**UNIVERSITY OF SINDH JOURNAL OF ANIMAL SCIENCES** 



DOI: <u>https://doi.org/10.57038/usjas.v7i04.6642</u> Uni. Sindh. J. Anim. Sci., vol. 7(4); 13-22, 2023 Email: <u>editors.usjas@usindh.edu.pk</u> ISSN (P): 2521-8328 ISSN (E): 2523-6067 Published by University of Sindh, Jamshoro.

# ZOONOTIC SPILLOVER AND THEIR RELATIVITY TO PANDEMICS

# MUHAMMAD ABDUL MANAN<sup>1</sup>, AYESHA KHAN<sup>1</sup>, AREEBA SAFDAR<sup>2</sup>, MUHAMMAD FARHAN KHAN<sup>3,4\*</sup>

<sup>1</sup>University of the Punjab, Department of Zoology, Lahore Pakistan <sup>2</sup>Department of Zoology, Bahauddin Zakariya University, Multan, Pakistan <sup>3</sup>College of international studies, Beibu gulf University, Qinzhou, China <sup>4</sup>Department of Chemistry, Gomal University Dera Ismail Khan, Pakistan

#### ARTICLE INFORMATION

## ABSTRACT

Article History: Received: 23<sup>rd</sup> September 2022 Accepted: 21<sup>st</sup> October 2023 Published online: 31<sup>st</sup> December 2023

#### Author's contribution

MAM and AK wrote manuscript, AS revised manuscript and MFK conceived idea and revised Manuscript.

#### Key words:

Anthropogenesis, Bat, Close contact, Pandemics, Prevent, Spillovers The study aims to emphasize the relativity of pandemics and their zoonotic spillover. During recent decades the chance of pandemics occurring has increased many folds. So, the study thoroughly reviewed seven zoonotic spillovers from bats in history that led us to the common feature that close contact of bats with other species including humans is the major reason for pandemics. Due to anthropogenesis bats started moving into buildings for survival, reproduction, and swarming became the susceptible source of disease transmission. The study aims to suggest measures that could help to prevent bat-borne viral outbreaks in the future.

# 1. INTRODUCTION

The ongoing wave of Coronavirus disease-19 (COVID-19) pandemic around the globe caused by SARS-CoV-2 became a global public health concern (Jin *et al.*, 2020, Rothan & Byrareddy, 2020). It is obvious that bats are the convenient reservoir of the CoVs and probably the etiological agent of COVID-19 (Sabir *et al.*, 2016).

In the past century, Asia, Africa, and Arabian states were the major hotspot regions related to bat-born viral pandemics in human society (Calisher *et al.*, 2006, Wolfe *et al.*, 2005, Morse, 2004). We studied seven zoonotic spillovers that are Marburg virus (Towner *et al.*, 2009, Nyakarahuka *et al.*, 2016), Ebola virus (Smith, 1978) Nipah virus (Paton *et al.*, 1999, Chua *et al.*, 2000), Hendra virus (Blum *et al.*, 2009), Severe Acute Respiratory Syndrome coronavirus in (Christian *et al.*, 2004) Middle East respiratory syndrome (Skariyachan *et al.*, 2019) and COVID-19 (Jin *et al.*, 2020).

\*Corresponding Author: <u>farhankhanbgu@gmail.com</u>

When we studied briefly these spillovers that resulted in the growth of the human population intensified the anthropogenic interference to wildlife (Plowright et al., 2011). Wild animals are the potential hosts of viral diseases as in the case of Hendra, Nipah, and SARS-CoVs (Chua et al., 2000, Philbey et al., 1998, Murray et al., 1995a). Similarly, the reason behind the EBOV and SARS-CoV outbreaks is the cultural traditions of eating wild animal meat or 'bush meat' in Africa and Asia (Dowell et al., 1999). Businesses in live markets such as the desire for eating wild bats conceded the susceptible animals to come into contact with bats and let the transmission of SARS-CoV into the human's population (Webster, 2004, Guan et al., 2003a, Lam et al., 2003). On the contrary, in the case of MERS-CoV outbreak civets dromedary camels were identified as major intermediate hosts that were got infected elsewhere from the reservoir host (Zaki et al., 2012).

Copyright 2017 University of Sindh Journal of Animal Sciences

Moreover, the anthropogenic activities including deforestation and urban development with the increasing human population selecting synanthropic bat species, and so far the bats are benefiting from living close to humans, thereby increasing their numbers and the risk of disease transmission to humans (McFarlane et al., 2012). The origin of the COVID-19 pandemic outbreak is still debatable but the very first cases of concentrated pneumonia were informed in Wuhan city of China and were linked with the wet animal market. Because of high human to human transmission, the COVID-19 rapidly spread to other countries of the world and global health emergency was announced by WHO (Jin et al., 2020, Rothan and Byrareddy, 2020). The aim of our study is to curb pandemics in future. So that, we thoroughly reviewed seven zoonotic spillovers from bats in history that led us to common feature that is close contact of bats with other species including human is the major reason of pandemics. Due to anthropization bats started moving into building for survival, reproduction and swarming that became the susceptible source of disease transmission. Our suggestions could help to prevent bat-borne viral outbreaks in future if seriously implemented.

#### Zoonotic Spillover from bats

Spread of viruses via bats causing EIDS (Emerging infectious disease) is a major issue. A list of bats born dreadful pandemics/epidemics outbreak in human populations all over the world is given in Table 1 and Figure 1. The mode of transmissions of viral pandemics/epidemics to the human population from different reservoirs is represented in Figure 2. This figure also explained that bat close contact in outbreak areas becomes the source of disease transmission. Our suggestions could help to prevent bat-borne viral outbreaks in future if seriously implemented listed below.

#### Why bat research always our week area?

Americans were on the belief that bats are violent enough to attack human and pets even organization which worked for bat conservation used to avoid bats because of fear (Tuttle, 1979). Therefore, deep research is much needed in broad spectrum which should be able to cover all aspects of bats.

#### Minimize anthropogenic activities

Area of forests has been lessened and environmental diversity is nearer to human population due to anthropization. Unlike their natural habitat, these landscapes provided a large number of bats to make their habitat in this anthropized environment. Bats found there niches in human populated areas which are their alternative environment for their survival (Walsh *et al.*, 2017a, Afelt *et al.*, 2018).In the name of urbanization, large scale deforestation has been done that threatens and disturbing the natural habitats, bats started changing their habitat and moved to urban areas that made close contact with humans. If we do reforestation bats can be revived to their original habitat and it will also help the ecosystem, whereas the problem of the increasing population can be controlled by the social campaign, social awareness, education, and proper planning.

#### Detecting synanthropic bat species

Most of the species are living in man-made buildings in which bats are using different spaces of these buildings which are good element for hibernation and reproduction (Lesinski, 2006). Searching for bats in the buildings and its premises is hard task even for the chiropterologists therefore infrared cameras, endoscopes and ultrasonic detectors can be used to find bats but more inovative technological equipment can be used with the help of chiroptrologists to capture bats.

#### Chiropterological inventory survey

Roofs, wall cracks, attics can be examined and inspected with use of modern equipment's. It is necessary step that should be practiced during any inventory of bats in buildings and its premises (Janus & Lesinski, 2018). A survey must take place with modern techniques and equipment's with high probability detection rate in new buildings as well as existing buildings and its premises including all spaces, roofs, wall cracks and all possible places.

#### Create rehabilitees for bats

Killing and capturing of bats has become extensive and not limited in buildings. Major colonies of bats (*Myotis* grisescens) were burnt in caves (Tuttle, 1979) when health officials conjectured i that these species of Bats were outbreak cause rabies in foxes (Fredrickson & Thomas, 1965). This information makes bats very unsafe. Killing and capturing of bats has become extensive and not limited in buildings. Major colonies of bats (*Myotis* grisescens) were burnt in caves (Tuttle, 1979) when health officials conjectured that these species of Bats were outbreak cause rabies in foxes (Fredrickson & Thomas, 1965). Historically, there are always reasons and objects of fear and conflict that came across many societies possibly due to their dark and evasive behavior (Kingston, 2016). Capturing and killing bats or keep them away from buildings is not the solution. We need to make rehabilitation areas for bats in forests, mountains, areas far away from human population where these captured bats can revive, live, hibernate and swarm. By reviving bats and saving its population is necessary as bats play important role in our ecosystem.

#### Management Implications

Our study can support wildlife management based on bats conservation and its rehabilitation.

## 2. CONFLICT OF INTEREST

All authors have declared that there is no conflict of interests regarding the publication of this article.

## REFRENCES

- Adjemian, J., Farnon, E. C., Tschioko, F., Wamala, J. F., Byaruhanga, E., Bwire, G. S., Kansiime, E., Kagirita, A., Ahimbisibwe, S., & Katunguka, F. (2011). Outbreak of Marburg hemorrhagic fever among miners in Kamwenge and Ibanda Districts, Uganda, 2007. *The Journal of Infectious Diseases*, 204, 796-799.
- Afelt, A., Lacroix, A., Zawadzka-Pawlewska, U., Pokojski, W., Buchy, P., & Frutos, R. (2018). Distribution of bat-borne viruses and environment patterns. *Infection, Genetics and Evolution*, 58, 181-191.
- Amman, B. R., Carroll, S. A., Reed, Z. D., Sealy, T. K., Balinandi, S., Swanepoel, R., Kemp, A., Erickson, B. R., Comer, J. A., & Campbell, S. (2012). Seasonal pulses of Marburg virus circulation in juvenile *Rousettus aegyptiacus* bats coincide with periods of increased risk of human infection. *PLoS Pathog*, 8, e1002877.
- Arunkumar, G., Chandni, R., Mourya, D. T., Singh, S. K., Sadanandan, R., Sudan, P., & Bhargava, B. (2019). Outbreak investigation of Nipah virus disease in Kerala, India, 2018. *The Journal of Infectious Diseases*, 219, 1867-1878.
- Bausch, D. G., Borchert, M., Grein, T., Roth, C., Swanepoel, R., Libande, M. L., Talarmin, A., Bertherat, E., Muyembe-Tamfum, J.-J., & Tugume, B. (2003). Risk factors for Marburg hemorrhagic

fever, Democratic Republic of the Congo. *Emerging Infectious Diseases*, 9, 1531.

- Beer, B., Kurth, R., & Bukreyev, A. (1999). Characteristics of Filoviridae: Marburg and Ebola viruses. *Naturwissenschaften*, 86, 8-17.
- Blum, L. S., Khan, R., Nahar, N., & Breiman, R. F. (2009). In-depth assessment of an outbreak of Nipah encephalitis with person-to-person transmission in Bangladesh: Implications for prevention and control strategies. *The American Journal of Tropical Medicine and Hygiene*, 80, 96-102.
- Bonaparte, M., Dimitrov, A., Bossart, K., Crameri, G., Mungall, B., Bishop, K., Choudhry, V., Dimitrov, D., Wang, L.-F., Eaton, B., & Broder, C. (2005). Ephrin-B2 Ligand is a Functional Receptor for Hendra Virus and Nipah Virus. *Proceedings of the National Academy of Sciences of the United States of America*, 102, 10652-7.
- Borba, M. G. S., Val, F. F. A., Sampaio, V. S., Alexandre, M. A. A., Melo, G. C., Brito, M., Mourão, M. P. G., Brito-Sousa, J. D., Baía-da-Silva, D., & Guerra, M. V. F. (2020). Effect of high vs low doses of chloroquine diphosphate as adjunctive therapy for patients hospitalized with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection: A randomized clinical trial. *JAMA Network Open*, *3*, e208857-e208857.
- Brearley, G., Rhodes, J., Bradley, A., Baxter, G., Seabrook, L., Lunney, D., Liu, Y., & McAlpine, C. (2013). Wildlife disease prevalence in humanmodified landscapes. *Biology Reviews of the Cambridge Philosophical Society*, 88, 427-442.
- Calisher, C. H., Childs, J. E., Field, H. E., Holmes, K. V., & Schountz, T. (2006). Bats: Important reservoir hosts of emerging viruses. *Clinical Microbiology Reviews*, 19, 531-545.
- CDC. (2020). MERS Clinical Features [Online]. Available: <u>https://www.cdc.gov/coronavirus/mers/clinical-features.html</u> [Accessed].
- Chan, J. F., Yuan, S., Kok, K. H., To, K. K., Chu, H., Yang, J., Xing, F., Liu, J., Yip, C. C., Poon, R. W., Tsoi, H. W., Lo, S. K., Chan, K. H., Poon, V. K., Chan, W. M., Ip, J. D., Cai, J. P., Cheng, V. C., Chen, H., Hui, C. K., & Yuen, K. Y. (2020). A familial cluster of pneumonia associated with the 2019 novel coronavirus indicating person-to-person transmission: A study of a family cluster. *The Lancet*, 395, 514-523.

- Christian, M. D., Poutanen, S. M., Loutfy, M. R., Muller, M. P., & Low, D. E. (2004). Severe acute respiratory syndrome. *Clinical Infectious Diseases*, 38, 1420-1427.
- Chua, K., Bellini, W., Rota, P., Harcourt, B., Tamin, A., Lam, S., Ksiazek, T., Rollin, P., Zaki, S., & Shieh, W.-J. (2000). Nipah virus: A recently emergent deadly paramyxovirus. *Science*, 288, 1432-1435.
- Dowell, S. F., Mukunu, R., Ksiazek, T. G., Khan, A. S., Rollin, P. E., & Peters, C. J. (1999). Transmission of Ebola hemorrhagic fever: A study of risk factors in family members, Kikwit, Democratic Republic of the Congo, 1995. *The Journal of Infectious Diseases*, 179, 87-91.
- Fredrickson, L. E., & Thomas, L. (1965). Relationship of fox rabies to caves. *Public Health Reports*, 80, 495.
- Giles, J. R., Eby, P., Parry, H., Peel, A. J., Plowright, R. K., Westcott, D. A., & McCallum, H. (2018). Environmental drivers of spatiotemporal foraging intensity in fruit bats and implications for Hendra virus ecology. *Scientific Reports*, 8, 9555.
- Guan, Y., Zheng, B., He, Y., Liu, X., Zhuang, Z., Cheung, C., Luo, S., Li, P., Zhang, L., & Guan, Y. (2003a). Isolation and characterization of viruses related to the SARS coronavirus from animals in southern China. *Science*, 302, 276-278.
- Guan, Y., Zheng, B. J., He, Y. Q., Liu, X. L., Zhuang, Z. X., Cheung, C. L., Luo, S. W., Li, P. H., Zhang, L. J., Guan, Y. J., Butt, K. M., Wong, K. L., Chan, K. W., Lim, W., Shortridge, K. F., Yuen, K. Y., Peiris, J. S., & Poon, L. L. (2003b). Isolation and characterization of viruses related to the SARS coronavirus from animals in southern China. Science, 302, 276-8.
- Hui, D. S., Azhar, E. I., Madani, T. A., Ntoumi, F., Kock, R., Dar, O., Ippolito, G., McHugh, T. D., Memish, Z. A., & Drosten, C. (2020a). The continuing 2019nCoV epidemic threat of novel coronaviruses to global health—The latest 2019 novel coronavirus outbreak in Wuhan, China. *International Journal of Infectious Diseases*, 91, 264-266.
- Hui, D. S., E, I. A., Madani, T. A., Ntoumi, F., Kock, R., Dar, O., Ippolito, G., McHugh, T. D., Memish, Z. A., Drosten, C., Zumla, A., & Petersen, E. (2020b). The continuing 2019-nCoV epidemic threat of novel coronaviruses to global health—The latest 2019 novel coronavirus outbreak in Wuhan, China.

International Journal of Infectious Diseases, 91, 264-266.

- Janus, K., & Lesinski, G. (2018). Birds and bats using buildings as a place of breeding or shelter. Annals of Warsaw University of Life Sciences-SGGW. Animal Science, 57.
- Jin, Y.-H., Cai, L., Cheng, Z.-S., Cheng, H., Deng, T., Fan, Y.-P., Fang, C., Huang, D., Huang, L.-Q., & Huang, Q. (2020). A rapid advice guideline for the diagnosis and treatment of 2019 novel coronavirus (2019nCoV) infected pneumonia (standard version). *Military Medical Research*, 7, 4.
- Jones, B. A., Grace, D., Kock, R., Alonso, S., Rushton, J., Said, M. Y., McKeever, D., Mutua, F., Young, J., McDermott, J., & Pfeiffer, D. U. (2013). Zoonosis emergence linked to agricultural intensification and environmental change. *Proc Natl Acad Sci U S A*, 110, 8399-404.
- Kakodkar, P., Kaka, N., & Baig, M. N. (2020). A Comprehensive Literature Review on the Clinical Presentation, and Management of the Pandemic Coronavirus Disease 2019 (COVID-19). *Cureus*, 12, e7560.
- Kandeil, A., Gomaa, M., Shehata, M., El-Taweel, A., Kayed, A. E., Abiadh, A., Jrijer, J., Moatasim, Y., Kutkat, O., & Bagato, O. (2019). Middle East respiratory syndrome coronavirus infection in noncamelid domestic mammals. *Emerging Microbes & Infections*, 8, 103-108.
- Kingston, T. (2016). Cute, Creepy, or Crispy—How values, attitudes, and norms shape human behavior toward bats. In *Bats in the Anthropocene: conservation of bats in a changing world* (pp. 571-588). Springer International AG, Cham.
- Lam, W., Zhong, N., & Tan, W. (2003). Overview on SARS in Asia and the world. *Respirology*, 8, S2-S5.
- Leligdowicz, A., Fischer, W. A., Uyeki, T. M., Fletcher, T. E., Adhikari, N. K., Portella, G., Lamontagne, F., Clement, C., Jacob, S. T., & Rubinson, L. (2016). Ebola virus disease and critical illness. *Critical Care*, 20, 217.
- Leroy, E. M., Epelboin, A., Mondonge, V., Pourrut, X., Gonzalez, J.-P., Muyembe-Tamfum, J.-J., & Formenty, P. (2009). Human Ebola outbreak resulting from direct exposure to fruit bats in Luebo, Democratic Republic of Congo, 2007. Vector-borne and Zoonotic Diseases, 9, 723-728.

- Lesinski, G. (2006). Wplyw antropogenicznych przekstalcen krajobrazu na strukture i funkcjonowanie zespolow nietoperzy w Polsce [Influence of anthropogenic landscape changes on the structure and functioning of bat communities in Poland]. SGGW.
- Li, W., Shi, Z., Yu, M., Ren, W., Smith, C., Epstein, J. H., Wang, H., Crameri, G., Hu, Z., Zhang, H., Zhang, J., McEachern, J., Field, H., Daszak, P., Eaton, B. T., Zhang, S., & Wang, L. F. (2005). Bats are natural reservoirs of SARS-like coronaviruses. *Science*, 310, 67-69.
- Lim, P. L., Kurup, A., Gopalakrishna, G., Chan, K. P., Wong, C. W., Ng, L. C., Se-Thoe, S. Y., Oon, L., Bai, X., Stanton, L. W., Ruan, Y., Miller, L. D., Vega, V. B., James, L., Ooi, P. L., Kai, C. S., Olsen, S. J., Ang, B., & Leo, Y. S. (2004). Laboratory-acquired severe acute respiratory syndrome. *N Engl J Med*, 350, 1740-1745.
- Lu, R., Zhao, X., Li, J., Niu, P., Yang, B., Wu, H., Wang, W., Song, H., Huang, B., Huang, L., Zhu, N., Bi, Y., Ma, X., Zhan, F., Wang, L., Hu, T., Zhou, H., Hu, Z., Zhou, W., Zhao, L., Chen, J., Meng, Y., Wang, J., Lin, Y., Yuan, J., Xie, Z., Ma, J., Liu, W. J., Wang, D., Xu, W., Holmes, E. C., Gao, G. F., Wu, G., Chen, W., Shi, W., & Tan, W. (2020). Genomic characterisation and epidemiology of 2019 novel coronavirus: implications for virus origins and receptor binding. *Lancet*, 395, 565-574.
- Luby, J. P., & Sanders, C. V. (1969). Green monkey disease ("Marburg virus" disease): a new zoonosis. *American College of Physicians*.
- Malvy, D., McElroy, A. K., De Clerck, H., Günter, S., & Van Griensven, J. (2019). Ebola virus disease. *The Lancet*, *393*, 936-948.
- McFarlane, R., Sleigh, A., & McMichael, T. (2012). Synanthropy of wild mammals as a determinant of emerging infectious diseases in the Asian– Australasian region. *EcoHealth*, *9*, 24-35.
- McMullan, L. K., Flint, M., Chakrabarti, A., Guerrero, L., Lo, M. K., Porter, D., Nichol, S. T., Spiropoulou, C.
  F., & Albariño, C. (2019). Characterisation of infectious Ebola virus from the ongoing outbreak to guide response activities in the Democratic Republic of the Congo: a phylogenetic and in vitro analysis. *The Lancet Infectious Diseases, 19*, 1023-1032.
- Morse, S. (2004). Factors and determinants of disease emergence. *Revue scientifique et technique-Office international des épizooties*, 23, 443-452.

- Murray, K., Selleck, P., Hooper, P., Hyatt, A., Gould, A., Gleeson, L., Westbury, H., Hiley, L., Selvey, L., & Rodwell, B. (1995a). A morbillivirus that caused fatal disease in horses and humans. *Science*, 268, 94-97.
- Murray, K., Selleck, P., Hooper, P., Hyatt, A., Gould, A., Gleeson, L., Westbury, H., Hiley, L., Selvey, L., Rodwell, B., et al. (1995b). A morbillivirus that caused fatal disease in horses and humans. *Science*, 268, 94-97.
- Negrete, O. A., Levroney, E. L., Aguilar, H. C., Bertolotti-Ciarlet, A., Nazarian, R., Tajyar, S., & Lee, B. (2005). EphrinB2 is the entry receptor for Nipah virus, an emergent deadly paramyxovirus. *Nature*, 436, 401-405.
- Nyakarahuka, L., Kankya, C., Krontveit, R., Mayer, B., Mwiine, F. N., Lutwama, J., & Skjerve, E. (2016). How severe and prevalent are Ebola and Marburg viruses? A systematic review and meta-analysis of the case fatality rates and seroprevalence. *BMC Infectious Diseases, 16*, 1-14.
- Nyakarahuka, L., Shoemaker, T. R., Balinandi, S., Chemos, G., Kwesiga, B., Mulei, S., Kyondo, J., Tumusiime, A., Kofman, A., & Masiira, B. (2019). Marburg virus disease outbreak in Kween District Uganda, 2017: Epidemiological and laboratory findings. *PLoS Neglected Tropical Diseases*, 13, e0007257.
- Ong, S. W. X., Tan, Y. K., Chia, P. Y., Lee, T. H., Ng, O. T., Wong, M. S. Y., & Marimuthu, K. (2020). Air, surface environmental, and personal protective equipment contamination by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) from a symptomatic patient. *Jama*, 323, 1610-1612.
- Paton, N. I., Leo, Y. S., Zaki, S. R., Auchus, A. P., Lee, K. E., Ling, A. E., Chew, S. K., Ang, B., Rollin, P. E., & Umapathi, T. (1999). Outbreak of Nipah-virus infection among abattoir workers in Singapore. *The Lancet*, 354, 1253-1256.
- Perlman, S., & Netland, J. (2009). Coronaviruses post-SARS: update on replication and pathogenesis. *Nature Reviews Microbiology*, 7, 439-450.
- Philbey, A. W., Kirkland, P. D., Ross, A. D., Davis, R. J., Gleeson, A. B., Love, R. J., ... & Hyatt, A. D. (1998).
  An apparently new virus (family Paramyxoviridae) infectious for pigs, humans, and fruit bats. *Emerging infectious diseases*, 4(2), 269.

- Plowright, R. K., Foley, P., Field, H. E., Dobson, A. P., Foley, J. E., Eby, P., & Daszak, P. (2011). Urban habituation, ecological connectivity and epidemic dampening: the emergence of Hendra virus from flying foxes (Pteropus spp.). *Proceedings of the Royal Society B: Biological Sciences, 278*(1725), 3703-3712.
- Rothan, H. A., & Byreddy, S. N. (2020). The epidemiology and pathogenesis of coronavirus disease (COVID-19) outbreak. *Journal of autoimmunity, 102433*.
- Sabir, J. S., Lam, T. T.-Y., Ahmed, M. M., Li, L., Shen, Y., Abo-Aba, S. E., Qureshi, M. I., Abu-Zeid, M., Zhang, Y., & Khiyami, M. A. (2016). Co-circulation of three camel coronavirus species and recombination of MERS-CoVs in Saudi Arabia. *Science*, 351(6268), 81-84.
- Saeed, A. A. B., Abedi, G. R., Alzahrani, A. G., Salameh, I., Abdirizak, F., Alhakeem, R., Algarni, H., El Nil, O. A., Mohammed, M., & Assiri, A. M. (2017). Surveillance and testing for Middle East respiratory syndrome coronavirus, Saudi Arabia, April 2015– February 2016. *Emerging infectious diseases*, 23(4), 682.
- Shehata, M. M., Gomaa, M. R., Ali, M. A., & Kayali, G. (2016). Middle East respiratory syndrome coronavirus: a comprehensive review. *Frontiers of medicine*, 10, 120-136.
- Sissoko, D., Duraffour, S., Kerber, R., Kolie, J. S., Beavogui, A. H., Camara, A.-M., Colin, G., Rieger, T., Oestereich, L., & Pályi, B. (2017). Persistence and clearance of Ebola virus RNA from seminal fluid of Ebola virus disease survivors: a longitudinal analysis and modelling study. *The Lancet Global Health*, 5(1), e80-e88.
- Skariyachan, S., Challapilli, S. B., Packirisamy, S., Kumargowda, S. T., & Sridhar, V. S. (2019). Recent aspects on the pathogenesis mechanism, animal models and novel therapeutic interventions for Middle East respiratory syndrome coronavirus infections. *Frontiers in microbiology*, 10, 569.
- Smith, D. (1978). Ebola haemorrhagic fever in Sudan, 1976. Bulletin of the World Health Organization, 56(2), 247-270.
- Symons, R. (2011). Re: Canine case of Hendra virus and Hendra virus vaccine. *Australian Veterinary Journal*, 89(11), 442.

- Timen, A., Koopmans, M. P., Vossen, A. C., van Doornum, G. J., Günther, S., van den Berkmortel, F., Verduin, K. M., Dittrich, S., Emmerich, P., & Osterhaus, A. D. (2009). Response to imported case of Marburg hemorrhagic fever, the Netherlands. *Emerging infectious diseases*, 15(8), 1171.
- Towner, J. S., Amman, B. R., Sealy, T. K., Carroll, S. A. R., Comer, J. A., Kemp, A., Swanepoel, R., Paddock, C. D., Balinandi, S., & Khristova, M. L. (2009). Isolation of genetically diverse Marburg viruses from Egyptian fruit bats. *PLoS Pathogens*, 5(7), e1000536.
- Tuttle, M. D. (1979). Status causes of decline, and management of endangered gray bats. *The Journal of Wildlife Management*, 43(1), 1-17.
- Walsh, M. G., Wiethoelter, A., & Haseeb, M. (2017a). The impact of human population pressure on flying fox niches and the potential consequences for Hendra virus spillover. *Scientific reports*, 7, 1-13.
- Walsh, M. G., Wiethoelter, A., & Haseeb, M. A. (2017b). The impact of human population pressure on flying fox niches and the potential consequences for Hendra virus spillover. *Scientific Reports*, 7, 8226.
- Wang, N., Shi, X., Jiang, L., Zhang, S., Wang, D., Tong, P., Guo, D., Fu, L., Cui, Y., & Liu, X. (2013). Structure of MERS-CoV spike receptor-binding domain complexed with human receptor DPP4. *Cell research*, 23(8), 986-993.
- Webster, R. G. (2004). Wet markets—a continuing source of severe acute respiratory syndrome and influenza? *The Lancet*, 363(9404), 234-236.
- World Health Organization. (2015). Preliminary Clinical Description of Severe Acute Respiratory Syndrome [Online]. Available: <u>https://www.who.int/csr/sars/clinical/en</u> [Accessed 23-11-2020].
- World Health Organization. (2018). Marburg virus disease [Online]. Available: <u>https://www.who.int/news-room/fact-</u> <u>sheets/detail/marburg-virus-disease</u>
- World Health Organization. (2020a). Coronavirus disease (COVID-19) [Online]. Available: <u>https://www.who.int/emergencies/diseases/novel-</u> <u>coronavirus-2019</u> [Accessed 23-11-2020].
- World Health Organization. (2020b). Ebola virus disease [Online]. Available: <u>https://www.who.int/newsroom/fact-sheets/detail/ebola-virus-disease</u> [Accessed 23-11-2020].

- Wolfe, N. D., Daszak, P., Kilpatrick, A. M., & Burke, D. S. (2005). Bushmeat hunting, deforestation, and prediction of zoonotic disease. *Emerging infectious diseases*, 11(12), 1822.
- Xu, L., Zhang, X., Song, W., Sun, B., Mu, J., Wang, B.,
  & Dong, X. (2020). Conjunctival polymerase chain reaction-tests of 2019 novel coronavirus in patients in Shenyang, China. *medRxiv*.
- Young, P. L., Halpin, K., Selleck, P. W., Field, H., Gravel, J. L., Kelly, M. A., & MacKenzie, J. S. (1996). Serologic evidence for the presence in Pteropus bats of a paramyxovirus related to equine morbillivirus. *Emerging infectious diseases*, 2(3), 239-240.
- Zaki, A. M., van Boheemen, S., Bestebroer, T. M., Osterhaus, A. D., & Fouchier, R. A. (2012). Isolation of a novel coronavirus from a man with pneumonia in Saudi Arabia. *New England Journal of Medicine*, 367(19), 1814-1820.



Figure 1: Visual map for the zoonotic spillover

Note: Yellow are representing the bat borne virus in maps

Virus Name	Origin/ Geographical location	Epidemic potential/ Time period	Target Species /Zoonosis	Symptoms	Incubation Period (Days)	Targeted human's receptor	Causes	Mode of Transmission	Death rate	References
MVD	Marburg and Frankfurt, Germany and Belgrade, Yugoslavia (present-day Serbia)	1967-2017	Bat, Primates, Human	hemorrhagic fever headache chills muscle ache	2-21	TIM-1	Visitor, Experimental study on infected monkeys	Infected Research Areas, Eco-tourism	81%	(Nyakarahuka <i>et al.</i> , 2019, Lam <i>et al.</i> , 2003, LUBY and SANDERS, 1969, Amman <i>et al.</i> , 2012, Sissoko <i>et al.</i> , 2017, Timen <i>et al.</i> , 2009, Adjemian <i>et al.</i> , 2011, Bausch <i>et al.</i> , 2003, Beer <i>et al.</i> , 1999, WHO, 2018)
EBOV	Sudan	1976-2018	Bat Primates Pigs Human	high fever malaise fatigue and body aches	2-21	TIM-1	Lovers of game meat, Dealing with wild animals	Hunters, Poor medical management	41.7%	(Arunkumar <i>et al.</i> , 2019) (Leroy <i>et al.</i> , 2009) (McMullan <i>et al.</i> , 2019) (Malvy <i>et al.</i> , 2019) (Leligdowicz <i>et al.</i> , 2016) (WHO, 2020b)
NiV	Malaysia/ Singapore	1990-2018	Bat Pig Human	neurological and respiratory	6-14	EFNB2	Agricultural intensification, Globalized economy	Workers on pig farms, Sale of infected Pigs to another region Persistence of pathogen increased the transmission in pigs and to humans.	39%	(Arunkumar <i>et al.</i> , 2019) (Negrete <i>et al.</i> , 2005) (Bonaparte <i>et al.</i> , 2005) (Arunkumar <i>et al.</i> , 2019, Paton <i>et al.</i> , 1999, Chua <i>et al.</i> , 2000)
HeV	Hendra, Queensland, Australi	1994-2012	Bat Horses Human	influenza-like illness multiorgan failure and	21	EFNB2	Climate change, Habitat loss,	Bats movement in agricultural land Urbanization	57%	(Murray <i>et al.</i> , 1995b) (Symons, 2011) (Brearley <i>et al.</i> , 2013) (Jones <i>et al.</i> , 2013) (Giles <i>et al.</i> , 2018)

# Table 1. A list of bat born dreadful pandemics / epidemics outbreak in human populations all over the world

Zoonotic Spillover: Bridging the Gap to Pandemics

				progressive encephalitis			Habitat fragmentation			(Walsh <i>et al.</i> , 2017b) (Young <i>et al.</i> , 1996)
SARS- CoV-1 (SARS)	Guangdong, China	November 2002-July 2003	Bat Civet cat Human	fever malaise myalgia headache diarrhea shivering cough and breath shortness	2-7	ACE2	Hunting, Dealers of wild animals in markets International travel	Butchers, Wildlife animal husbandry, Infected Research Areas	10.9%	(Hui <i>et al.</i> , 2020b) (Kakodkar <i>et al.</i> , 2020) (Lu <i>et al.</i> , 2020) (Chan <i>et al.</i> , 2020) (Li <i>et al.</i> , 2005) (Webster, 2004) (Guan <i>et al.</i> , 2003b) (Lim <i>et al.</i> , 2004) (WHO, 2015)
MERS- CoV (MERS)	Saudi Arabia	July 2012- 2018	Bat Camel Human	fever cough and shortness of breath common cold chills headache joint pain	5	DPP4	International travel, contact with camels	Consumption of camel meat or milk	35%	(Skariyachan et al., 2019) (Kakodkar et al., 2020) (Zaki et al., 2012) (Wang et al., 2013) (Perlman and Netland, 2009) (Saeed et al., 2017) (Shehata et al., 2016) (Kandeil et al., 2019) (CDC, 2020)
SARS- CoV-2 (COVID -19)	Wuhan, China	December 2019- Not controlled yet	Bat Pangolin Human	fever cough and shortness of breath	2-14	ACE2	sale of bush meat in seafood market	Fecal-oral path, Droplet transmission, Conjunctiva Fomites	4.9%	(Jin <i>et al.</i> , 2020) (Borba <i>et al.</i> , 2020) (Hui <i>et al.</i> , 2020a) (Ong <i>et al.</i> , 2020) (Zhang <i>et al.</i> , 2020) (WHO, 2020a)

*Manan et al.*, (2023)



Figure 2: The mode of transmissions of viral pandemics/epidemics to the human population from different reservoirs is represented.