenesis, and epidemiology. Efforts are ongoing to develop vaccines and antiviral therapies, which could prove instrumental in preventing future outbreaks and improving patient outcomes.

Conclusion: Nipah Contagion outbreak should be suspected in applicable epidemiological settings in clusters of cases presenting with acute encephalitis with(or) without ARDS, high secondary attack rate and veritably high mortality The cases should be managed with applicable infection control. Measures. The part of medicines like ribavirin must be easily established with the help of duly designed trials Until the time when newer medicines are developed for their effective treatment. To help its transmission from creatures to humans in conditions Prone areas must instibuted.

Reference:

- 1. Clayton BA, Wang LF and Marsh GA (2013) Henipaviruses: an updated review focusing on the pteropid reservoir and features of transmission. Zoonoses and Public Health **60**, 69–83. [PubMed] [Google Scholar]
- Institute of Epidemiology, Disease Control and Research (IEDCR). Available at <u>http://www.iedcr.org/</u> (Accessed 8 October 2018)
- 3. WHO (2018) WHO | Nipah Virus Infection. World Health Organization; Available at <u>http://www.who.int/csr/disease/nipah/en/</u> (Accessed 17 June 2018). [Google Scholar]
- 4. Luby SP (2013) The pandemic potential of Nipah virus. Antiviral Research **100**, 38–43. [PubMed] [Google Scholar]
- 5. Chatterjee P (2018) Nipah virus outbreak in India. The Lancet **391**, 2200. [Google Scholar]
- 6. Chua KB (2003) Nipah virus outbreak in Malaysia. Journal of Clinical Virology **26**, 265–275. [PubMed] [Google <u>Scholar</u>]
- 7. Chadha MS et al. (2006) Nipah virus-associated encephalitis outbreak, Siliguri, India. Emerging Infectious Diseases **12**, 235–240