



## Nipah Virus Infection in Asia Region: A Review Update



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### Abstract

The Nipah virus is extremely virulent and has been linked to neurological and respiratory diseases in both humans and animals. Fruit bats, virus-containing fruit and date juice consumed by animals, as well as airborne droplets, are the main sources of the virus's dissemination. Over the previous decades, Asian countries have experienced several epidemics and significant fatality rates. Additionally, the Nipah virus infection disrupted the socio-economic progress of the affected nations, which hinders the achievement of the SDGs. [*Bangladesh Journal of Infectious Diseases*, June 2022;9(1):31-37]

**Keywords:** Nipah virus; NiV infection; Asia; SDGs; Socio-economy

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### Introduction

Ten major outbreaks occurred by various virus types that were highly infectious and deadly from 1967 to 2020<sup>1</sup>. Marburg had the highest fatality rate of 80.0% among these ten outbreaks, which caused an outbreak<sup>2</sup>. The novel Coronavirus infected around 270.2 million people until December 2021, and more than 5.3 million deaths occurred, which had a fatality rate of 2.2% in January 2020<sup>3</sup>. The second-highest fatality rate was 77.6 % for the Nipah virus in 1998<sup>4</sup>. Nipah virus (NiV) causes Nipah virus infection, a zoonotic disease transmitted by an animal to humans through contaminated food and human to human through direct transmission<sup>5</sup>. Nipah virus is a profoundly virulent, bat-borne virus widespread throughout tropical climates<sup>6</sup>. *Nipah henipavirus* belongs to the Mononegavirales order, the Paramyxoviridae family and several species of Pteropus fruit bats are the natural sources of infection<sup>7</sup>. It can be

asymptomatic and infectious, which causes respiratory illness, seizures, and fatal severe encephalitis<sup>8</sup>. This study demonstrates Nipah virus infection as a public health issue by knowing theories, models, strategies, and ethical aspects. This assignment critically explains the importance of the Nipah virus as a health-related problem by using epidemiological data and principles of population health that relate to socio-economic development. It also addresses the ethical issues and implications done for this highly infectious disease and critically discusses the different measures taken by the community to control the disease in Asian countries.

### Critically Examines Nipah Virus Infection as a Contemporary Public Health Issue

A total of 700 instances of Nipah virus infection have been detected from 1998 to 2018 worldwide, and among them, 50.0 to 75.0% died after being

infected with the Nipah virus<sup>9</sup>. Due to NiV infection, 17 deaths were reported in India in May 2018<sup>10</sup>. Even though only a few cases have been detected so far in several Asian nations, because of the virus's highly infectious features, it has the potential to infect many animals and humans, posing a threat to public health<sup>11</sup>. Symptoms and further laboratory investigation can diagnose the disease<sup>12</sup>. There is no specific treatment for NiV infection; only symptomatic management can be given. Up until this point, no vaccine has been developed<sup>13</sup>. The Nipah virus is classified as a priority disease on the World Health Organization's research and development roadmap list<sup>14</sup>. The organization is worried that NiV infection is a public health issue that requires close monitoring, research, and development<sup>15</sup>.

### Historical Perspective of the Nipah Virus

The virus was officially titled "Nipah" after the name of the Nipah river village (Kampung Sungai Nipah)<sup>16</sup>. Nipah Virus was first discovered in a pig farm around September 1998 in Ipoh, the northern side of Peninsular, Malaysia. Moreover, some farmers of Ipoh sold the pigs in a cut-rate sale, and some pigs crossed the border, including some areas of the South such as Sikamat, Kampung Sungai Nipah, Kampung Sawah, and Bukit pelanduk<sup>17</sup>. The third severe outbreak occurred in Bukit Pelanduk in December 1998<sup>18</sup>. However, the cases were initially diagnosed as Japanese B encephalitis (JE) with previous outbreak history in Malaysia, and treatments and prevention measurements were taken accordingly. Later, it revealed that most cases had a contact history with pigs and members of the same household, which was not like Japanese B encephalitis<sup>19</sup>. The rate of infectivity was more than Japanese B encephalitis which was 1:300<sup>20</sup>. Additionally, the affected persons were adult males and held an immunization history against JE and anti-JE. Furthermore, it was reported that pigs had severe coughs and fatality<sup>17</sup>. Meanwhile, The University of Malaya Hospital discovered the Nipah virus from the Paramyxoviridae family, which reacted with the Hendra virus antibody in March 1999<sup>21</sup>.

### Demography and Epidemiology of Nipah Virus

**Malaysia/Singapore:** During the first outbreak of Nipah in Malaysia, in 1998, among five hundred thousand population, NiV infected 27 people and 15 people died. After the post-mortem, it was confirmed that 9 of the people died because of NiV infection<sup>22</sup>. The following outbreaks occurred in South and Bukit Pelanduk, where more than 180

people were infected<sup>22</sup>. Serum was collected from 28 patients, where Japanese encephalitis (JE) specific IgM and JE nucleic acids were found in 4 serums<sup>23</sup>. Hence, to prevent the occurrence of Japanese encephalitis, the government took measures such as JE immunization and killing mosquitoes<sup>24</sup>. Meanwhile, 500 cases were detected in Singapore who worked in a slaughterhouse with symptoms of fever and respiratory illness or neurological disease<sup>25</sup>. The Ministry of Health tracked down the cause of the outbreak that revealed pigs imported from a Malaysia farmhouse spreading the infection and slaying 2 to 3 weeks prior to the spread of the disease<sup>26</sup>. However, it was revealed by University Hospital, Kuala Lumpur, that the cause of outbreaks was Nipah virus infection<sup>27</sup>. Malaysia was a predominantly Muslim country that was unlikely to have a contact history with pig and pig-containing food. After thorough investigations, a natural reservoir for the virus was found: Pteropodid fruit bats<sup>28</sup>. Bats usually eat fruits. Partially eaten fruits may be dropped down into pigsties, and pigs can be infected by eating contaminated food<sup>29</sup>.

**Bangladesh:** In 2001, the outbreaks occurred in 20 districts of Bangladesh located in the central and north-western regions during the winter season. Drinking date juice is the primary derivation of spreading the virus<sup>27</sup>. Date juice is ubiquitous and famous in the winter season in Bangladesh. The date juice is collected by snipping the husk of the date palm tree at the top, and then a pot attached to the tree collects drip of the nectar overnight<sup>30</sup>. Pteropus species bat consumes the nectar and infects it with saliva, urine, and excreta. At the same time, a human consumes the same nectar the virus transmits from bat to human<sup>31</sup>. Moreover, the virus can be transmitted by people climbing the poisonous date tree, eating contaminated fruits, and contacting sick animals and humans<sup>32</sup>. Domestic animals are being infected by eating contaminated saliva and mixing fruits<sup>33</sup>. A retrospective investigation was done in two villages in Bangladesh to identify clinical features, causative agents, risk factors, and natural and animal reservoirs. Immunoglobulin M for IgM antibodies and indirect EIA for IgG antibodies were tested in the Centers for Disease Control and Prevention (CDC) to affirm the diagnosis. Samples were taken from severely ill patients having encephalitis and identified Nipah virus<sup>34</sup>. The critical source of spreading the disease in Bangladesh is man-to-man transmission<sup>35</sup>. The largest outbreak occurred in 2004 by droplet infection and was identified in saliva<sup>36</sup>. The Nipah virus has been a flare up in Bangladesh since 2001. In 2009 another outbreak

occurred in Rajbari and Manikganj, and cases were reported of encephalitis and severe respiratory illness<sup>37</sup>. Between 2010 to 2012, multiple outbursts occurred in different places in northern Bangladesh, such as Faridpur and Lalmonirhat. The experts could not control the death from the disease; hence the fatality rate was high<sup>38</sup>. About 24 patients were detected having Nipah virus infection in 2013 and 21 of them died where male and female patients were 16 and 8 respectively, and the age was between 8 months to 60 years<sup>39</sup>.

**Table 1: Outbreaks of Nipah Virus<sup>40</sup> from 1998-2013**

Year	Country	State or District	Cases	Deaths	Case fatality
1998-1999	Malaysia	Perak, Selangor, Negeri Sembilan states	265	105	40%
1999	Singapore	Singapore	11	1	9%
2001	India	Siliguri district, West Bengal	66	49	74%
2001	Bangladesh	Meherpur district	13	9	69%
2003	Bangladesh	Naogaon district	12	8	67%
2004	Bangladesh	Faridpur and Rajbari districts	67	50	75%
2005	Bangladesh	Tangail district	12	11	92%
2007	Bangladesh	Thakurgaon, Naoga and Kushtia districts	18	9	50%
2007	India	Nadia district, West Bengal	5	5	100%
2008	Bangladesh	Manikganj, Rajbari and Faridpur district	11	9	82%
2009	Bangladesh	Rajbari, Gaibandha, Rangpur and Nilphamari districts	4	1	25%
2010	Bangladesh	Faridpur, Rajbari, Gopalganj and Madaripur districts	16	14	88%
2011	Bangladesh	Lalmonirhat, Dinajpur, Comilla, Nilphamari and Rangpur districts	44	40	91%
2012	Bangladesh	Joypurhat Rajshahi, Natore, Rajbari and Gopalganj districts	12	10	83%
2013	Bangladesh	Gaibandha, Jhainaidaha, Kurigram, Kushtia, Magura, Manikganj, Mymensingh, Naogaon, Natore, Nilphamari, Pabna, Rajbari and Rajshahi districts	24	21	87%

**India:** Two separate incidents of the Nipah virus were reported from 2001-2009 and occurred across the demarcation line from Bangladesh's Nipah zone<sup>41</sup>. Initially, in 2001, 11 infected patients were admitted into the hospital; for further treatment, the patient got transferred to another hospital, which caused the further infection, and 25 hospital staff and 8 visitors got infected<sup>42</sup>. During the first outbreak, 45 cases died among 66 patients, and during the subsequent outbreak, 4 cases were infected in 2007, with a 100.0% fatality rate<sup>43</sup>. In 2007, the first case had a history of taking alcohol made of date palm, and other subjects were infected by the first case<sup>38</sup>. The recent outbreak happened in 2018 in districts of Kerala named Kozhikode and Malappuram; 17 people dead among 18 affected people<sup>10</sup>. There was no history of Nipah virus infection in this area before and separate from the areas previously affected by the virus. The date juice was not typical in that area, and the spread of the infection was by a contacted health care professional<sup>44</sup>. The affected groups mainly were

from economic aged and different age groups. Moreover, all the outbreaks that occurred in India were human-to-human transmission<sup>45</sup>.

**Philippines:** Seventeen persons were identified for NiV infection in an island of the Philippines named Mindanao in 2014, where two deaths, eleven cases of encephalitis, five influenza-like illnesses, and one meningitis were recorded<sup>46</sup>. The Philippines National Epidemiology centre confirmed the testing for henipavirus, where antibodies against NiV and IgM against NiV were confirmed. The fatality rate was 53.0%, the rate was high (82.0%) in patients with acute encephalitis<sup>47</sup>. Around 10 cases had contact history with horses or meat of horses. At the same time, ten horses died having neurological symptoms; however, the horses were not tested for NiV infection. Five cases were infected through man-to-man transmission; among them, 2 cases were health care professionals<sup>48</sup>. Later, it revealed that the strain was like the Malaysian strain, either a mutation strain or collaboration or multiple strains<sup>49</sup>.

### Achieving Goal of SDGs

The United Nations agreed on an agenda 2030, Sustainable Development Goals (SDGs), consisting of 17 goals that focus on peace and prosperity in developing and developed countries<sup>50</sup>. SDGs aim to end poverty, inequality, and deprivation and improve the education system and economy. All the members will conserve the forest and ocean along with lives under oceans and intercept climate change<sup>51</sup>. Most of the emerging infectious diseases are arisen from wild animals and become infectious due to intermingling with domestic birds or animals. The occurrence of infectious diseases depends on the population size and wildlife varieties<sup>52</sup>. Nowadays, due to the destruction of forests and the growth of industries and agricultural land, the build-up of the animal farm hampering the climate and wildlife. Wild animals and birds are traveling to the urban area, searching for food, and transmitting the disease to local animals, birds, and humans<sup>53</sup>. For example, the beginning of the Nipah virus occurrence in 1998 happened in a pig farm located in the countryside area, near to tropical rainforest where the fruit bats (reservoir of Nipah virus) resided<sup>54</sup>. The bats are the reason for the emergence of the Nipah virus, Ebola virus, SARS, and areas affected by those viruses were the place for haunting wildlife and living place of wild animals<sup>55</sup>. Humans are being infected due to contact with wildlife or infected domestic animals. Multiple socioeconomic developments must be required,

including the strategy of SDGs need to imply, to minimize this problem<sup>56</sup>.

The goal 3 implies to confirm healthy lives and promoting health and well beings<sup>57</sup>. Infectious diseases like the Nipah virus increase the rate of hospital admission, morbidity, and mortality that impairs the goal three strategies<sup>58</sup>. Hence, to achieve goal three, infectious diseases need to be controlled. Moreover, there is a strong relation between infectious disease with environmental policy<sup>59</sup>. The destruction of forest and the extermination of wild animals from their inhabited region increases the chance of transmitting the virus and hamper the achievement of SDGs goal 15 that focuses on the existence of living species on land<sup>60</sup>. The main aim of this goal is to conserve and re-establish the forest that encourages earth ecologies<sup>61</sup>.

The purpose of SDGs 2 is to increase the productivity of crops and livestock that will improve the agricultural system, reduce food hunger, and increase food security among the people<sup>62</sup>. Around 2.37 billion people cannot have a healthy diet in the world<sup>63</sup>. During any pandemic, the rate of impoverished people increases; as a result, it is challenging for people to meet their basic needs<sup>64</sup>. Hence, a vast number of people encounter hunger. In 2020, almost 70 to 161 million had experienced starvation during the pandemic. The most vulnerable groups were the reproductive age group and children. One-third of total reproductive age women had anaemia due to having less nutritious food<sup>65</sup>. Additionally, some pig farms were destroyed previously due to the transmission of the Nipah virus from bat to pig and other domestic animals, which affected livestock productivity<sup>66</sup>. Hence, it can be said that infectious disease can obstruct achieving goal 2. However, to increase the productivity of crops and livestock, it may need to destroy the forest, wildlife life may be curtailed, and the risk of infectious diseases like the Nipah virus may increase<sup>67</sup>.

### **Socio Economic Determinants of Nipah virus Infection**

Workers from China and other Asian countries ran the farm in Peninsular Malaysia in 1998 and 1999. Farmers lived in the farmhouse, passed down through the family<sup>68</sup>. They sold the meat and food from the pigs in the local market and shipped them overseas to Singapore and other Asian countries. They also supplied the local markets<sup>69</sup>. This is on top of them lending a helping hand to the neighbourhood's bank and shops. Young adults

began to work in banks, shops, and farmhouses due to increased employment options<sup>70</sup>. The outbreak of the Nipah virus infection, on the other hand, had destroyed the company. Eight thousand five hundred direct pig farm workers, 9400 supported workers in pig farm industries, and 3,00,000 workers from other industries were affected during the Nipah Virus outbreak in Malaysia<sup>71</sup>. The pigs were infected and died because of the infection. It was recommended that the pigs be slaughtered, and the export of pigs to other countries was put on hold<sup>72</sup>. The virus had the most significant impact on farmers, who were either admitted to the hospital or died due to the infection. The death of the farmers had an impact on the family member who died and influenced the economy of the region<sup>73</sup>. The government encouraged to start off another agricultural and animal farmhouse and provided free medical services. The government offered a donation of 32 USD to the families of the pig farm; however, the amount was insufficient to stock up the pig farm business. Hence, the pig farm business and other local businesses were closed<sup>74</sup>. Local authorities collected funds from the public to help the devastated family. Around 91.0% of patients had received USD 6950, while half of the dead patients received 1400 USD for their funeral<sup>11</sup>.

The drink of date juice is popular among children and adults in Bangladesh during the winter season. Date juice is commonly used to make date jaggery and pithas a type of pancakes or fritters<sup>75</sup>. Local markets and other districts in Bangladesh used to be flooded with date juice, date jaggery, and pithas, which were sold by locals<sup>76</sup>. People became ill due to the Nipah virus outbreak and were admitted to hospitals, placing a significant financial strain on their families. The Nipah virus was also extremely contagious, as previously stated<sup>35</sup>. It had an impact on other family members and healthcare workers, which resulted in an increase in hospital admissions and a burden on hospitals due to the limited number of beds available<sup>77</sup>. Bangladesh is a country with a population that is moderately educated<sup>78</sup>. People began to isolate themselves, which had a negative impact on their families. Rather than assisting, they began to avoid family members, which had a negative impact on their social and psychological well-being<sup>38</sup>.

Domestic animals (cows and goats) were also affected and died due to the effects of the Nipah virus, which had an adverse economic impact on the local market<sup>15</sup>. Bats are known to consume fruits from the surrounding area<sup>79</sup>. Children in Bangladesh died after consuming bat fruit that had been partially digested<sup>80</sup>. Farmers and vendors at

the fruit market have also reported a low selling rate for their products<sup>81</sup>.

## Conclusion

According to aetiology and demography, it implies that, Nipah virus infection is a highly contagious zoonotic disease circulating for the past two decades. After thorough analysis it is discovered that, even though bats spread the initial outbreak to pigs, it later spread from bats to humans and from humans to humans. Epidemiological data shows that, many pig farms were destroyed due to NiV because of the high mortality and morbidity rates among animals and humans. Acute encephalitis and respiratory illness necessitate hospitalization and treatment on an emergency basis. To reduce the frequency of Nipah virus infection outbreaks, WHO, government agencies, and various social work organizations collaborate. However, outbreaks persist due to inadequate awareness about the spread of infection, poor education, a shortage of surveillance, an inappropriate isolation strategy, and poor hygiene maintenance.

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## Conflict of Interest

We declare that we have no conflict of interest.

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## Data Availability

Any inquiries regarding supporting data availability of this study should be directed to the corresponding author and are available from the corresponding author on reasonable request.

## Ethics Approval and Consent to Participate

None

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## References

1. Wang L, Crameri G. Emerging zoonotic viral diseases. *Rev Sci Tech*. 2014;33(2):569-81.
2. Pigott DM, Golding N, Mylne A, Huang Z, Weiss DJ, Brady OJ, et al. Mapping the zoonotic niche of Marburg virus disease in Africa. *Transactions of the Royal Society of Tropical Medicine and Hygiene*. 2015;109(6):366-78.
3. Roser M, Ritchie H, Ortiz-Ospina E, Hasell J. Coronavirus disease (COVID-19)—Statistics and research. *Our World in data*. 2020;4.
4. Xin H, Xue A, Liang J, Qiu B, Dong L, Pan B, et al. Global epidemiology and risk factors of nipah virus infection in human, 1998–2018. *Disease Surveillance* 2019;34(1):89-92.
5. Chua K, Bellini W, Rota P, Harcourt B, Tamin A, Lam S, et al. Nipah virus: a recently emergent deadly paramyxovirus. *Science*. 2000;288(5470):1432-5.
6. Epstein JH, Field HE, Luby S, Pulliam JR, Daszak P. Nipah virus: impact, origins, and causes of emergence. *Current infectious disease reports*. 2006;8(1):59-65.
7. Rathish B, Vaishnani K. Nipah Virus. *StatPearls [Internet]*. 2021.
8. Lam SK, Chua KB. Nipah virus encephalitis outbreak in Malaysia. *Clinical Infectious Diseases*. 2002;34(Supplement\_2):S48-S51.
9. Gupta B, Ahmed KM, Gupta R. Nipah Virus Research: A Scientometric Assessment of Global Publications Output during 1999-2018. *International Journal of Medicine and Public Health*. 2018;8(2).
10. Thayyil J, Padmanabhan A, Gangadharan A, Salim S, Jayakrishnan T. Nipah outbreak in Kerala, South India: Ethical challenges in the deployment of healthcare workers. *Indian Journal of Medical Ethics*. 2020;5(4).
11. Chua KB. Nipah virus outbreak in Malaysia. *Journal of Clinical Virology*. 2003;26(3):265-75.
12. Mazzola LT, Kelly-Cirino C. Diagnostics for Nipah virus: a zoonotic pathogen endemic to Southeast Asia. *BMJ global health*. 2019;4(Suppl 2):e001118.
13. Geisbert TW, Mire CE, Geisbert JB, Chan Y-P, Agans KN, Feldmann F, et al. Therapeutic treatment of Nipah virus infection in nonhuman primates with a neutralizing human monoclonal antibody. *Science translational medicine*. 2014;6(242):242ra82-ra82.
14. WHO. Nipah Virus Infection: World Health Organization; 2022 [Available from: [https://www.who.int/health-topics/nipah-virus-infection#tab=tab\\_1](https://www.who.int/health-topics/nipah-virus-infection#tab=tab_1)].
15. Hughes JM, Wilson ME, Luby SP, Gurley ES, Hossain MJ. Transmission of human infection with Nipah virus. *Clinical Infectious Diseases*. 2009;49(11):1743-8.
16. Bossart KN, Zhu Z, Middleton D, Klippel J, Crameri G, Bingham J, et al. A neutralizing human monoclonal antibody protects against lethal disease in a new ferret model of acute nipah virus infection. *PLoS pathogens*. 2009;5(10):e1000642.
17. Ang BSP, Lim TCC, Wang L, Kraft CS. Nipah Virus Infection. *Journal of Clinical Microbiology*. 2018;56(6):e01875-17.
18. Homaira N, Rahman M, Hossain MJ, Epstein JH, Sultana R, Khan MSU, et al. Nipah virus outbreak with person-to-person transmission in a district of Bangladesh, 2007. *Epidemiology and Infection*. 2010;138(11):1630-6.
19. Lam S-K. Nipah virus—a potential agent of bioterrorism? *Antiviral Research*. 2003;57(1):113-9.
20. Paton NI, Leo YS, Zaki SR, Auchus AP, Lee KE, Ling AE, et al. Outbreak of Nipah-virus infection among abattoir workers in Singapore. *Lancet*. 1999;354(9186):1253-6.
21. Av R, Sadanandan S, Thulaseedharan NK, Kg SK, Pallivalappil B, As AK. Nipah Virus Infection. *J Assoc Physicians India*. 2018;66(12):58-60.

22. Soman Pillai V, Krishna G, Valiya Veetil M. Nipah Virus: Past Outbreaks and Future Containment. *Viruses*. 2020;12(4).
23. Looi LM, Chua KB. Lessons from the Nipah virus outbreak in Malaysia. *Malays J Pathol*. 2007;29(2):63-7.
24. Yong MY, Lee SC, Ngui R, Lim YA, Phipps ME, Chang LY. Seroprevalence of Nipah Virus Infection in Peninsular Malaysia. *J Infect Dis*. 2020;221(Suppl 4):S370-s4.
25. Gomez Román R, Wang LF, Lee B, Halpin K, de Wit E, Broder CC, et al. Nipah@20: Lessons Learned from Another Virus with Pandemic Potential. *mSphere*. 2020;5(4).
26. Chew MH, Arguin PM, Shay DK, Goh KT, Rollin PE, Shieh WJ, et al. Risk factors for Nipah virus infection among abattoir workers in Singapore. *J Infect Dis*. 2000;181(5):1760-3.
27. Kulkarni DD, Tosh C, Venkatesh G, Senthil Kumar D. Nipah virus infection: current scenario. *Indian J Virol*. 2013;24(3):398-408.
28. Sharma V, Kaushik S, Kumar R, Yadav JP, Kaushik S. Emerging trends of Nipah virus: A review. *Rev Med Virol*. 2019;29(1):e2010.
29. Goh GK, Dunker AK, Foster JA, Uversky VN. Nipah shell disorder, modes of infection, and virulence. *Microb Pathog*. 2020;141:103976.
30. Banerjee S, Gupta N, Kodan P, Mittal A, Ray Y, Nischal N, et al. Nipah virus disease: A rare and intractable disease. *Intractable Rare Dis Res*. 2019;8(1):1-8.
31. Clayton BA. Nipah virus: transmission of a zoonotic paramyxovirus. *Curr Opin Virol*. 2017;22:97-104.
32. Hsu VP, Hossain MJ, Parashar UD, Ali MM, Ksiazek TG, Kuzmin I, et al. Nipah virus encephalitis reemergence, Bangladesh. *Emerging infectious diseases*. 2004;10(12):2082.
33. Luby SP, Rahman M, Hossain MJ, Blum LS, Husain MM, Gurley E, et al. Foodborne transmission of Nipah virus, Bangladesh. *Emerging infectious diseases*. 2006;12(12):1888.
34. Mahmood-ul-Hassan M, Salim M. Public perceptions about the fruit bats in two horticulturally important districts of Pakistan. *Journal of Animal and Plant Sciences*. 2011;21(2).
35. Lo MK, Lowe L, Hummel KB, Sazzad HM, Gurley ES, Hossain MJ, et al. Characterization of Nipah virus from outbreaks in Bangladesh, 2008–2010. *Emerging infectious diseases*. 2012;18(2):248.
36. Nikolay B, Salje H, Hossain MJ, Khan AD, Sazzad HM, Rahman M, et al. Transmission of Nipah virus—14 years of investigations in Bangladesh. *New England Journal of Medicine*. 2019;380(19):1804-14.
37. Rahman M, Chakraborty A. Nipah virus outbreaks in Bangladesh: a deadly infectious disease. *WHO South-East Asia Journal of Public Health*. 2012;1(2):208.
38. Dhillon J, Banerjee A. Controlling Nipah virus encephalitis in Bangladesh: Policy options. *Journal of Public Health Policy*. 2015;36(3):270-82.
39. Nahar N, Asaduzzaman M, Sultana R, Garcia F, Paul RC, Abedin J, et al. A large-scale behavior change intervention to prevent Nipah transmission in Bangladesh: components and costs. *BMC research notes*. 2017;10(1):1-13.
40. Giangaspero M, Al Ghafri M. Poaching: A Threat for Vulnerable Wild Animal Species in Oman. *Tropical Medicine & Surgery*. 2014;02.
41. Paul L. Nipah virus in Kerala: a deadly Zoonosis. *Clin Microbiol Infect*. 2018;24(10):1113-4.
42. Mourya D, Yadav P, Rout M, Pattnaik B, Shete A, Patil D. Absence of Nipah virus antibodies in pigs in Mizoram State, North East India. *Indian J Med Res*. 2019;149(5):677-9.
43. Raza A, Awrejcewicz J, Rafiq M, Mohsin M. Breakdown of a Nonlinear Stochastic Nipah Virus Epidemic Models through Efficient Numerical Methods. *Entropy (Basel)*. 2021;23(12).
44. Epstein JH, Anthony SJ, Islam A, Kilpatrick AM, Ali Khan S, Balkey MD, et al. Nipah virus dynamics in bats and implications for spillover to humans. *Proc Natl Acad Sci U S A*. 2020;117(46):29190-201.
45. Singh RK, Dhama K, Chakraborty S, Tiwari R, Natesan S, Khandia R, et al. Nipah virus: epidemiology, pathology, immunobiology and advances in diagnosis, vaccine designing and control strategies - a comprehensive review. *Vet Q*. 2019;39(1):26-55.
46. Donaldson H, Lucey D. Enhancing preparation for large Nipah outbreaks beyond Bangladesh: Preventing a tragedy like Ebola in West Africa. *Int J Infect Dis*. 2018;72:69-72.
47. Ambat AS, Zubair SM, Prasad N, Pundir P, Rajwar E, Patil DS, et al. Nipah virus: A review on epidemiological characteristics and outbreaks to inform public health decision making. *J Infect Public Health*. 2019;12(5):634-9.
48. Broder CC, Weir DL, Reid PA. Hendra virus and Nipah virus animal vaccines. *Vaccine*. 2016;34(30):3525-34.
49. Aditi u, Shariff M. Nipah virus infection: A review. *Epidemiology and Infection*. 2019;147:e95.
50. Loomis RJ, Stewart-Jones GB, Tsybovsky Y, Caringal RT, Morabito KM, McLellan JS, et al. Structure-based design of Nipah virus vaccines: a generalizable approach to paramyxovirus immunogen development. *Frontiers in immunology*. 2020;11:842.
51. Di Marco M, Baker ML, Daszak P, De Barro P, Eskew EA, Godde CM, et al. Opinion: Sustainable development must account for pandemic risk. *Proceedings of the National Academy of Sciences*. 2020;117(8):3888-92.
52. ADR ADR, Act AH. International Case Studies in the Management of Disasters. *Crisis*. 2020;10:11.
53. Prada D. Viral ecology of Western Australian microbat communities: Murdoch University; 2020.
54. Hauser N, Gushiken AC, Narayanan S, Kottilil S, Chua JV. Evolution of Nipah Virus Infection: Past, Present, and Future Considerations. *Tropical Medicine and Infectious Disease*. 2021;6(1):24.
55. Wood JL, Leach M, Waldman L, MacGregor H, Fooks AR, Jones KE, et al. A framework for the study of zoonotic disease emergence and its drivers: spillover of bat pathogens as a case study. *Philosophical Transactions of the Royal Society B: Biological Sciences*. 2012;367(1604):2881-92.
56. Sun B, Jia L, Liang B, Chen Q, Liu D. Phylogeography, transmission, and viral proteins of Nipah virus. *Virologica Sinica*. 2018;33(5):385-93.
57. Budhathoki SS, Pokharel PK, Good S, Limbu S, Bhattachan M, Osborne RH. The potential of health literacy to address the health related UN sustainable development goal 3 (SDG3) in Nepal: a rapid review. *BMC Health Services Research*. 2017;17(1):237.
58. Guegana J-F, Suzán G, Kati-Coulibaly S, Bonpangue DN, Moatti J-P. Sustainable Development Goal# 3, “health and well-being”, and the need for more integrative thinking. *Veterinaria México*. 2018;5(2):1-18.
59. Perrings C, Castillo-Chavez C, Chowell G, Daszak P, Fenichel EP, Finnoff D, et al. Merging Economics and Epidemiology to Improve the Prediction and Management of Infectious Disease. *EcoHealth*. 2014;11(4):464-75.
60. Yonehara A, Saito O, Hayashi K, Nagao M, Yanagisawa R, Matsuyama K. The role of evaluation in achieving the SDGs. *Sustainability Science*. 2017;12(6):969-73.
61. Liu S, Bai J, Chen J. Measuring SDG 15 at the county scale: Localization and practice of SDGs indicators based on geospatial information. *ISPRS International Journal of Geo-Information*. 2019;8(11):515.
62. Gil JDB, Reidsma P, Giller K, Todman L, Whitmore A, van Ittersum M. Sustainable development goal 2: Improved targets and indicators for agriculture and food security. *Ambio*. 2019;48(7):685-98.
63. Perez-Escamilla R. Food security and the 2015–2030 sustainable development goals: From human to planetary health: Perspectives and opinions. *Current developments in nutrition*. 2017;1(7):e000513.

64. Ahmed S, Byker Shanks C. Supporting Sustainable Development Goals Through Sustainable Diets. In: Leal Filho W, Wall T, Azul AM, Brandli L, Özuyar PG, editors. *Good Health and Well-Being*. Cham: Springer International Publishing; 2020. p. 688-99.
65. Ezirigwe J, Ojike C, Amechi E, Adewopo A. 'COVID-19/Food Insecurity Syndemic': Navigating the Realities of Food Security Imperatives of Sustainable Development Goals in Africa. *Law and Development Review*. 2021;14(1):129-62.
66. Mohd Nor MN, Gan CH, Ong BL. Nipah virus infection of pigs in peninsular Malaysia. *Revue scientifique et technique* (International Office of Epizootics). 2000;19(1):160-5.
67. Plowright RK, Eby P, Hudson PJ, Smith IL, Westcott D, Bryden WL, et al. Ecological dynamics of emerging bat virus spillover. *Proceedings of the royal society B: biological sciences*. 2015;282(1798):20142124.
68. Uppal P. Emergence of Nipah virus in Malaysia. *Annals of the New York Academy of Sciences*. 2000;916(1):354-7.
69. Chua K. Epidemiology, surveillance and control of Nipah virus infections in Malaysia. *The Malaysian journal of pathology*. 2010;32(2):69-73.
70. Hosono H, Kono H, Ito S, Shirai J. Economic impact of Nipah virus infection outbreak in Malaysia. *Proceedings of the 11th ISVEE*. 2006.
71. Johara MY, Field H, Rashdi AM, Morrissy C, van der Heide B, Rota P, et al. Nipah virus infection in bats (order Chiroptera) in peninsular Malaysia. *Emerging Infectious Diseases*. 2001;7(3):439-41.
72. Tan K-S, Tan CT, Goh KJ. Epidemiological aspects of Nipah virus infection. *Neurol J Southeast Asia*. 1999;4(1):77-81
73. Lim C, Lee W, Leo Y, Lee K, Chan K, Ling A, et al. Late clinical and magnetic resonance imaging follow up of Nipah virus infection. *Journal of Neurology, Neurosurgery & Psychiatry*. 2003;74(1):131-3.
74. Ahmad K. Malaysia culls pigs as Nipah virus strikes again. *The Lancet*. 2000;356(9225):230.
75. Vengaiiah P, Ravindrababu D, Murthy G, Prasad K. Jaggery from Palmyrah palm (*Borassus flabellifer* L.)-Present status and scope. 2013.
76. Chowdhury MSH, Halim MA, Muhammed N, Haque F, Koike M. Traditional utilization of wild date palm (*Phoenix sylvestris*) in rural Bangladesh: an approach to sustainable biodiversity management. *Journal of forestry research*. 2008;19(3):245-51.
77. Hossain MJ, Gurley ES, Montgomery JM, Bell M, Carroll DS, Hsu VP, et al. Clinical presentation of nipah virus infection in Bangladesh. *Clinical infectious diseases*. 2008;46(7):977-84.
78. Rahman T, Nakata S, Nagashima Y, Rahman M, Sharma U, Rahman M. Bangladesh tertiary education sector review. 2019.
79. Giangaspero M. Nipah virus. *Trop Med Surg*. 2013;1(129):2.
80. Siddique AB, Fardows J, Farhana N, Mazumder M. Nipah Virus: a public health concern. *Journal of Enam Medical College*. 2016;6(2):101-5.
81. Nahar N, Paul RC, Sultana R, Sumon SA, Banik KC, Abedin J, et al. A controlled trial to reduce the risk of human Nipah virus exposure in Bangladesh. *Ecohealth*. 2017;14(3):501-17