

How Scientists Look for New Pathogens Before They Spread

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Abstract

Emerging infectious illnesses are a persistent danger to global health. They often come from unexpected places, such as wildlife reservoirs, changes in the climate, or human activities. Finding new diseases before they create epidemics is one of the biggest problems in modern epidemiology. This article talks about how scientists use genetic monitoring, field sampling, and real-time data processing to find, track, and describe new infectious organisms in a way that works together. Researchers can now find viral and bacterial signatures in environmental and clinical samples long before symptoms show up in the general population. This is thanks to improvements in metagenomic sequencing, artificial intelligence, and global health networks. Case studies, ranging from the initial identification of SARS-CoV-2 variations to pathogen surveillance in isolated habitats, demonstrate how proactive monitoring can avert epidemics and inform public health strategies. In the end, early pathogen identification is not only a scientific goal, but it is also a key part of being ready for a pandemic in a world that is connected.

Keywords: Pathogen monitoring, new infectious illnesses, metagenomic sequencing, epidemiology, readiness for a pandemic

INTRODUCTION

Pathogens can spread swiftly across the planet. Scientists can warn people about a problem before it gets worse if they find it early. Finding pathogens is particularly important for controlling infectious diseases, supporting clinical microbiology, and protecting public health. Experts have improved pathogen discovery techniques, technologies, and strategies to quickly find new infections since COVID-19. Working together on a global scale has made networks around these detection attempts stronger. Genetic changes or the spread of diseases between species can cause new illnesses to appear, which can be a problem for health and the economy around the world. Monitoring environmental samples, such as animal specimens from humans, instruments, and labs, helps keep track of these changes, find any hazards, and give people time to prepare [1].

WHAT PATHOGENS ARE AND HOW EARLY DETECTION HELPS

Pathogens are microorganisms that go into living things and make them sick. Viruses, bacteria, fungus, and parasites are some of them. Most infections can't go after new hosts by themselves. They depend on their first host migrating to a different place. The pathogen could receive another chance to infect more hosts if the host moves to a different nation. Zoonoses are pathogens that can spread from animals to people. They can move from one type of animal to another. Rabies, the West Nile virus, and SARS-CoV-2 are all examples. Finding infections early, before they have a chance to spread, helps keep people, crops, and animals safe. The "One Health" idea is something that the World Health Organisation keeps an eye on. A novel infection can hurt all three when it spreads.

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Finding a crop disease early can help keep it from getting into a field or barn. It stops a new infection from getting into a music hall. It stops a dangerous fungus before a farmer can send birds to the store. Scientists seek for novel infections in three key places: labs, communities, and ecosystems. They also tell the public about their discoveries and let partners know about them.

Pathogens don't care about borders. They travel around the world. Tools and methods for keeping an eye on infections must also keep up. There's never enough time. If you don't stop an invader, they can get a foothold before anyone notices. Once it has settled in, getting rid of it is almost impossible. More than 80% of the germs that make people sick come from other animals. Globalisation, land use, and climate change all make the spread happen faster. The World Organisation for Animal Health uses the term "potential zoonotic pathogen" to mean a new arrival that people should keep an eye on. Researchers can suggest a reaction if they know the invader's history and biology [2].

WAYS THAT PEOPLE WATCH EACH OTHER TODAY

Pathogens are germs that make people, animals, and plants sick. Viruses, bacteria, and fungi are some of them. You can stop these germs from spreading to other people by killing or inhibiting them in people who are sick or in the environment. A lot of the time, finding and treating a virus early stops a few cases from turning into major outbreaks that are harder and more expensive to control. When you talk about a major outbreak, it doesn't indicate that one thing caused the rise. Open and honest reporting and exchange of data can help set up quick alert systems that spread the latest information broadly and make people more aware of certain circumstances that need particular attention.

There are more than one way to keep track of intriguing pathogens. You can look for certain types of reported test results and link them to certain viruses from certain places of the world. Some tests often identify a restricted number of pathogens in particular samples from specific contexts. If infections from a monitored pathogen or organisms not typically associated with such illnesses manifest at an area where the pathogen is not endemic to that organism, it suggests that the infection likely originates from a different source. These signals can be gathered and communicated. By monitoring the date and location of notifications regarding discovered Norrbottens or other outbreaks, it is feasible to ascertain whether these incidents occurred at a time and place associated with subsequent outbreaks of the same or a different pathogen [3].

People from health agencies, wildlife agencies, and other partners can collect samples from the environment. This could mean taking samples of soil, water, or weather and analysing them for pathogens that are of interest. You can also take samples from animals, including wild birds if there is avian flu. Along with the results, information about the samples taken, like where and when, is submitted [4].

Specific dangers in 1-12 life forms that can carry many diseases in their intestines may be sampled in certain situations according to a careful sampling schedule. Comparing field and laboratory work keeps pathogens that are not usually important in the spotlight. Wilder, farms, and other animals are watched for signs of pathogens that are known to infect people, those that cause unusual clustering of unknown illnesses or are spreading at the same time as injuries that are easy to see in ecosystems that are already in place in certain areas where similar pathogens have not yet stabilised, and those that are linked to unrelated human injury data. Detecting pathogens before, during, or after temporary human-introduced occupancy events provides valuable information for assessing potential future actions [5].

Testing in Labs That Are Careful

Lab-based vigilant testing looks for new germs that might spread from animals to people. Scientists put together information about animals and reports of diseases in people. They figure out the best places, times, and activities for animals to move from one person to another. When this happens, communities collect samples from high-risk species in the area to look for new infections.

Not all hazards can be found in lab tests. People interact with animals outside of areas with a lot of risk. Facilities that have been chosen test animals and their surroundings for germs that can cause disease. They look for biological signs in water and food sources, waste streams, and animal products that could mean new infections are on the way, like genetic material, toxins, and traces of vaccine candidates. They check samples for known risks. Networks communicate information about these activities, the species that were tested, and the presence and spread of pathogens to help with larger surveillance operations.

Field Sampling in Communities

Scientists gather information about individuals and their environments. They do this by working with different groups of individuals in communities and the people that live there. Community groups and those who are connected to them are gathering information. Researchers want to know about diseases and the germs that cause them. They also collect weather data around the samples they collect.

The health data collected from people includes ailments and health complaints like fever, rashes, vomiting, diarrhoea, cough, ear infections, joint discomfort, eye infections, and more. This information helps find infections that are making people sick as soon as feasible [6].

Keeping an Eye on Animals and Environments

Scientists need to keep an eye on animals and ecosystems to find infections before they spread. An outbreak usually happens after a spillover. Nonetheless, the development of a novel disease may transpire undetected in humans [7]. So, keeping an eye on wildlife, pets, and the environment can help find signs of a novel pathogen before it spreads to people. This proactive step is still very important for quick and effective control measures [8].

Several biotic and abiotic elements contribute to monitoring activities among various ecological indicators. Species composition and diversity offer insights into ecological equilibrium, wildlife population dynamics, habitat degradation, biodiversity decline, and climate change. Ecosystem stability is associated with higher latitudes and altitudes, resulting in unique ecosystems that predate the previously recognised diversity [9].

Sharing Data and Sending Alarms

Scientists recognise that speed is very important when novel germs threaten people's health. They can intervene before a new infection spread if they can find it quickly. They may tell people what to do, make vaccines, and get hospitals, farms, and other locations ready. So, surveillance is really important. Scientists and people who work in public health keep an eye out for such risks. They quickly communicate information with each other and with the public.

Sharing data is very important for finding pathogens. A lot of governments, labs, and businesses keep an eye on possible risks. When testing, analysis, or monitoring gives new information, it is added to public databases. Anyone in the globe can get these datasets for free. The World Health Organisation (WHO) [10] runs websites that list confirmed diseases. Official notifications talk about possible hazards, such animals dying in strange ways or people getting sick in ways that are linked to a recognised pathogen. Public health officials can swiftly investigate the situation and act because these signals get out so quickly, sometimes in less than 48 hours [11].

THINGS SCIENTISTS USE

To stop epidemics, it's important to keep people, crops, and animals from meeting new viruses. To find new infections, you need to watch high-risk areas, take samples, and study them [1]. They test samples of many different kinds. They could have the pathogen itself or things that it leaves behind when it infects its host. Scientists use these kinds of indications to help them figure out what to do next.

The goal is to find a new pathogen in time to do something about it that will protect people, crops, or animals. Viruses, bacteria, fungus, parasites, and prions are all pathogens of interest. A new virus that

can spread between people has the potential to spark an outbreak. Being able to find it promptly could provide you the time you need to stop or slow its spread. The same is true for various sorts of germs that can spread widely from animals or plants or only among certain people.

Sequencing of Genes

Genetic sequencing is a necessary tool for keeping an eye on germs. Genetic sequencing shows the DNA or RNA of a pathogen. It shows what kind of organism the pathogen is. It also reveals how closely related a newly found pathogen is to other infections that are already recognised. Scientists can figure out how harmful a pathogen is and where it can spread next by looking at its genetic code.

Most newly found infections can make people, plants, or animals sick. The first thing that scientists investigate are pathogens that impact people. It's crucial to find these pathogens early. When scientists find a pathogen, they can start to figure out where it has been and where it might go next [12].

Models for Computers

Pathogens can spread swiftly, affecting many people, crops, or animals before anybody notices. This might happen even when you're looking for fresh ones. Before they spread, scientists try to find new infections. They want to keep people, animals, and crops safe, so they focus on the ones that affect people, livestock, or valuable crops. They employ surveillance, sharing alerts, finding problems early, and responding quickly.

Computer models can help us figure out how a new infection might spread. These simulations help scientists get ready for what will happen. They indicate how fast the virus spreads, which groups of people are most at danger, how fast the pandemic is growing, and where it might spread in the future. Models employ information from outbreaks of other pathogens or species that are similar. Scientists are always making them better by learning new things [13, 14].

Quick Tests

Researchers have more time to plan since rapid testing quickly discover some infections.

It can take days to weeks to find a new pathogen. Scientists are in a hurry to figure out what the pathogen is. Rapid tests that seek for bits of genetic material help speed up the early work [2]. The quick tests give unambiguous results. It's easy to share them. Because all tests give the same information, numerous teams can test at the same time. Scientists compare results from different tests and locations. The extra time helps plans [15].

HOW RESULTS LEAD TO ACTION

Finding pathogens early helps stop their spread and find them quickly. Public health takes steps to protect people from infections such as sampling, making vaccines, developing therapies, and moving people to safer places [1]. Vaccines have problems include finding the right target, choosing the right platform, designing the right candidate, running trials, and scaling up production. There may be extra safety measures at farms, workplaces, and transportation hubs [3].

Responses to Public Health

Scientists move swiftly when they find a new pathogen. Early reactions lower the number of deaths and illnesses. They also stop food shortages and losses in the economy. It helps to have smart public health programs. It is possible to stop outbreaks. Vaccines and treatments are made [1]. Workplaces and farms are ready.

When discoveries are communicated, plans change. Health officials are notified. Being ready is the most important thing. Plans that are already in place get new action steps. The first response usually focusses on farming. The assessments of pathogen risk are done. People know about the risks at work. Other nations are notified [16, 17].

Creating Vaccines and Treatments

The purpose of early detection is to find a disease as soon as feasible and make a fast vaccine, antiviral treatment, or antibody treatment to halt it. It took 131 days from the time a new disease was found to the time a vaccine was first used to treat the H1N1 pandemic flu strain. It took only 15 days for the first test of monoclonal antibody therapy against the Ebola virus. These initiatives rely on early pathogen detection, viral isolate characterisation, pathogen library screening against circulating strains, and the expedited advancement of novel vaccination and monoclonal antibody technologies [1].

Getting Farms and Workplaces Ready

Research indicates that pathogens account for 60% of newly emerging infectious disease outbreaks that pose risks to public health, agriculture, and animals. Closing workplaces and farms lowers the risk of disease appearing and spreading. Specific places are picked depending on where infectious disease epidemics have happened before. It is also possible to map out the most likely ways that pathogens could get into the site. When possible, data from monitoring household animals and wildlife reservoirs is combined [4].

PROBLEMS AND LIMITS

Finding new viruses is a race against time. Finding outbreaks early is the most important thing to do. Pathogens are bad organisms that cause diseases as COVID-19, Ebola, and many infections in animals. It is easier to stop these germs from spreading to more people if they are found early. People's health is better, and the economy can grow when they are well [1]. Scientists use three main methods to find novel pathogens: sampling, sequencing, and sharing [3].

Infections that kill come from animals. The first several months of the COVID-19 pandemic show how quickly germs spread when they get into people. People still don't know all the harmful infections that can harm plants, animals, and people.

Data Gaps

There are holes in the collection and reporting of pathogens. No existing detection efforts stop new diseases from happening. These big data gaps make it harder to understand and act.

There are holes in the collection and reporting of pathogens. Pathogen detection happens when there are gaps in technology, monitoring, and awareness [14]. Sampling ceases or diminishes with limited reports. Fewer reporting led to fewer infections being found and less awareness of connected topics or events. No current efforts to find diseases stop new ones from happening. New viruses come out all the time, but there is only one pathogen that has a plan that is ready to go: the H17N10 virus. Agencies keep track of daily reports, but current outbreak reports don't focus on early warnings. These big holes in the data make it harder to interpret and act.

Privacy and Morals

Even while epidemic prediction systems have been shown to work, they can raise several ethical and privacy issues [18]. For instance, the mere possibility of being recognised can pose privacy threats in the context of detecting a developing virus. People who respond to early alarms (even if they aren't unwell) may also face scrutiny or discrimination after that. After a suspected outbreak, these problems often cause certain demographic groups or entire geographic areas to be unfairly stigmatised. Scientists talk about and try to figure out how to deal with the moral problems that come up when an epidemic case or virus of concern gets more attention.

Working Together Around the World

Surveillance is important all around the world. Pathogens move swiftly. The way infections are spread in one location can often give you a heads-up about epidemics in other areas. But keeping an eye on things costs time and money. Countries value financial aid, technical support, and access to new ways to find things. The United States gives money to assist upgrade existing programs. Scientists set

up international seminars to share tools for reporting data. The World Organisation for Animal Health oversees setting higher standards for sharing information between animal health bodies [11].

Sharing pathogens makes things clearer but also makes people more worried. During the 2013 H7N9 influenza outbreak in China, specialists from around the world requested samples to keep an eye on how it was spreading. China agreed and set up a public website to keep track of new instances. Countries then told the World Health Organisation about H7N9 activity but didn't give any information. The extra viral knowledge sped up the assessments of danger. During the Ebola outbreak in 2015, countries reported tested bodies and empty buildings, which sped up the process of finding dangerous remains and contact-tracing sites. The Global Health Security Agenda was used by the United States and non-governmental organisations. The desire to prevent epidemics unites different political conditions to share data on infectious diseases [19].

WHAT PEOPLE CAN DO

Early discovery keeps diseases from spreading, which is good for people, animals, and plants. A pathogen is a virus, parasite, or organism that makes you sick. You can get pathogens from the air, water, soil, and animals. Early detection finds an outbreak before it spreads too far. For instance, finding a cow virus, like foot-and-mouth disease, early can stop an outbreak that could stop trade between countries. Finding the bird flu virus or another disease early lets people take steps to keep people from getting sick before the sickness spreads from animals to people [16].

Scientists use community-based testing and monitoring to keep an eye on new pathogens as they move from one group to another. This helps them find and stop new pathogens before they can spread widely. Early detection activities in the community are especially crucial for viruses because they might modify their genes a lot as they migrate from one host to another [3].

To keep people safe, laboratory tests look for infections in samples. Scientists get samples from different places and check them for germs. Some of the samples are air filters, wastewater, and crops. A lab with biosafety level-3 (BSL-3) equipment checks for new viruses in samples that come from outside the lab. Scientists utilise genomic sequencing to figure out what a new virus is like. The results show what steps to do next [4].

Community sampling collects information about the health of people and animals in communities. Community health institutions collect data on probable outbreaks, including fever. Organisations that study science look at reports from local farms about crop cases. Community-based biosecurity programs keep an eye on ill fish to find out which diseases need more testing. People also tell if their cattle get sick after eating certain kinds of animal feed.

Animals can show us new viruses that can hurt people or crops. Changes in ecosystems, the temperature, and the way people farm can all influence how animals behave and where they live. Scientists also watch mosquitoes, flies, and ticks because they could be carriers of viruses. Watching these signals can show that there is a higher chance of infections spreading to people.

Researchers discuss information on new discoveries, new germs, and possible outbreaks. They use an internet mechanism to disseminate information quickly. Quick notice helps public health officials take measures to stop large epidemics.

CONCLUSION

Finding infections early is very important for stopping epidemics. A pathogen is a living thing that can make you sick. An infectious agent is also called a pathogen. This word derives from the Latin word meaning "to infect." Pathogens are things like viruses, bacteria, fungi, and parasites. A lot of infections can kill people, animals, and plants. Finding viruses early lowers the risk of their spreading, which helps save lives and crops.

Before they spread, scientists look for new infections. It takes a lot of steps to find novel infections early. First, scientists gather data about pathogens. Next, they search for new infections in animals, plants, and the environment. Finding a novel pathogen usually means taking more specific steps. Researchers put monitoring first in places with a lot of connections, such as cities and airports. They also tell each other about fresh discoveries as they happen. News about probable threats spreads swiftly around the world.

REFERENCES

1. Guo C, Wu JY. Pathogen discovery in the post-COVID era. 2024. Available from: <https://www.ncbi.nlm.nih.gov>.
2. Mastin AJ, van den Bosch F, van den Berg F, Parnell SR. Measuring the hidden costs of inaccurate detection for early detection surveillance. *Philos Trans R Soc B Biol Sci*. 2019 Jul 8;374(1776):20180261. doi: 10.1098/rstb.2018.0261. PMID:31104597; PMCID:PMC6558562.
3. Koopmans M. Surveillance strategy for early detection of unusual infectious disease events. 2013. Available from: <https://www.ncbi.nlm.nih.gov>.
4. Chiu KH, Sridhar S, Yuen KY. Getting ready for the next pandemic: Problems with strengthening surveillance systems. 2023. Available from: <https://www.ncbi.nlm.nih.gov>.
5. Downie DL, Rao P, David-Ferdon C, Courtney S, et al. Surveillance for emerging and reemerging pathogens using pathogen-agnostic metagenomic sequencing in the United States: A critical role for federal government agencies. 2024. Available from: <https://www.ncbi.nlm.nih.gov>.
6. Ferguson JM, Langebrake JB, Cannataro VL, Garcia AJ, et al. Optimal sampling strategies for detecting zoonotic disease epidemics. 2014. Available from: <https://www.ncbi.nlm.nih.gov>.
7. Guberti V, Stancampiano L, Ferrari N. Surveillance, monitoring, and assessment of wildlife diseases: A public health and conservation perspective. *Hystrix Ital J Mammal*. 2014;25(1). doi: 10.4404/hystrix-25.1-10114.
8. Vourc'h G, Bridges VE, Gibbens J, De Groot BD, et al. Detecting emerging diseases in farm animals through clinical observations. 2006. Available from: <https://www.ncbi.nlm.nih.gov>.
9. Childs JE. Pre-spillover prevention of emerging zoonotic diseases: Targets and available tools. 2007. Available from: <https://www.ncbi.nlm.nih.gov>.
10. Elbe S, Buckland-Merrett G. Data, disease, and diplomacy: GISAID's innovative contribution to global health. 2017. Available from: <https://www.ncbi.nlm.nih.gov>.
11. Morse SS. Global surveillance of infectious diseases and health intelligence. *Health Aff (Millwood)*. 2007 Jul–Aug;26(4):1069–1077. doi: 10.1377/hlthaff.26.4.1069. PMID:17630449.
12. Grubaugh ND, Ladner JT, Lemey P, Pybus OG, et al. Tracking virus outbreaks in the twenty-first century. 2018. Available from: <https://www.ncbi.nlm.nih.gov>.
13. Siettos CI, Russo L. Mathematical modelling of infectious disease dynamics. 2013. Available from: <https://www.ncbi.nlm.nih.gov>.
14. Kraemer MUG, Cummings DAT, Funk S, Reiner RC, Faria NR, Pybus OG, et al. Reconstruction and prediction of viral disease epidemics. *Epidemiol Infect*. 2018 Nov 5;147:e34. doi: 10.1017/S0950268818002881. PMID:30394230; PMCID:PMC6398585.
15. Marx V. The coronavirus pushes laboratories to accelerate research efforts. 2020. Available from: <https://www.ncbi.nlm.nih.gov>.
16. Petrone ME, Holmes EC, Harvey E. Through an ecological lens: An ecosystem-based approach to zoonotic risk assessment. 2023. Available from: <https://www.ncbi.nlm.nih.gov>.
17. Cassetti MC, Pierson TC, Patterson LJ, Bok K, et al. Prototype pathogen approach for vaccine and monoclonal antibody development: A critical component of the NIAID plan for pandemic preparedness. 2022. Available from: <https://www.ncbi.nlm.nih.gov>.
18. Keuleyan E. Liberty to decide on dual-use biomedical research: An acknowledged necessity. 2010. Available from: <https://www.ncbi.nlm.nih.gov>.
19. Switzer S, Morgera E, Tsioumani E, Burci GL. Biodiversity, pathogen sharing, and international law. 2019. Available from: <https://www.academia.edu>.