

Human activity is responsible for animal viruses crossing over and causing pandemics

May 13 2020, by Narveen Jandu



Credit: AI-generated image ([disclaimer](#))

Viruses are naturally occurring entities. [Viruses have existed on Earth long before humans](#) and [vastly outnumber humans](#). There are [more viruses on Earth](#) than there are [stars in the universe](#) or [cells in the human body](#).

As a cellular microbiologist who has studied [the origin and development of infectious diseases and their prevention](#), understanding where infectious agents come from is as important, if not more important, to understanding how to combat the rampant spread of diseases within the [human population](#).

Fortunately, only a small fraction—[about 200](#)—of this vast array of [viruses](#) can infect humans. Some of the better-known [human](#) viral infections include measles, varicella, polio, [human papilloma virus](#), influenza and rhinoviruses, which are typically responsible for the common cold.

An even smaller number of viruses are responsible for the deadliest human infections that we have experienced. In recent decades these include rabies virus, [human immunodeficiency virus](#) (HIV), Ebola virus and now, infamously, coronaviruses.

Viruses that circulate in other animals can enter a human population when [a variety of human activities](#) allow for consistent and regular interaction with naturally occurring reservoirs. These events involve repeated and routine [interaction of humans with these animal hosts](#).

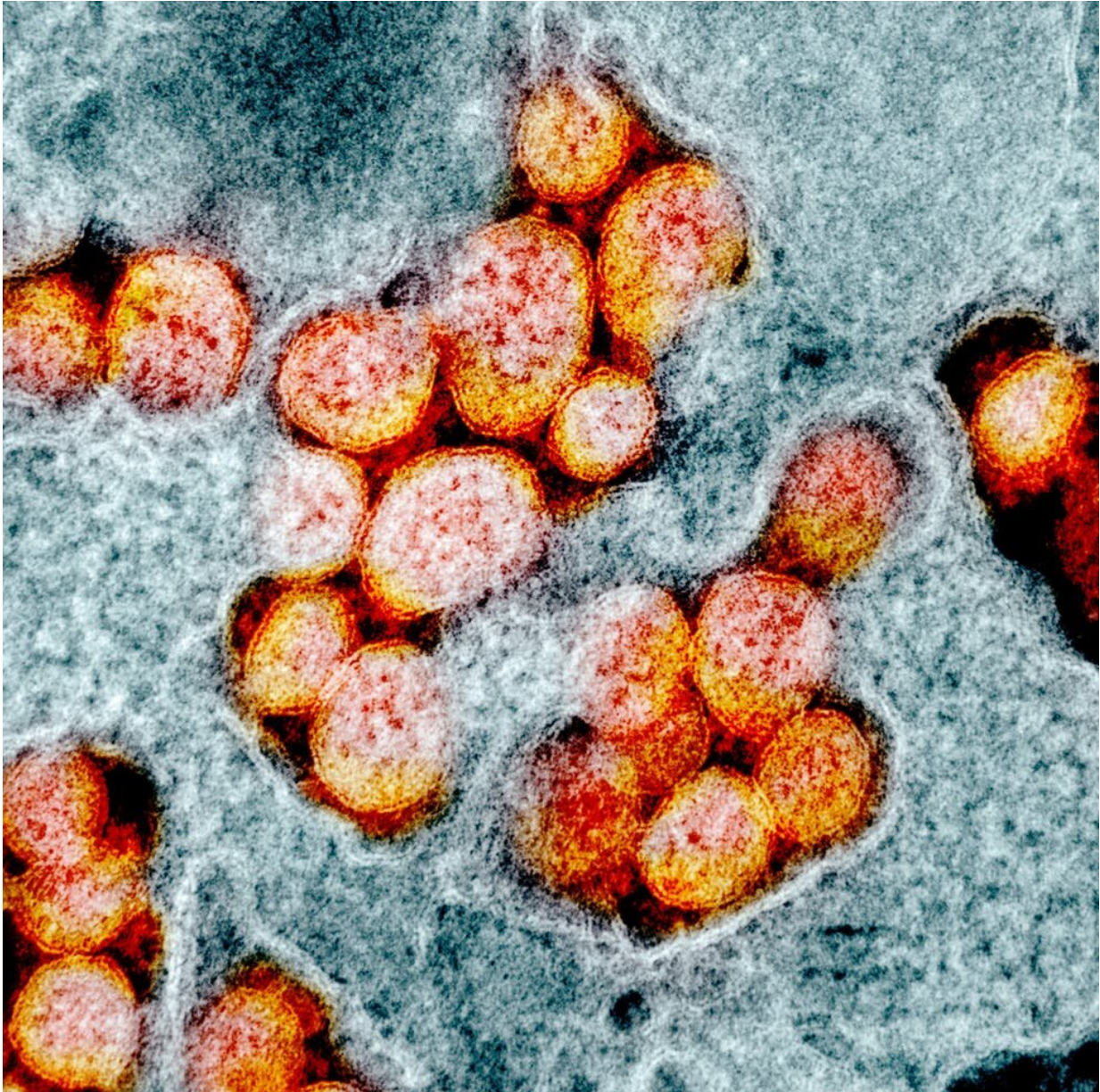
Some of these interactions take place through the following [human activities](#): hunting, butchering and farming (husbandry), as well as the global trade of animals and domestication of exotic animals as pets. Population growth, global travel and [climate change](#) that cause the disruption of habitats further provide opportunities for cross-species transfer.

Cross-species transfer

Many of the viruses that have affected us over the past 20 years have emerged from non-human reservoirs. Reservoirs are the source of

viruses and other pathogens and can be located in animal populations or the natural environment. What is of note is that [the original reservoir for many of these viruses are bats.](#)

Bats have been shown to be the [natural reservoir](#) of numerous deadly human viruses. The specialized immune systems of bats allow multiple different types of viruses to persist within these hosts. The coexistence of these [viruses within the same host](#) in combination with the molecular makeup of viruses further allows for the emergence of deadly human diseases. Bats aren't all bad though; they have an important role in our ecosystem.



A colour-enhanced image of SARS-CoV-2 virus particles, isolated from a patient. Credit: National Institute of Allergy and Infectious Diseases Integrated Research Facility

Current COVID-19 pandemic

The current [coronavirus pandemic has been traced to a wet market, called the Huanan Seafood Wholesale Market, in Wuhan, China](#). Within these wet markets, numerous animals are held together in confined spaces and small cages. This setting is inherently stressful and can allow for the exchange and mixing of multiple different bodily fluids. Human handling of these animals allows for [the viruses contained within these reservoirs to spill over into the human population](#). This is further facilitated by the long working hours and stress of the handlers themselves, as well as cuts, scratches, bites or other wounds inflicted by the animals on their handlers.

Read more: [Coronavirus: live animals are stressed in wet markets, and stressed animals are more likely to carry diseases](#)

A similar confluence of factors was attributed to the 2002 outbreak of SARS, [which has been attributed to the virus crossing over from bats to civets](#). The related MERS-CoV was [attributed to a bat reservoir in Saudi Arabia](#).

Ebola outbreaks

Outbreaks of Ebola in West Africa (2014-16 and 2018-present) have been attributed [to human interaction with bats](#). Three subfamilies of fruit bats—*Hypsignathus monstrosus*, *Epomops franqueti* and *Myonycteris torquata*—[have been identified as natural reservoirs for the Ebola virus](#).

These bats [inhabit caves deep within the Ebola River Valley](#). Cross-species transmission of the Ebola virus from bats to humans can occur either directly or via intermediary hosts such as non-human primates, horses or pigs.

Global travel allowed for the virus to [spread from Central Africa to West Africa, and subsequently Europe and North America](#).



Credit: AI-generated image ([disclaimer](#))

The Hendra virus

The Hendra virus (HeV) has been a persistent infection of [horses and humans in Australia since 1994](#). Transmission into the equine population has been attributed to [food or water contaminated with bat feces, urine or saliva](#). HeV originated in black flying foxes, and its transmission to horses and humans results in a severe respiratory disease.

Subsequent transmission from horses to humans involves the close contact of humans with the [bodily fluids of infected horses](#). Fortunately, [horse-to-horse, human-to-horse and bat-to-human transmission of hendra virus \(HeV\) is currently limited](#). As a result, [this viral infection has not lead to widespread infections or fatalities](#).

The Nipah virus

Nipah virus (NiV) infections have been occurring [in Bangladesh and Malaysia since 1998](#). This virus causes respiratory disease and swelling of the brain—encephalitis—in pigs. Human contact with infected pigs results in severe encephalitis, fever and eventually death. This [virus](#) also emerged from fruit bats through a spill-over event with the [close proximity of living conditions](#) between humans, domesticated [animals](#) and the natural [reservoir](#).

Altogether, it is clear that human activities play a role in the emergence or re-emergence of infectious diseases. The role of [bats as natural reservoirs](#) to numerous deadly human diseases is also very clear.

The uncertainty of subsequent outbreaks or pandemics is not related to how or why but rather a question of when.

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Citation: Human activity is responsible for animal viruses crossing over and causing pandemics (2020, May 13) retrieved 4 May 2024 from <https://medicalxpress.com/news/2020-05-human-responsible-animal-viruses-pandemics.html>

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