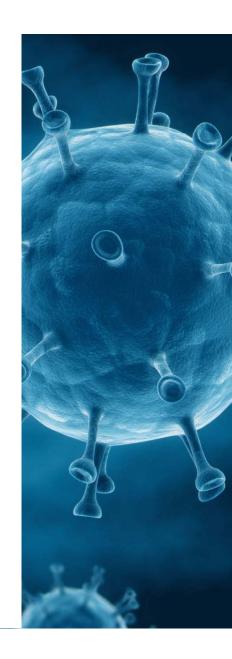
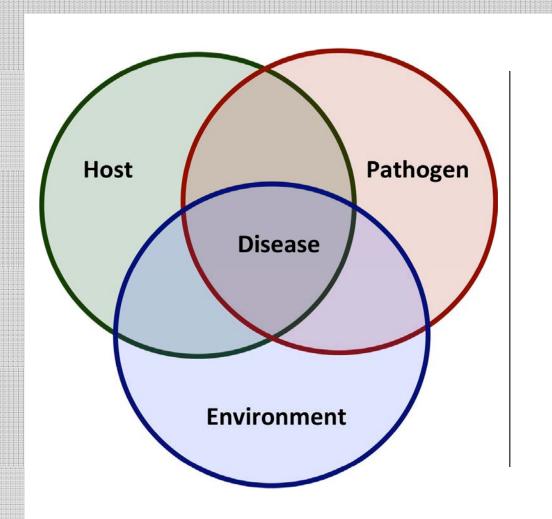
Environmental Impact on the Viruses & the diseases

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Mangla Hospital & Research Center,
BIJNOR-2646701 (UP)





Pathogen factors

- ✓ Infectious dose
- ✓ Shedding before symptoms
- ✓ Immune evasion

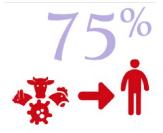
Human factors

- ✓ Crowding
- ✓ Promiscuity

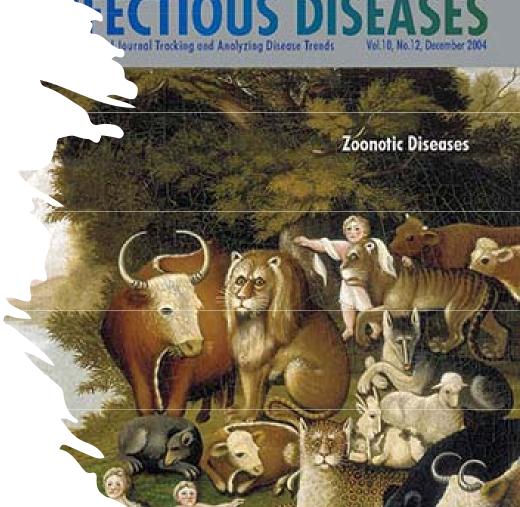
Environmental factors

✓ Poor health care





of emerging diseases in human are of animal origin



https://wwwnc.cdc.gov/eid/article/10/12/ac-1012_article

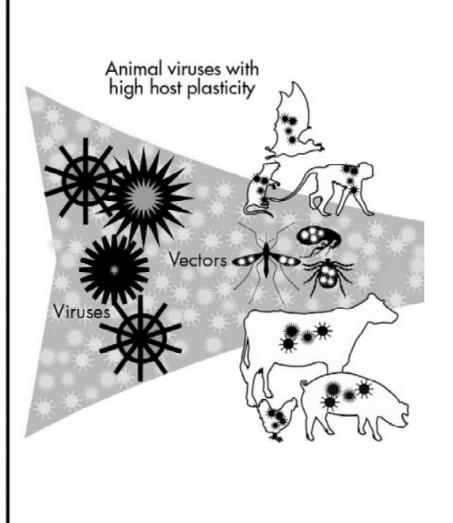
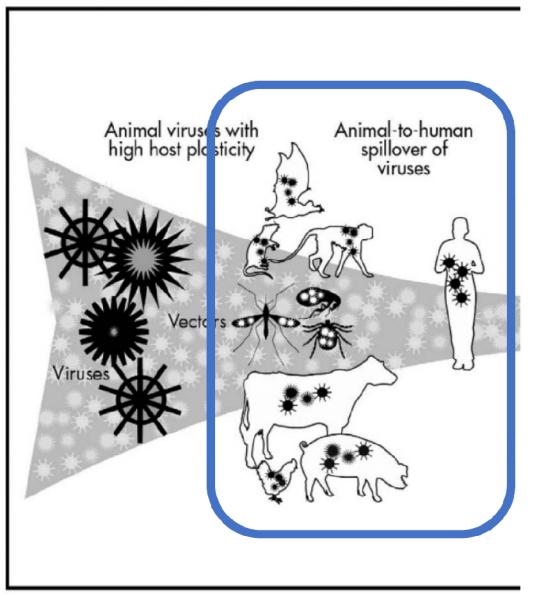
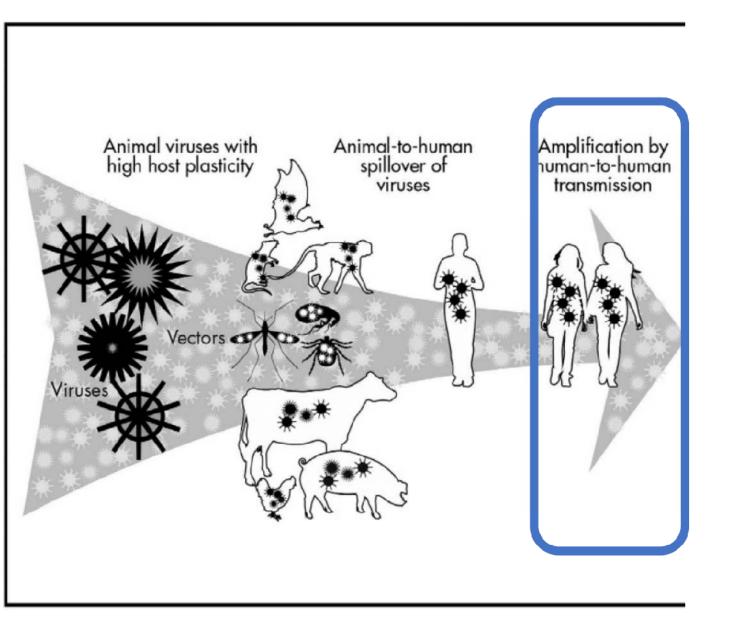
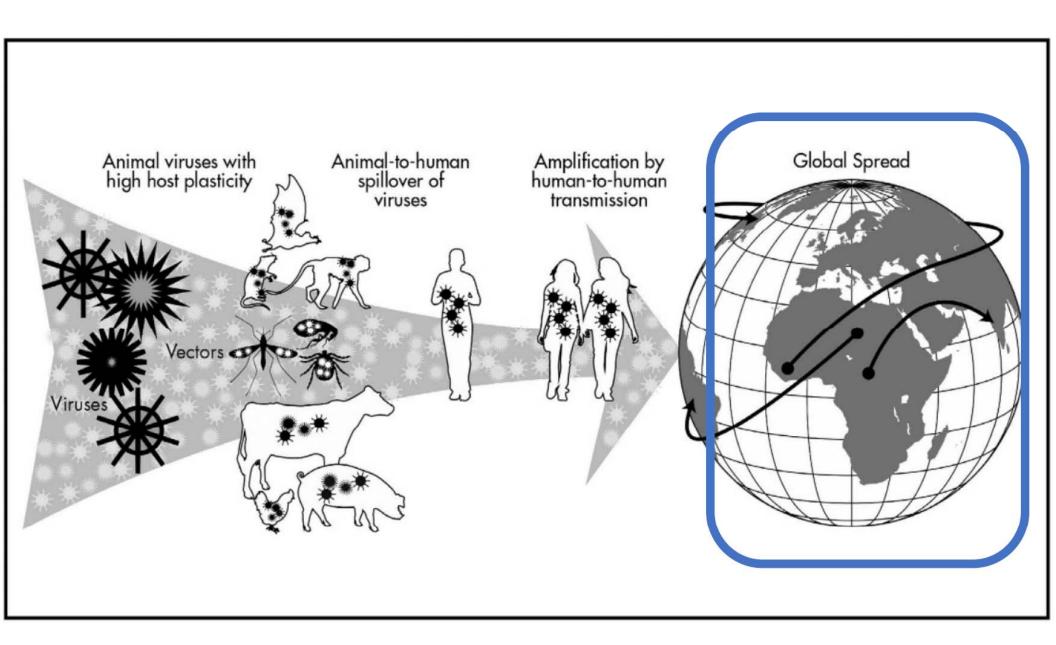


Figure 1. Pandemic properties of zoonotic viruses that spill over from animals to humans and spread by secondary transmission among humans. Key characteristics of pandemic potential that were evaluated

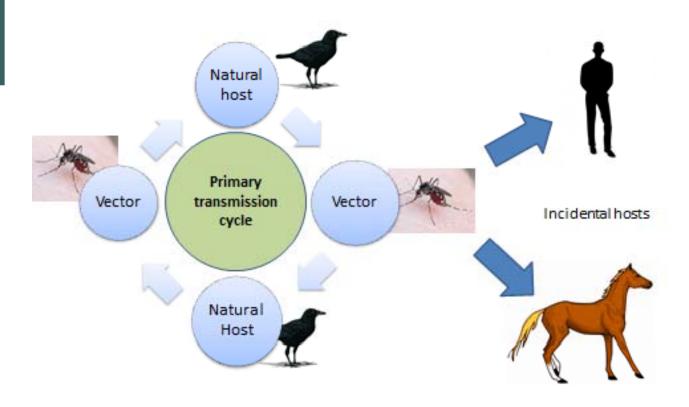
Kreuder et al 2015 Scientific Report







Transmission
Cycle of Vector
borne zoonoses

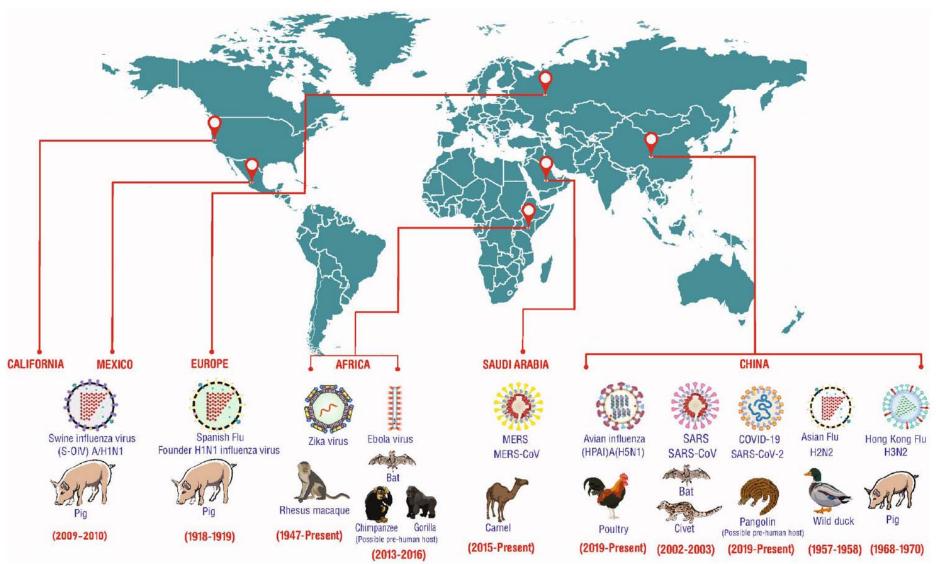


Major global & regional vector-borne diseases

Disease	Pathogen	Primary vector(s)	Primary non-human reservoir (competent) hosts
Malaria	Plasmodium parasite	Anopheles mosquito	Non-human hosts of minor concern
Dengue*	Flavivirus	Aedes aegypti and Aedes albopictus mosquitoes	Non-human hosts of minor concern
Yellow fever	Flavivirus	A. aegypti and A. albopictus mosquitoes	Non-human primates
Zika	Flavivirus	A. aegypti and A. albopictus mosquitoes	Non-human hosts of minor concern
Chikungunya*	Alphavirus	A. aegypti and A. albopictus mosquitoes	Non-human hosts of minor concern
Lymphatic filariasis*	Various filarial nematodes	A variety of mosquito genera	Non-human hosts of minor concern
Schistosomiasis*	Schistosoma trematode	Snail	Non-human hosts of minor concern
Onchocerciasis*	Onchocerca volvulus nematode	Simulium (black fly)	None
Chagas disease*	Trypanosoma cruzi parasite	Triatomine bug	Mammals
Leishmaniasis*	Leishmania parasite	Sand fly	Rodents, dogs, other mammals
Japanese encephalitis	Flavivirus	Culex mosquitoes	Pigs, birds
African trypanosomiasis*	Trypanosoma brucei parasite	Glossina (tsetse fly)	Wild and domestic animals
Lyme disease	Borrelia spirochete	Ixodes ticks	White-footed mouse and other small mammals, birds
Tick-borne encephalitis	Flavivirus	Ixodes ticks	Small rodents
West Nile fever	Flavivirus	Culex mosquitoes	Birds

https://www.nature.com/articles/s41590-020-0648-y.pdf

Major pandemic/outbreaks in the recorded history



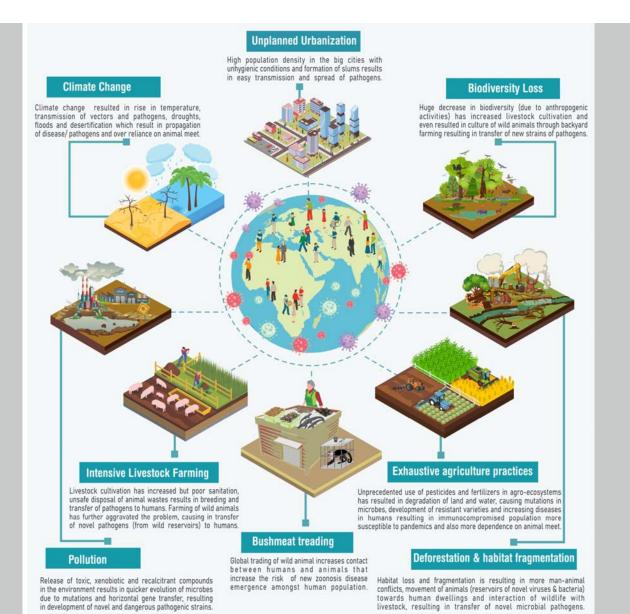
https://link.springer.com/article/10.1007/s42398-021-00165-x

Why is there frequent emergence of novel zoonotic outbreaks?

- Changing global environment
- Destruction of natural habitats of wild animals, possible reservoirs of novel human pathogen
- An increase in man-animal conflicts is being directly

In one of its report, FAO mentioned that >70% of human diseases are of animal origin, & have a direct relation with humans' reliance on animals for food

https://link.springer.com/article/10.1007/s42398-021-00165-x



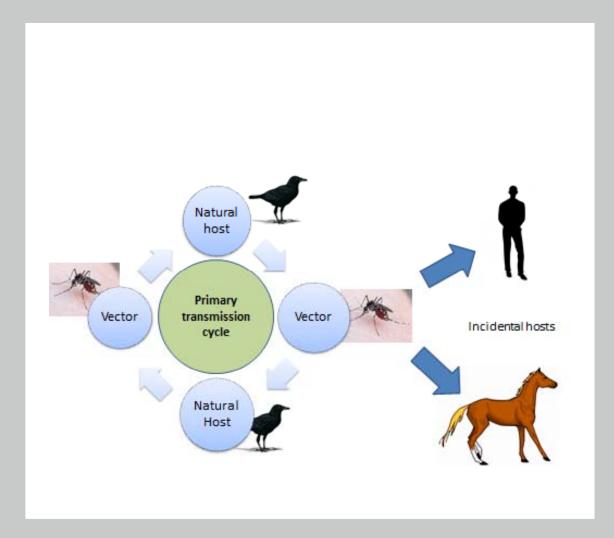
Environmental issues that lead to the pandemic

https://link.springer.com/article/10.1007/s42398-021-00165-x

How are the environmental factors becoming important in relation to the emergence & spread of novel viruses?

Climatic conditions

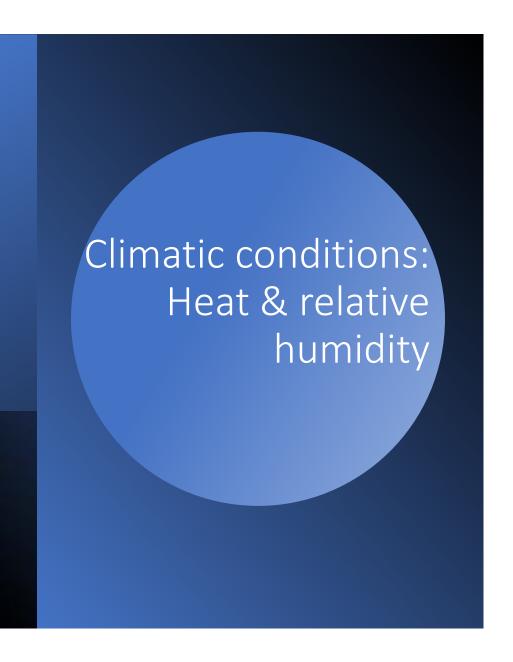
 In tropics change in climatic conditions have led to the intensive spread of the carriers/vectors of zoonotic viruses which increased the transition of viruses from their carrier to the animal host



Armon R, Cheruti U (2012) Environmental aspects of zoonotic diseases. IWA Publishing,

Impact of climatic conditions

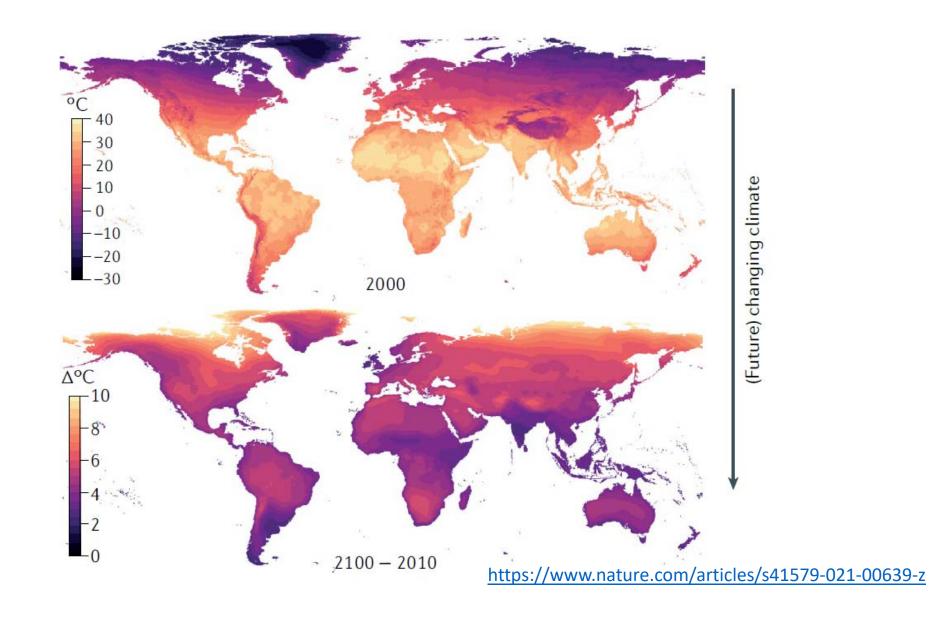
- WNV: Changes in temperature caused an adaptive mutation in the West Nile virus (WNV) in North America to efficiently propagate virus at elevated temperatures
- Brault AC (2009) Changing patterns of West Nile virus transmission: altered vector competence and host susceptibility. Vet Res 40:43
- MERS-CoV: 56.1% of the positive cases aroused between April to August, & the viability of the virus was favored by high temperature, high ultraviolet index, low wind speed, and low relative humidity
- Altamimi A, Ahmed AE (2019). J Infect Public Health 13:704–708



SARS-CoV-2:

 Most studies on the effects of temperature, and relative humidity on disease incidence and spread remain to be contradictory or inconclusive, with data showing positive as well as negative relation or in some researches no relationship at all

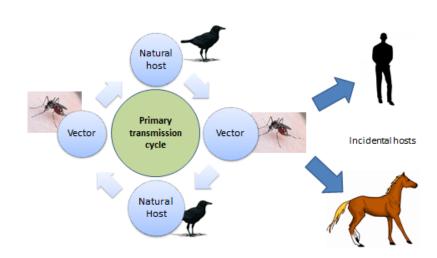
Prata et al. 2020; Sobral et al. 2020; Yao et al. 2020.



Impact of global warming

 Increase in global temperature in the past few decades & future projections (of ~ 3°C rise by 2050) have & will result in the transfer of vectors of human pathogens & reservoirs (as well) in unreported regions of the world!

• Spread of dengue virus in southern Europe is a classic example!



Caminade C, et al. (2019) Ann N Y Acad Sci 1436):157-173; Parkinson AJ, et al. (2014) Int J Circumpolar Health

Impact on Influenza epidemics

 Spread of Flu from one continent to another is linked with climate-driven changes.

 In case of H5N1 transmission, it was found that climatic factors (like temperature, relative humidity, & wind speed) changed the migration patterns in migratory birds which increased avian flu outbreaks in some regions in Egypt



Naguib MM, et al. (2019). Infect Ecol Epidemiol

Impact of Heat stress on viral genome

- Sequencing revealed that thermotolerance occurred via a **key mutation in gene** P5 (viral lysis protein), previously shown to be associated with heat shock survival in the virus.
- robust populations adapted faster to the environmental change, and more easily accessed mutations of large benefit.
-genetic robustness can play a role in determining the relative ability for microbes to adapt to changing environments
- N501Y substitution of SARS2 provides higher fitness against tropical high temperature

Extreme weather extremes: floods and droughts

- The occurrence of floods, due to unpredictable & heavy rains has been indirectly linked with high incidences of vector-borne disease.
- For instance, there were more outbreaks of **Dengue, WNV, & malaria** due to heavy rainfall and flooding in Romania (1996–97), Czech Republic (1997), Italy (1998), southern Africa (2000), and some parts of Southeast Asia (2011)

https://link.springer.com/article/10.1007/s42398-021-00165-x

Biodiversity loss

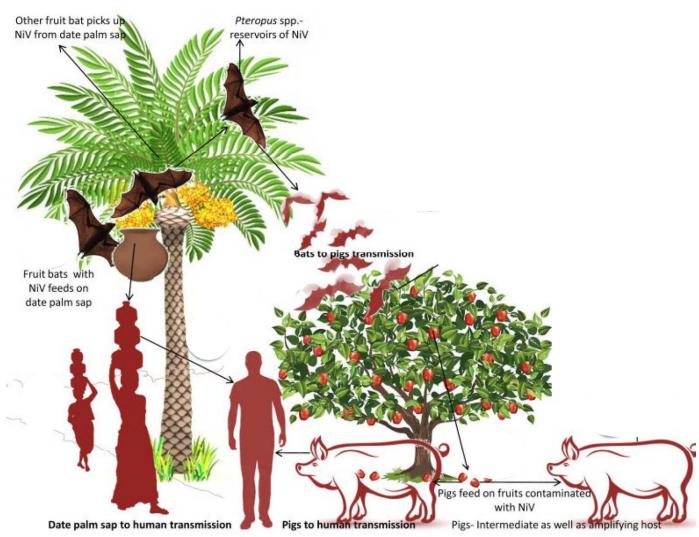
- Biodiversity loss makes an ecosystem more fragile and possible breach by the pathogens to humans becomes much easier.
- Biodiversity loss is correlated with a decrease in forest cover, increased human-animal conflicts, upsurge in livestock farming, and cultivation of wild animals.
- Sudden changes in species diversity can also result in an upsurge of novel pathogens.
 - Nipah virus outbreak in Malaysia in the late 1990s is an ideal example
 - Hantavirus outbreaks followed after a decline in the diversity of small mammals' reservoir (rodent) increased the prevalence of viruses

Suzán G, et al. (2009). PLoS ONE 4(5):e5461

Virus Nipah, Malaysie 1998

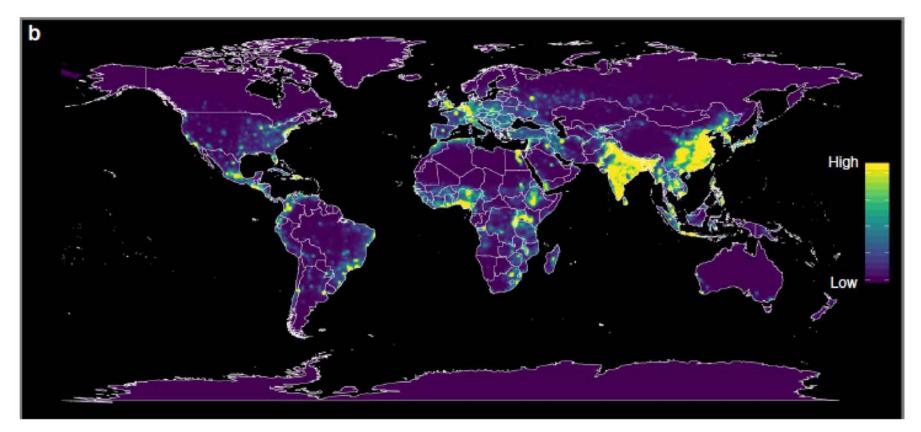
Deforestation for palm tree plantation \rightarrow





Singh et al 2019 VetQuater

Risk of emerging zoonosis in humans



- ☐ Tropical forest regions
- \square High species richness in mammals
- \square Links with changes in land use

Allen et al 2017 Nature communication

Anthropogenic factors

Intensive livestock production

Exhaustive agricultural practices

Urbanization

Globalization

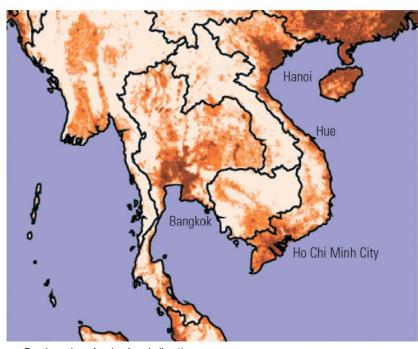
Pollution

Intensive poultry production in Southeast Asia

 Intensive poultry production in Southeast Asia led to the highly pathogenic Avian Influenza upsurge early in 2004

Slingenbergh J, et al. Ecological sources of zoonotic diseases. Rev Sci Tech 23:467–484

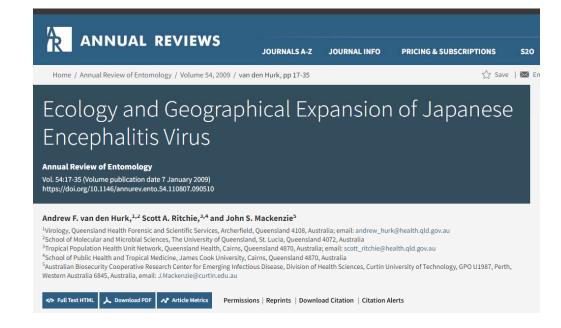
Estimated poultry density in Thailand, Laos and Vietnam (animals per km2)





Japanese encephalitis virus (JEV) outbreaks in Southeast Asia

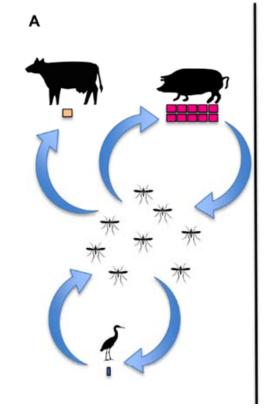
 In the case of JE virus outbreaks in Southeast Asia, increased production of irrigated rice and pig farming were reported as major anthropogenic factors to induce disease prevalence

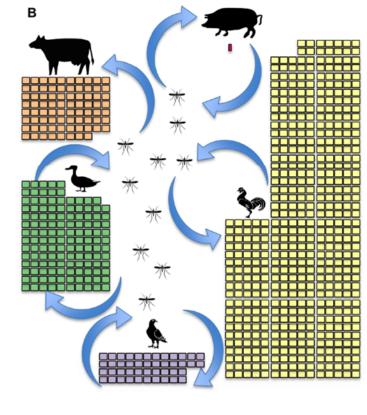


van den Hurk AF, et al. Annu Rev Entomol. 2000:54:17-35

JE Virus Transmission: Changing trends

Outbreaks of JE in the 1950s in Saitama Prefecture, Japan



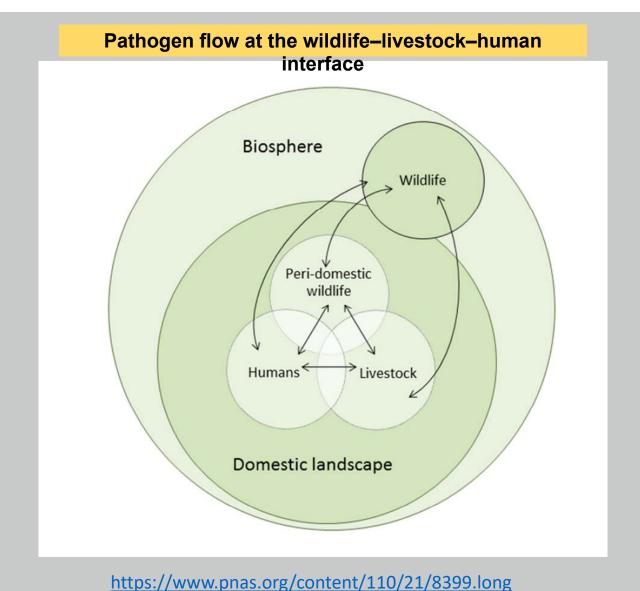


JE-endemic regions of India & B'Desh

Role of agricultural intensification &/or environmental changes

 "In the near future rate of zoonotic disease emergence or re-emergence will be closely linked to the evolution of the agriculture environment nexus"

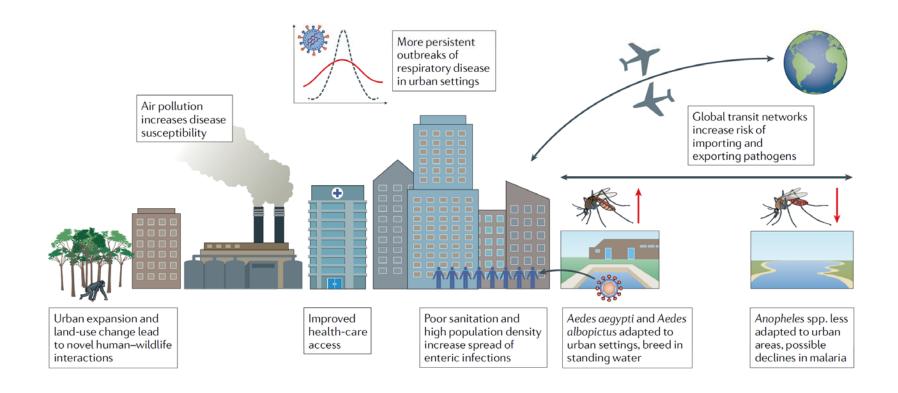
Proc Natl Acad Sci USA 110:8399–8404

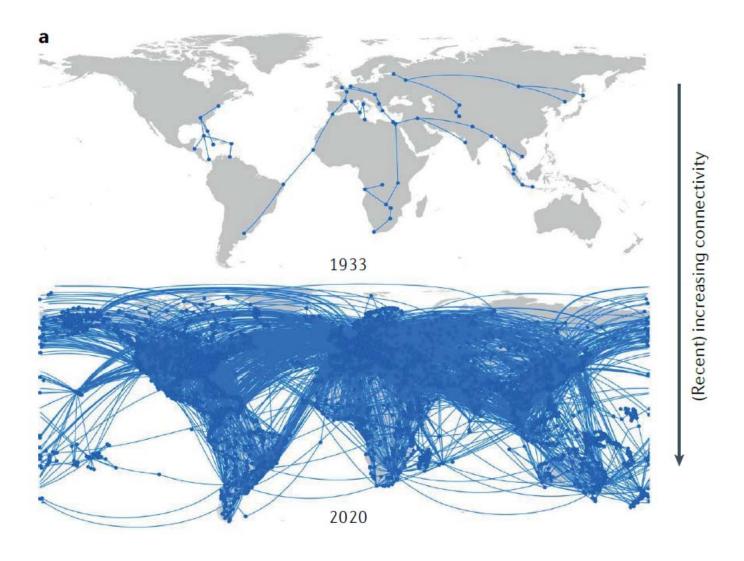


Extensive use of agrochemicals

- The extensive use of agrochemicals in agriculture has resulted in the loss of biodiversity, an increase of desertification & marginal lands, and mutation among microbial populations in soil and water bodies (Arora 2018).
- It has been realized that pollution due to dangerous recalcitrant and xenobiotic pollutants including **pesticides** can be significant for the evolution of bacterial & viral pathogens & their transfer to humans.

Impacts of urbanization on infectious disease





Impacts of urbanization on viral diseases

 Rapid urbanization affects peak timing, prevalence, & bimodality of influenza pandemics in Australia.

 Slums in the cities/peri-urban areas have become COVID hotspots in many countries and have become major areas for the spread of SARS2in big cities

SCIENCE ADVANCES | RESEARCH ARTICLE

SOCIAL SCIENCES

Urbanization affects peak timing, prevalence, and bimodality of influenza pandemics in Australia: Results of a census-calibrated model

Cameron Zachreson¹*, Kristopher M. Fair¹, Oliver M. Cliff¹, Nathan Harding¹, Mahendra Piraveenan¹, Mikhail Prokopenko^{1,2}

We examine salient trends of influenza pandemics in Australia, a rapidly urbanizing nation. To do so, we implement state-of-the-art influenza transmission and progression models within a large-scale stochastic computer simulation, generated using comprehensive Australian census datasets from 2006, 2011, and 2016. Our results offer a simulation-based investigation of a population's sensitivity to pandemics across multiple historical time points and highlight these notable trends in pandemic patterns over the years: increased peak prevalence, faster spreading rates, and decreasing spatiotemporal bimodality. We attribute these pandemic trends to increases in two key quantities indicative of urbanization: the population fraction residing in major cities and international air traffic. In addition, we identify features of the pandemic's geographic spread that we attribute to changes in the commuter mobility network. The generic nature of our model and the ubiquity of urbanization trends around the world make it likely for our results to be applicable in other rapidly urbanizion attorions.





Preventing COVID-19 Amid Public Health and Urban Planning Failures in Slums of Indian Cities

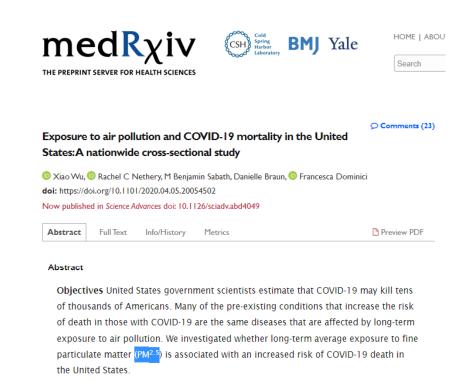
Amit Patel ①

The COVID-19 pandemic has brought renewed attention to the lack of urban planning and its public health implications in developing countries. Share momentations for the dual challenges of diventically poor residential environments and the acute effects of a paniencia and the preventive measure that follows. In this paper, I assess the effectiveness and implications of social distancing, frequent hand-auxiling, and wholeven in the context of plannin in falami cities, where occronology, lack of across to souter and sanitation, and dependence on daily sugges for sustenance and livelihoid are common. Using data from multiple sources, I demonstrate that an only will tiltoe measures be hard to articles in slams in the short term due to specific characteristics of these habitats, but they will bring near challenges in the lange term due to disproportionate impacts on the sharin port, casson formed from this pandemic will require us to retinize patients but the responses and urban planning practices that could better properse our clinks for future pandemics.

KEY WORDS: COVID-19, slums, India, urban health, urban plannin

Air pollution impact on Covid19

- Cities with high air pollution levels are more affected with COVID19 cases and deaths
- Indoor air pollution aggravates the severity of COVID-19.
- A slight increase of 1 µg/m3 in PM^{2.5} can result in 8% increment in COVID19 death rate!!



Wu X, et al. medRxiv 20202004200520054502. https://doi. org/ 10. 1101/ 20200 40520054502

Bushmeat (Wildlife) consumption and trade

Wet markets



Bush meat consumption



Kurpierset al 2016 *In* Problematic Wildlife, Springer

Bushmeat consumption & emergence of novel zoonotic disease

- Every four months one new infectious disease emerges in humans & is likely to be linked to the consumption of NHP and other exotic animals (UNEP frontiers 2016 report)
- HIV 1 & 2 reached in humans by multiple crossspecies transmission of simian immunodeficiency viruses (SIVs) through bushmeat hunting & trade
- **SARS-CoV** is reported to be originated from bats then transmitted to humans, while **MERS-CoV** passed on from bats to camels to humans

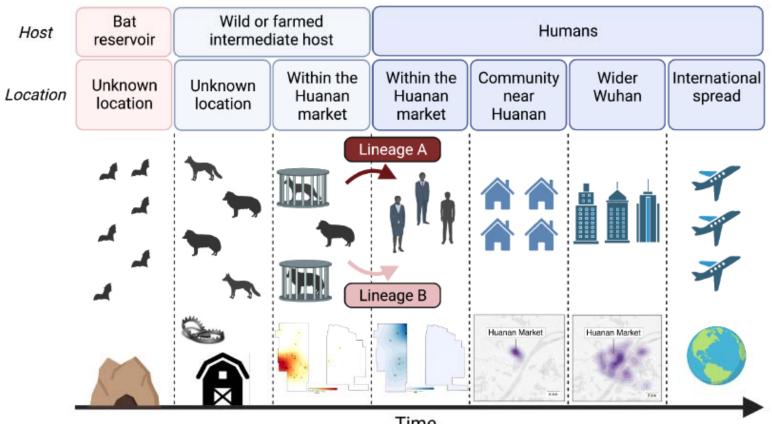


Abstract

Acquired immunodeficiency syndrome (AIDS) of humans is caused by two lentiviruses, human immunodeficiency viruses types 1 and 2 (HIV-1 and HIV-2). Here, we describe the origins and evolution of these viruses, and the circumstances that led to the AIDS pandemic. Both HIVs are the result of multiple cross-species transmissions of simian immunodeficiency viruses (SIVs) naturally infecting African primates. Most of these transfers resulted in viruses that spread in humans to only a limited extent. However, one transmission event, involving SIVcpz from chimpanzees in southeastern Cameroon, gave rise to HIV-1 group M—the principal cause of the AIDS pandemic. We discuss how host restriction factors have shaped the emergence of new SIV zoonoses by imposing adaptive hurdles to cross-species transmission and/or secondary spread. We also show that AIDS has likely afflicted chimpanzees long before the emergence of HIV. Tracing the genetic changes that occurred as SIVs crossed from monkeys to apes and from apes to humans provides a new framework to examine the requirements of

Sharp & Hahn BH. Cold Spring Harb Perspect Med. 2011(1):a000

Schematic diagram of key milestones in the emergence of SARS-CoV-2



Time

https://zenodo.org/record/6299600#.YhxLmDhBzIW

Conclusions

- 75% of emerging viral diseases in humans are from animal origin
- Human pressure on the environment intensifies exposure of wildlife to human
- Spread of diseases is accelerated through globalization/urbanization

What can we do?

- Early warning and action -> preparedness, cooperation
- Limit wildlife—human interface market/ trade/ bushmeat—> balance between prohibition, regulation and support
- Human disturbance of habitats -> reduce human footprint, relocalisation
- Need to conserve the ecosystem! Conservation of natural resources and control of environmental pollution
- Need biotechnological tools & study of the microbiome of possible reservoirs--identifying the possible novel
 pathogens or their genes in advance
- A paradigm shift is required in the thinking and working to let the nature breathe and sustain itself.

Thank You for your attention!