Human activities and zoonotic epidemics: A two-way relationship. The case of the COVID-19 pandemic

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Non-technical summary

We humans have the tendency to damage the natural environment in many ways. Deforestation and conversion of forests for residential, industrial development, and expansion of agricultural crops, as well as the burning of fossil fuels, are some activities that disrupt natural ecosystems and wildlife and contribute to climate change. As a result, the life cycles of pathogens and intermediate hosts (insects, rodents, mammals) as well as biodiversity are affected.

Through these activities, humans meet wild animals that transmit pathogens, resulting in their infection by zoonoses and causing epidemics-pandemics, the effects of which have as their final recipient himself and his activities.

Technical summary

This article aims to highlight the two-way relationship between those human activities and the occurrence of epidemics-pandemics. We will try to elaborate this two-way relationship, through the overview of the current pandemic (origin of SARS-CoV-2, modes of transmission, clinical picture of the disease of COVID-19, influence of weather and air pollution on prevalence and mortality, pandemic effects, and treatments). They are used as primary sources, scientific articles, literature, websites, and databases (Supplementary Appendix) to analyze factors involved in the occurrence and transmission of zoonotic diseases in humans (Ebola, influenza, Lyme disease, dengue fever, cholera, AIDS/HIV, SARS-CoV, MERS-CoV). The present paper concluded that humanity today faces two major challenges: controlling the COVID-19 pandemic and minimizing the risk of a new global health crisis occurring in the future. The first can be achieved through equitable access to vaccines and treatments for all people. The second needs the global community to make a great change and start protecting the natural environment and its ecosystems through the adoption of prevention policies.

Summary of social media

This article will try to highlight the two-way relationship of human activities with the emergence of epidemics-pandemics, through a multi-level approach to the COVID-19 pandemic, the challenges that humanity faces, and the need to adopt policies at a global level to prevent future pandemics.

Keywords: human activities; natural environment; two-way relationship; zoonoses; pandemic COVID-19

1. Introduction

Emerging infectious diseases are a threat to world health, according to the World Health Organization (WHO). Over the years, infectious diseases have plagued humanity (Ebola, plague, Asian cholera, yellow fever, Spanish flu). However, the contribution of medical science is enormous, given that severe and infectious diseases like smallpox and polio have been eliminated or reduced, while at the same time there are positive developments for diseases such as malaria, tuberculosis, and HIV/AIDS (Gresham et al., 2013; Hays, 2005). These achievements should not reassure the international community though, as new diseases emerge and deadly diseases of the past are re-emerging, causing epidemics and pandemics. But how do human activities relate to the emergence and spread of communicable zoonotic diseases? The present paper aims to highlight this existing relationship.

Humans with their activities invade the wild and disrupt the natural ecosystems, thus fueling the emergence of zoonotic diseases, as they come into direct contact with wild animals infected with viruses. These diseases rapidly spread to the human population through travel, trade, urbanization, migration, and human behavior, causing epidemics or pandemics that in turn result in human losses and huge social and economic implications. In the present study, the bidirectional relationship between human activities and epidemics/pandemics is highlighted through the multi-layered review of the COVID-19 pandemic (origin of SARS-CoV-2, clinical picture of COVID-19, modes of transmission, relationship of weather conditions and air pollution in the spread of disease, effects, and therapeutic developments).

2. Zoonoses: Definitions, classification, and typology of zoonotic dynamics.

2.1. Zoonosis

Disease or infection transmitted to humans by vertebrate animals (mammals, birds, reptiles, amphibians, and fish) that carry germs such as viruses, bacteria, fungi, and parasites (CDC, 2021).
2.2. Transmission of zoonoses

Transmission occurs through various ways (direct contact, indirect contact, carrier, consumption of contaminated food and water) and in multiple environments (urban centers, rural areas, trips, zoos, and open-air markets for the sale and consumption of wild animals) (Kock & Caceres-Escobar, 2022; Wegner et al., 2022). Table 1.

**Table 1. Means of transmission of zoonoses**

<table>
<thead>
<tr>
<th><strong>Direct contact:</strong> Bile, or contact with body fluids of an infected animal (saliva, blood, urine, feces)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Indirect contact:</strong> In places where animals live and roam, or there are objects contaminated with germs (aquarium water, coops, barns, plants, soil, food, pet food utensils)</td>
</tr>
<tr>
<td><strong>Carrier:</strong> Bitten by ticks or insects such as mosquitoes, fleas, lice, flies</td>
</tr>
<tr>
<td><strong>Consumption of contaminated food:</strong> Unpasteurized milk, meat and eggs not cooked well, raw fruits and vegetables contaminated with feces of infected animals</td>
</tr>
<tr>
<td><strong>Consumption or contact of contaminated water:</strong> Contaminated water with feces of contaminated animals</td>
</tr>
</tbody>
</table>

A dominant role in the transmission of zoonoses is played by "natural" hosts that act as reservoirs of pathogens, such as various species of bats, rodents, and birds that are natural reservoirs of viruses (chanterelles, arenaviruses, arboviruses, and coronaviruses). A key link in the transmission chain is the "intermediate" hosts (wild, domestic animals, and arthropods) through which pathogens can evolve and pass from the "natural" host to humans. Zoonotic diseases are also transmitted to animals and humans through the bite of arthropod vectors (mosquitoes, ticks, fleas) (Wegner et al., 2022). (Table 2 Table 3, Figure 1).

**Table 2. Transmission of zoonoses through direct, and indirect contact with wild or domestic animals and consumption of contaminated water or food**
<table>
<thead>
<tr>
<th>Disease</th>
<th>Pathogen</th>
<th>Natural Host</th>
<th>Intermediate Host</th>
<th>Transmission</th>
</tr>
</thead>
<tbody>
<tr>
<td>Brucellosis</td>
<td>Bacterium Brucella</td>
<td>Cattle, pigs, goats, sheep, dogs, etc.</td>
<td></td>
<td>Direct contact, consumption of contaminated animal products.</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>Inhalation of airborne contaminants.</td>
</tr>
<tr>
<td>Ebola</td>
<td>Virus</td>
<td>Bat (Pteropus)</td>
<td>Ape, Monkey Antelope</td>
<td>Direct contact with the body fluids of an infected animal</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>(duiker)</td>
<td></td>
</tr>
<tr>
<td>Leptospirosis</td>
<td>Leptospira</td>
<td>Rat</td>
<td></td>
<td>Direct contact with an infected animal</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>Indirect contact through soil or water contaminated with the urine of an infected animal</td>
</tr>
<tr>
<td>HIV/AIDS</td>
<td>Monkey immunodeficiency</td>
<td>Chimpanzean</td>
<td></td>
<td>Direct contact with the blood of an infected animal</td>
</tr>
<tr>
<td></td>
<td>virus (SIV)</td>
<td></td>
<td></td>
<td>Sexual activity</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>Intravenous drug use</td>
</tr>
<tr>
<td>MERS-COV</td>
<td>Coronavirus</td>
<td>Bat (Rhinolophus)</td>
<td>Camel (dromedary)</td>
<td>Direct contact</td>
</tr>
<tr>
<td>SARS-COV</td>
<td>Coronavirus</td>
<td>Bat (Rhinolophus)</td>
<td>Civet (poguma larvata)</td>
<td>Direct contact</td>
</tr>
<tr>
<td>SARS-COV-2</td>
<td>Coronavirus</td>
<td>Bat (Rhinolophus)</td>
<td>Mammals (e.g., pangolins, raccoon dogs, red foxes, minks, snakes) and domestic animals (e.g., cats)</td>
<td>Direct contact</td>
</tr>
<tr>
<td>Cholera</td>
<td>Bacterium vibrio cholaera</td>
<td></td>
<td></td>
<td>Consumption of contaminated water or food</td>
</tr>
<tr>
<td>Salmonellosis</td>
<td>Bacterium Salmonella</td>
<td>Cattle, chicken, turkey, pig.</td>
<td></td>
<td>Consumption of contaminated water or food</td>
</tr>
<tr>
<td>Escherichia coli</td>
<td>Bacterium Escherichia coli</td>
<td>Cows, deer, calves, goats, sheep</td>
<td></td>
<td>Consumption of contaminated water or food</td>
</tr>
</tbody>
</table>

(1) potential intermediate hosts

**Table 3. Transmission of zoonoses through vector**

<table>
<thead>
<tr>
<th>Disease</th>
<th>Pathogen</th>
<th>Host</th>
<th>Carrier</th>
</tr>
</thead>
<tbody>
<tr>
<td>Plague</td>
<td>Bacterium (Versinia pestis)</td>
<td>Rodents (e.g., black rats)</td>
<td>Flea</td>
</tr>
<tr>
<td>Chagas Disease</td>
<td>Parasite (Trypanosoma cruzi)</td>
<td>Humans and animals</td>
<td>Triatome bug</td>
</tr>
<tr>
<td>Malaria</td>
<td>Parasite (Plasmodium)</td>
<td></td>
<td>Mosquito Anopheles</td>
</tr>
<tr>
<td>Dengue Fever</td>
<td>Virus</td>
<td></td>
<td>Mosquito Aedes</td>
</tr>
<tr>
<td>Yellow Fever</td>
<td>Virus</td>
<td>Humans and animals</td>
<td>Mosquito Aedes</td>
</tr>
<tr>
<td>Lyme Disease</td>
<td>Bacterium (Borrelia burgdorferi)</td>
<td>Wild and domestic animals,</td>
<td>Tick ixodes ricinus</td>
</tr>
<tr>
<td>(Borreliosis)</td>
<td></td>
<td>birds, and reptiles</td>
<td></td>
</tr>
<tr>
<td>Leishmaniasis</td>
<td>Parasite (Leishmania)</td>
<td>Mammals (e.g., marsupials, rodents, edentates, monkeys, canids, and domestic dogs)</td>
<td>Sandfly Phlebotomus papatasi</td>
</tr>
</tbody>
</table>
2.3. Classification of zoonoses

Zoonoses are classified based on the causative agents (pathogens) responsible for their occurrence, the transmission cycle (orthozoonoses, cyclozoonoses, metazoonoses, saprozonoses), and reservoir hosts (anthropozaoons, zoonanthropozaoons, amphixenoses, human diseases) (Kock & Caceres-Escobar, 2022; Leal Filho, Ternova, Parasnis, Kovaleva, & Nagy, 2022). Table 4

Table 4. Classification of zoonoses
2.4. Dynamics of zoonosis typology

The course of a zoonosis depends on the dynamics of the pathogen. Wolfe et al. (2007) proposed a five-stage classification by which a pathogen that initially infects only animals (Stage I) goes on to develop into a pathogen that infects only human populations (Stage V).

In intermediate stages (II-IV) pathogens based on their behavior are divided into (a) pathogens transmitted from animals to humans causing "primary" infections, but do not show secondary transmission from human to humans (Stage II), such as West Nile virus or Brucella abortus, (b) animal-borne pathogens that spread to human populations, causing limited transmission cycles until extinction (Stage III), such as Ebola, Marburg and monkeypox viruses and (c) pathogens retained in animal tanks and can cause self-sustaining transmission chains in human populations (Stage IV), such as Yersinia pestis (plague) and pandemic influenza.

Over the years pathogens shaped the ability to change behavior and while in the past some of them may have caused limited cycles of transmission in human populations until they disappeared (Stage III), they have now re-emerged causing major epidemic outbreaks. An example is the Ebola virus that seems to have evolved into a pathogen that mainly infects human populations (Stage V). Ebola has caused many epidemic outbreaks since 1976 when it first appeared, with the largest one in West Africa (2014-2016) and recently with the outbreak in Uganda affecting the Country for the first time since 2012 (WHO, 2022c) Figure 2.

<table>
<thead>
<tr>
<th>Etiological factors</th>
<th>Transmission cycle</th>
<th>Tank hosts</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bacterial, mycotic, viral, rickettsial, chlamydial, parasitic, protozoal</td>
<td>Orthozoanoses: They are transmitted from one infected vertebrate host to another through direct contact, contact with a fomite or mechanical transmission. (eg rabies, brucellosis).</td>
<td>Human zoonoses: Diseases transmitted to humans by vertebrate animals (animal reservoir) through direct contact or indirectly through food, fomite or carrier (arthropods) (eg Lyme disease, Malaria, MERS-COV, SARS-COV, SARS-COV-2, Chagas disease)</td>
</tr>
<tr>
<td></td>
<td>Cyclozoanoses: The presence of more than one vertebrate animal is required for the development of the zoonosis. The animal acts as an intermediate host for the transmission of the disease to humans (eg echinococcosis)</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Metazoanoses: Transmitted by a host invertebrate where the pathogen multiplies, evolves and is transmitted to a vertebrate host (eg plague, Lyme disease)</td>
<td>Zooanthropoanoses: Diseases affecting humans, which act as a reservoir / host and are transmitted from animals to animals (eg SARS-CoV-2, Influenza A, yellow fever)</td>
</tr>
<tr>
<td></td>
<td>Saprozoanoses: Diseases that require the presence of a vertebrate host and another type of environmental reservoir, such as food, soil, plants (eg listeriosis, histoplasmosis).</td>
<td>Amphixeanoses: Diseases that are transmitted either from animals to humans or vice versa (eg Salmonellosis, Streptococcus)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Human diseases: Diseases that are naturally conserved and transmitted from person to person, with or without vectors (eg Malaria, SARS-COV, MERS-COV, SARS-CoV-2)</td>
</tr>
</tbody>
</table>
Similar behavior is shown by other viruses, such as that of monkeypox which was discovered in 1970 and has caused many epidemic episodes in the past. The virus strongly re-emerged in 2022, as it spread to many countries (mainly in Europe and America), outside the endemic areas of Africa, recording from the beginning of the year to October (26-10-2022) more than 76,500 cases and 36 deaths in 109 countries worldwide (WHO 2022d).

3. Human Activities and Epidemics of Zoonoses: A Two-Way Relationship

The occurrence and transmission of zoonoses in the human population is related to the existence of several factors, of which human activities seem to play a decisive role. Deforestation for the purpose of residential and industrial activity and the development of agricultural crops, the trade and consumption of wild animals are some activities with which man directly interferes with the natural environment, disrupting biodiversity and natural ecosystems.

A similar disturbance is caused by climate change (a result of human activities) and extreme weather events (heatwaves, floods, etc.), affecting the life cycles of pathogens and their vectors (insects, rodents, mammals), as well as the geographical their allocation. Human either directly or indirectly with his activities disrupts the balance of wildlife and fuels the emergence of zoonotic diseases (Loh et al., 2015; UNEP, 2020) Figure 3.
These activities lead many animals that are potential carriers of pathogens to explore new natural habitats, thus forcing them closer to inhabited areas. Such is the case with some species of bats that manage to continue their life cycle and grow in new acceptable habitats such as humanized environments (Afelt et al., 2018; Plowright et al., 2015; Reuter, Wills, Lee, Cordes, & Sewall, 2016). This increases the risk of infectious diseases as humans and pets come closer to wild animals carrying pathogens (Halliday & Rohr, 2019; Karesh et al., 2012; Keesing et al., 2010).

In addition to human activities, microbial resistance to antibiotics, inadequate health systems, the lack of specialized health personnel, the poor quality of public health, the low level of health services provided, and the absence of preventive measures and infectious disease surveillance systems contribute to the increase in communicable diseases.

These emerging diseases due to travel, trade, urbanization, population migration, and human behavior rapidly spread across the planet, causing epidemics and pandemics (CDC, 1994), which sequentially affect people and their activities with great impact on public health, health systems, the economy, tourism, education, and society.

The relationship between human activities and epidemics/pandemics could be characterized as a two-way relationship. A chain process whose beginning and the end are human and his activities. 

**Figure 3. The impact of direct and indirect human activities on biodiversity and natural ecosystems**

Figure 4. [https://doi.org/10.1017/sus.2022.18](https://doi.org/10.1017/sus.2022.18) Published online by Cambridge University Press
4. Categorization of factors favoring the emergence and transmission of zoonotic diseases in humans.

These factors could be distinguished into those that favor the emergence of zoonotic diseases (direct and indirect human activities, antimicrobial resistance to antibiotics) and those that contribute to their transmission to the human population (travel, trade, urbanization, human behavior). Figure 5, Figure 6
4.1. Factors that favor the appearance of zoonotic diseases

4.1.1. Human activities (direct)

A series of activities, such as deforestation, reforestation, changing land use for agriculture, urban and industrial development, exploitation-trade, and consumption of wild animals, directly affect the natural environment by contributing to the degradation of biodiversity and by disturbing the balance of wildlife.

Extensive deforestation in the Amazon, Southeast Asia, Central and West Africa has been linked to the spread of animal-borne diseases. Human intervention within the rainforests of Central and West Africa is reportedly linked to Ebola outbreaks. Oliveiro et.al (2017) observed such outbreak of the disease in populated areas created by extensive deforestation between 2001-2014.

Reforestation, another form of ecosystem disruption, is responsible for the emergence of borreliosis in the United States and Europe. This reverse process has led to an increase in deer population. The Ixodes Ricinus sea bream, which is responsible for transmitting the borreliosis disease, exists in deer’s skin. Human activity in these areas has resulted in the direct contact of a large part of the population with the disease carrier (Fineberg & Wilson, 2010; Spielman, 1994).

The re-emergence of leishmaniasis in parts of Brazil is also related in the change of the use of the forest areas, where the natural trees are replaced by others suitable for the paper industry. This change in the natural ecosystem has favored the growth of foxes, through which the disease was transmitted to the human population (Patz, Graczyk, Geller, & Vittor, 2000).

Furthermore, the exploitation marketing and consumption of wild animal meat is associated with the occurrence of zoonotic diseases in humans. In China, for example, research has shown that both SARS-CoV (2002-2003) and SARS-COV-2 coronaviruses; related to bat viruses, may have been transmitted to humans through wild animals (vectors: Paguma lavrata for SARS-CoV and propably pangolin, raccoon dog, red fox, mink, snake etc for SARS-CoV-2) sold for consumption in popular markets in China (Mackenzie & Smith, 2020; Song et al., 2005; Wong, Javornik Cregeen, Ajami, & Petrosino, 2020).

4.1.2. Human activities (indirect)

Human activities (burning fossil fuels, cutting down trees, forest fires) are a key factor in climate change, due to the increase in greenhouse gas emissions (CO2 N2O, CH4, and chlorofluorocarbons) into the atmosphere. The result of climate change is the rise of global warming and extreme weather events (floods, heat waves, droughts).

Climate change contributes to the reduction of biodiversity, affects ecosystems, the life cycles of pathogens and vectors (insects, rodents, mammals), their geographical distribution and enhances the occurrence of diseases.

On one hand, temperature rise has been found to extend the geographical distribution of pathogen habitats (mosquitoes, ticks), causing shifts in infectious diseases such as malaria, dengue fever, yellow fever, and Lyme disease, from areas of lower latitude to higher altitudes. On the other hand, heavy rains and floods favor malaria and leptospirosis as environmental conditions (humidity, stagnant water) influence the female mosquito’s reproduction, which is the primary vector of these diseases (Harvell et al., 2002).

In parts of Asia and South America, periodic floods due to El Nino (ENSO) have been associated with malaria epidemics (Kovats, Bouma, Hajat, Worrall, & Haines, 2003) like in the coastal region of northern Peru (Bayer et al., 2014) and in Brazil, where the malaria epidemic of 2000 may be due to high temperatures and heavy rainfall (Vasconcelos et al., 2001). Likewise, Karande et.al (2003) believe that leptospirosis occurred in Mumbai, India after the 2000 floods.

Sunshine and warm temperatures combined favor the multiplication of pathogens as is the case with the bacterium Vibrio cholerae, the causative agent responsible for cholera (Islam et al., 2009).

Heatwaves, as a result of global warming, combined with prolonged periods of drought due to the El Nino phenomenon, make forests extremely flammable and vulnerable to catastrophic fires, which accelerate deforestation (Lenton et al., 2008; Nobre et al., 2016). Anthropogenic deforestation processes and fires work collaboratively to reduce biodiversity and disturbance of natural ecosystems.

4.1.3. Antimicrobial resistance to antibiotics and public health systems
Antimicrobial resistance to antibiotics combined with poor hygiene conditions in health facilities, contribute to the appearance of infections. O’Neil (2014) has reached the conclusion that by year 2050, approximately 10,000,000 people will die from antimicrobial resistance, while the cost to health systems will exceed one hundred trillion dollars.

Similar estimates for public health and health systems are given in a recent report by 13 international organizations, which said the economic damage would be as catastrophic as the global financial crisis of 2008-2009, and by 2030, antimicrobial resistance could force up to 24 million people in extreme poverty (IAGG, 2019).

In addition, the risk of developing infectious diseases and causing epidemics increases especially in poor countries with limited health resources, such as sub-Saharan African countries. For example, the 2014 Ebola outbreak in West Africa quickly escalated into an epidemic due to inadequate public health infrastructure and logistical problems, lack of effective disease surveillance systems, hospital infection control and contact tracing, isolation, and patient care (Buseh et al. 2015).

4.2. Factors contributing to the spread of zoonotic diseases

4.2.1. International travels and trade

International travels and trade have in the past contributed to the spread of infectious diseases in humans. Yellow fever was transmitted from Africa to America in the 16th-17th centuries during the transportation of black slaves (Hays, 2005). The Asian cholera started in 1817 from the Ganges (Calcutta) plain and spread with merchant ships and British Navy ships, in the Middle East, Africa, and Europe. Additionally, the Spanish flu of 1918-1919, which appeared towards the end of World War I was transmitted worldwide by warships carrying troops (Vinet Freddy, 2020). In the recent past, the SARS-CoV epidemic (2002-2003) appeared to have spread to 31 countries through travel and trade; factors that are also responsible for the spread of SARS-CoV-2 coronavirus from China to all countries worldwide, causing the COVID-19 pandemic.

4.2.2. Intense urbanization

Intense urbanization, poor sanitation, and shortages of clean drinking water contribute to the growth of pathogens and the spread of infectious diseases. Dengue fever occurs in cities with tropical and subtropical climate, where the Aedes Aegypti mosquito, a carrier of the virus, accumulates in piles of waste, in pots and open water storage containers and infects humans (Fineberg & Wilson, 2010). Forecasts of increasing urbanization in the future are ominous and raise concerns about the development of new infectious diseases. It is estimated that by 2050, 66% of the world’s population will live in cities and the total area of urban areas worldwide will increase by more than 1,5 million square kilometers by 2030 (Seto, Güneralp, & Hutyra, 2012).

4.2.3. The migration of populations

The migration of populations from countries with a high epidemiological burden is a means of invading infectious diseases in host countries where these diseases are not endemic. The recurrence of brucellosis in Germany in recent years is associated with increased migration flows from Turkey where the disease is endemic (Al Dahouk et al., 2007). Also, Chagas disease (CD), a parasitic disease endemic to Latin American countries, now appears in non-endemic areas such as Europe. It is estimated that 68,000 to 120,000 people carriers of the disease live in European countries and that 30% may develop gastrointestinal disorders (Requena-Méndez et al., 2015; Troy, Rickman, & Davis, 2005).

4.2.4. Human behavior

Human behavior (sexual activity, intravenous drug use, etc.) contributes to the development of sexually transmitted diseases such as HIV / AIDS, hepatitis, and tuberculosis (Lederberg, Shope, & Oaks, 1992).

A typical example is a COVID-19 pandemic, where human behavior is directly related to the spread of the disease. When protective measures are applied (mask use, social distancing) then the disease recedes, while when these are circumvented, the virus spreads faster in the human population.

5. The two-way relationship between human activities and zoonotic diseases in the case of the COVID-19 pandemic
The COVID-19 pandemic caused by the coronavirus SARS-CoV-2 that is currently affecting the planet is a much typical example of the two-way relationship between human activities and zoonotic epidemics/pandemics, as shown in the figure below. Figure 7.

**Figure 7. The two-way relationship between human activities and pandemic COVID-19**


5.1.1. Natural and intermediate host

The SARS-COV-2 coronavirus, appeared in December 2019 in China, belongs to the same family (Coronaviridae-b) of the SARS-CoV and MERS-CoV coronaviruses that caused epidemics in 2002 and 2012, respectively (Docea et al., 2020; Mousavizadeh & Ghasemi, 2021). It has a positive polarity RNA genome, encodes a non-structural replicase polyprotein and four structural proteins (spike S, envelope E, membrane M, and nucleocapsid N), and it uses the S protein, which is located on its surface to enter the host cells (Carfì, Bernabei & Landi, 2020; Datta, Talwar, & Lee, 2020; del Rio, Collins, & Malani, 2020).

The high sequence identity of the two coronavirus genes (96.2% for Rhinolophus affinis - RaTG13 and 93.3% for Rhinolophus malayanus - RmYN02) of bats suggests that bats may be the natural reservoir of SARS-CoV-2 (Chan, To, Tse, Jin, & Yuen, 2013; Nguyen et al., 2022; Ren et al., 2020; Zhou, H. et al., 2020; Zhou, P. et al., 2020). Although direct infection of humans with bat viruses has not yet been documented, it could, however, occur in mixed unsanitary environments such as bazaars, where people mix with domestic or wild animals that carry viruses and exchange them with each other, making them more contagious, and easier to be transmitted to humans. (Menachery et al., 2015).

Human interaction with bats or any other wild animal (intermediate host) may be the cause of transmission of SARS-CoV-2 to humans, as the first confirmed cases of SARS-CoV-2 coronavirus were associated with patients, who developed severe pneumonia after visiting the Wuhan Seafood and Wildlife Flea Market. The intermediate host of SARS-CoV-2 may be related to some species of pangolins from Malaysia or Bangladesh. Researchers have identified coronavirus (CoV pangolin) of the same family (Coronaviridae-b) as SARS-CoV-2 and found a close genetic relationship (Xiao et al., 2020) and
a high similarity of the viral sequence\textsuperscript{1} between the two coronaviruses (Lam et al., 2020; Liu et al., 2020; Zhang, Wu, & Zhang, 2020).

In addition to the pangolin, other animals in the Wuhan flea market may have been infected with SARS-CoV-2 and transmitted the virus to humans, such as raccoon dogs and red foxes (Worobey et al. 2022).

According to Shi et al. (2020) and Kim et al. (2020) minks are highly sensitive to SARS-CoV-2 and prone to infection.

An example is the appearance of SARS-CoV-2 on mink farms in Denmark\textsuperscript{1} (Enserink, 2020; Oreshkova et al., 2020; Oude Munnink et al., 2021), in Spain, Sweden, Lithuania, Greece\textsuperscript{11}, Italy, and the USA (WHO, 2020b). Other research has shown that the snakes might also be potential vectors, where SARS-CoV-2 could recombine and acquire the ability to infect humans (Ji, Wang, Zhao, Zai, & Li, 2020). Similarly, the potential susceptibility of domestic pets was also investigated (cats, dogs, pigs, chickens, ducks) in SARS-Cov-2 infection due to their close contact with humans. According to Shi et al. (2020), cats are very susceptible to SARS-CoV-2, dogs have low susceptibility, while pigs, chickens, and ducks are not susceptible to the virus.

5.1.2. COVID-19 disease caused by SARS-CoV-2 and transmission among the human population

Symptoms and clinical picture of COVID-19 disease

COVID-19 is a disease of the respiratory system with mild to very severe symptoms (cough, myalgia, fever, fatigue, and shortness of breath). The clinical picture of the disease ranges from asymptomatic or mild respiratory infection to uncontrolled pneumonia with acute respiratory distress syndrome, multiorgan failure, and death (Guan et al., 2020; Huang et al., 2020; Wang et al., 2020). Complications following the acute phase of the infection, such as rare multisystem inflammatory disease in children and adults, 2 to 5 weeks after initial infection (Datta et al., 2020) is also identified, mainly from the cardiovascular and gastrointestinal systems. The manifestations of those complications were dermatological and mucosal, resembling the Kawasaki disease in children.

Modes of transmission of SARS-CoV-2 and the cause of the pandemic

The SARS-CoV-2 virus is mainly transmitted mainly by respiratory droplets produced by persons infected with the virus (Dhand & Li, 2020) through human contact, or by touching infected surfaces (van Doremalen et al., 2020). Asymptomatic carriers also contribute to the transmission of the disease (Zhao, Lu, Deng, Tang, & Lu, 2020), while fecal transmission is rare (Zuo, Uspal, & Wei, 2020). It however, needs further investigation, since the virus has been detected in fecal samples of symptomatic and asymptomatic patients (Gao, Chen, & Fang, 2020; Zhang, Li, Zhang, Wang, & Molina, 2020). Also, in some patients that were free of respiratory symptoms, the virus was still present in their feces (Wu, Y. et al., 2020).

Close monitoring of wastewater has been shown to contribute to the early detection of SARS-CoV-2, the identification of the true magnitude of the pandemic, and the design of appropriate measures to prevent the transmission of the disease (Mallapaty, 2020; Medema, Heijnen, Elsinga, Italiaander, & Brouwer, 2020). The disease is highly contagious\textsuperscript{12} as it has been proven since the beginning of the pandemic\textsuperscript{13} until today.

The effect of weather and air pollution on the spread and mortality of COVID-19 disease

The effect of weather on the spread and mortality of COVID-19 disease has been the subject of many studies. Some have shown that the rate of spread of the disease decreased with increasing temperature and humidity (Chatziprodromidou, Apostolou, & Vantarakis, 2020; Wang & Di, 2020).

Research in 166 countries (\textbf{Error! Reference source not found.}), showed that the increase in temperature (+ 1 °C) and relative humidity (+ 1%) was associated with a decrease in daily epidemiological data\textsuperscript{1}. Similarly, according to Qi et al. (2020) daily incidence\textsuperscript{1} in China decreased significantly when the average daily temperature and relative humidity increased. The critical temperature value leading to a decrease in the rate of exponential transmission of the disease was also studied and it was found that at an ambient temperature of 30 °C, the basal rate of reproduction (R0) was about 1, while it increased significantly (R 0 =2.5) when the temperature reached 0 °C (Livadiotis, 2020).

However, according to Paraskevis et al. (2021), climatic conditions alone cannot lead to a reduction in cases and prevent the emergence of new outbreaks of COVID-19, in the absence of public health protection measures. Other studies also reached this conclusion when they incorporated public health
protection measures into their research models (Jüni et al., 2020; Oliveiros, Caramelo, Ferreira, & Caramelo, 2020).

Many researchers have focused on the possible relationship between air pollution and COVID-19 mortality. According to Conticini et al. (2020), high mortality from COVID-19 (> 12%) in Northern Italy (Lombardy, Emilia-Romagna, Piedmont) was associated with high levels of air pollution and increased rates of population aging. Ogen's study (2020) found that very high mortality rates (83%) in northern Italy and Madrid, Spain were associated with high concentrations of carbon dioxide (> 100 μmol/m2).

A similar conclusion was reached by studies (Travaglio et al., 2021) for the urban areas in England (London, Midlands) and for the USA (Wu et al., 2020), which highlighted the relationship between long-term exposure of the human population to harmful particulate matter (PM2.5) and mortality risk from COVID-19.

The pandemic from its outbreak until today has had a huge impact on health and all areas of human activities. The impact was spectacular within the first year of the pandemic, as the virus was unknown, and there were no treatments to prevent and suppress COVID-19. Two years later, the pandemic continues to plague the world's population as the virus mutates and becomes more contagious. But societies are gradually opening, and human activities are returning to more normal levels, as they now have more knowledge about the virus, there are vaccines, medicines, early detection methods and tried and tested crisis management policies.

5.2. Public health and health systems

5.2.1. Infections and deaths from COVID-19, mortality rates, excess mortality

Public health has been hit hard by the COVID-19 pandemic, with millions of cases and deaths worldwide (553,77 million confirmed cases, 6.35 million confirmed deaths, as of July 7, 2022). The regions most affected are Europe and the Americas, followed by Asia, Africa, and Oceania (Our World in Data, 2022).

However, this image is not representative of the real situation. An estimate of the pandemic’s health footprint could be derived from the case fatality rate (CFR), which translates as the ratio between confirmed deaths and cases, but this approach also appears to be precarious. The actual impact of the pandemic may differ greatly from the picture that emerges from the reported data (infections and deaths). The reasons for this difference may be due to limited controls, the way cases and deaths are recorded by each country and the problems of assigning the cause of death. The countries of the African continent, for example, are characterized by inadequate health systems and incomplete infectious surveillance structures. According to the WHO, the measurement of excess mortality is the parameter that can essentially contribute to understanding the impact of the pandemic and to the comparison of mortality estimates between countries, even when data are incomplete or unavailable. The first estimate of excess mortality emerged under an innovative methodology developed by WHO and the United Nations and showed that it was approximately 14.9 million between 1 January 2020 and 31 December 2021 (WHO, 2022b).

5.2.2. SARS-CoV-2 variants and the waves of the pandemic

Two and a half years since the start of the pandemic, the SARS-CoV-2 virus, in its attempt to survive, is constantly mutating and appearing with new variants which, according to the WHO, are distinguished into variants of interest (VOI) and variants of concern (VOCs). Some of them caused strong waves of infections, with the main characteristics being high transmissibility, increasing virulence and reducing the effectiveness of therapeutic methods and vaccines. Table 5.

Table 5. Variants of SARS-COV-2
The prevailing variant of concern today is Omicron (B.1.1.529) which appeared in November 2021 and between 10 June-10 July 2022 (WHO, 2022a) accounted for 84% of viral sequences. In the past, four other variants of concern (Alpha, Beta, Gamma, and Delta) have emerged, causing less or equally significant impact on global public health. The Alpha variant (B.1.1.7) mainly affected Europe, the United Kingdom where it appeared in September 2020, North America, Asia and to a lesser extent Oceania and South America. The Betta (B.1.351) that appeared in South Africa in May 2020 mainly affected Africa, as did the Gamma (P.1) that appeared in Brazil in 2020 and affected South America. Particularly aggravating was the contribution of the Delta variant (B.1.617.2) to public health (GISAID, 2022). The variant first appeared in India in October 2020 and affected all continents equally, causing a strong wave of infections and deaths. By October 2021 it was the dominant variant accounting for 90% of viral sequences (WHO, 2022a) Table 6 (GISAID, 2022).

Table 6. Spread of variants of concern (VOCs) by continent

<table>
<thead>
<tr>
<th>Variants of concern (VOCs)</th>
<th>Currently circulating</th>
<th>Variants of interest (VOI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gamma (P.1): Brazil, November 2020</td>
<td></td>
<td>Theta (P.3): Philippines, January 2021</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Kappa (B.1.617.1): India, October 2020</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Lambda (C.37): Peru, December 2020</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Mu (B.1.621): Colombia, January 2021</td>
</tr>
</tbody>
</table>

The prevailing variant of concern today is Omicron (B.1.1.529) which appeared in November 2021 and between 10 June-10 July 2022 (WHO, 2022a) accounted for 84% of viral sequences. In the past, four other variants of concern (Alpha, Beta, Gamma, and Delta) have emerged, causing less or equally significant impact on global public health. The Alpha variant (B.1.1.7) mainly affected Europe, the United Kingdom where it appeared in September 2020, North America, Asia and to a lesser extent Oceania and South America. The Betta (B.1.351) that appeared in South Africa in May 2020 mainly affected Africa, as did the Gamma (P.1) that appeared in Brazil in 2020 and affected South America. Particularly aggravating was the contribution of the Delta variant (B.1.617.2) to public health (GISAID, 2022). The variant first appeared in India in October 2020 and affected all continents equally, causing a strong wave of infections and deaths. By October 2021 it was the dominant variant accounting for 90% of viral sequences (WHO, 2022a) Table 6 (GISAID, 2022).

Table 6. Spread of variants of concern (VOCs) by continent
In the first months of the pandemic, the resilience of health systems was shaken. Many countries, underestimating the risk of the disease (e.g., USA) or adopting the theory of “herd immunity” (e.g., United Kingdom, Netherlands, Belgium, Luxembourg, Sweden), did not take immediate precautionary and restrictive measures (Doumas, Imprialos, Patoulias, Katsimardou, & Stavropoulos, 2020).

In contrast, countries that adopted timely measures such as lockdown and large-scale case detection tests have succeeded in reducing the spread of the disease. According to Moris & Schizas (2020), the time between the onset of the first confirmed case per million population and the imposition of the quarantine measure proved to be a critical factor.

Greece, a country with the lowest funded health system (8.45% of GDP) in Europe, imposed an immediate (within six days) general lockdown and in one month had only 63 deaths and 1,613 confirmed cases. On the other hand, the delay in imposing the lockdown (13 to 18 days), contributed to the rapid spread of the disease, in countries (Italy, Spain) with gross public health costs, like Greece. In the first 30 days, Italy recorded 63,927 cases and 6,077 deaths, while Spain recorded 87,956 cases and 7,716 deaths.

The countries of East Asia with the immediate imposition of a general lockdown significantly reduced the spread of the disease in Europe and North America. China, for example, imposed a lockdown one day after the first confirmed case per million population, resulting in one-month mortality at just 1.86 per million population. In the USA, on the other hand, unprecedented chaos occurred and the world’s most prepared country for infectious disease management, according to the Global Health Insurance Index (Nalabandian et al., 2019), was hit hard.

According to WHO (2020a), significant disruptions were caused to health services provided which were reduced, suspended or discontinued (emergency: 22%, emergency procedures: 19%, blood transfusions: 23%, suspension of dental care, outpatient follow-up: 76%, routine examinations: 70%, diagnostic treatments: cancer 55%, mental health 61%, non-communicable diseases 69%, malaria 46%, tuberculosis 42%, antiretroviral therapy 32%, family planning and contraception 8%).

### 5.2.3. Health Systems

<table>
<thead>
<tr>
<th>Continent</th>
<th>Variants duration (V, d) based on smoothing rate (&gt;0.1% - &lt;0.1%) / Period and value of maximum prevalence (P, Vmaxp)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Europe – U. K</td>
<td>V: d: 10/2020-10/2021 not affected not affected 3/2021-4/2022 10/2021-7/2022 P: vmax: 3/2021 (96.9%) 4/2021 (0.7%) 4/2021 (0.2%) 10/2021 (99.9%) 7/2022 (99.9%)</td>
</tr>
<tr>
<td>North America</td>
<td>V: d: 10/20-10/2021 not affected 2/2021-10/2021 10/2020-5/2022 10/2021-7/2022 P: vmax: 5/2021 (59.9%) 3/2021 (0.1%) 2/21 (13.6%) 11/2021 (99.4%) 7/2022 (99.2%)</td>
</tr>
<tr>
<td>South America</td>
<td>V: d: 12/2020-10/2021 not affected 12/2020-12/2021 11/2020-6/2022 10/2021-7/2022 P: vmax: 3/2021 (8.4%) 4/2021 (0.8%) 6/2021 (76.5%) 11/2021 (95.6%) 7/2022 (99.5%)</td>
</tr>
<tr>
<td>Asia</td>
<td>V: d: 11/2020-11/2021 12/2020-9/2021 not affected 10/2020-5/2022 10/2021-7/2022 P: vmax: 5/21 (46.6%) 3/2021 (3.9%) 3/2021 (0.2%) 11/2021 (97.9%) 7/2022 (98.8%)</td>
</tr>
<tr>
<td>Oceania</td>
<td>V: d: 10/2020-10/2021 11/2020-8/2021 not affected 1/2021-5/2022 10/2021-7/2022 P: vmax: 2/2021 (34%) 1/2021 (7.3%) 4/2021 (1%) 11/2021 (99.9%) 7/2022 (99.5%)</td>
</tr>
<tr>
<td>Africa</td>
<td>V: d: 10/2020-12/2021 10/2020-12/2021 not affected 10/2020-5/2022 10/2021-7/2022 P: vmax: 4/2021 (23.5%) 1/2021 (49.7%) 9/2021 (92.6%) 7/2022 (99.6%)</td>
</tr>
</tbody>
</table>

- **Alpha:** affected all regions of the world differently
- **Beta:** significantly affected only Africa
- **Gamma:** significantly affected only South America
- **Delta:** significantly affected all regions of the world
- **Omicron:** dominates in all regions of the world

### 5.3. Global and European economy
The COVID-19 pandemic has led the world economy into a recession which, according to the World Bank, is unique as it is the first to be caused exclusively by a pandemic in the last 150 years, but also deeper than that of World War II.

According to the Organization for Economic Co-operation and Development (OECD), world GDP shrank by 4.2% in 2020, while the global economy is expected to recover in the next two years (OECD, 2020b). Government debt has risen to almost 100% of GDP due to the huge cost of policies to rescue national economies and reduce tax revenues (OECD, 2020a).

In Europe, the pandemic negatively affected all indicators of the European economy (GDP, unemployment, public debt). According to the European Commission (2020), the economies of the EU and the euro area countries shrank by 7.4% and 7.8% respectively in 2020 and will gradually recover over the next two years (2021: + 4.1% and + 4.2% respectively, 2022: + 3%).

Unemployment has risen dramatically, and it is estimated that 88-115 million people will be plunged into extreme poverty, mainly in South Asia and sub-Saharan Africa (Blake & Wadhwa, 2020).

In Europe, the figures in 2020 and 2021 are overwhelming (EU: 7.7% in 2020, 8.6% in 2021, euro area: 8.3% in 2020 and 9.4% in 2021) while the expected improvement in 2022 will lag pre-pandemic levels (EU 8.0% vs. 6.7% in 2019, euro area: 8.9% vs. 7.5% in 2019). Deficits and public debt will also increase due to increased expenditures and reduced tax revenues (euro area deficit: 0.6% in 2019, 8.8% in 2020, 6.4% in 2021 and 4.7% in 2022, debt to GDP ratio: 85.9% in 2019, 101.7% in 2020, 102.3% in 2021 and 102.6% in 2022) (European Commission, 2020).

5.4. Global air transport


In 2020, air passenger seats offered for international and domestic travel fell by 50% compared to 2019, which corresponds to 2.699 million fewer passengers, while airlines recorded losses of $ 371 billion in gross revenue. For the first quarter of 2021, the estimates translate into a reduction of the offered passenger seats from 26% to 45% which corresponds to a total reduction of 393 to 635 million passengers, and the loss in gross revenue will amount to 57 to 90 billion dollars (ICAO, 2021).

5.5. Global tourism industry.

In 2020, travel restrictions stopped tourism around the world. In the first eight months, international tourist arrivals fell by 70% and revenue losses skyrocketed to $ 730 billion, eight times higher than in the global financial crisis of 2009.

The picture of international arrivals in the northern hemisphere gradually improved in July (-81%) and August (-79%), compared to the first half (decrease> 90%) of the year (UNWTO, 2020). The viability of mainly small and medium-sized enterprises and jobs was greatly affected (OECD, 2020a). According to the World Travel and Tourism Council (WTTC, 2020), 142.6 million jobs were lost worldwide. Cruise lines lost $ 50 billion in revenue and 334,000 jobs from mid-March to the end of September (CLIA, 2020), while the impact on travel agents was also significant. In the first half of 2020, the travel group TUI reported revenue losses of 75 million euros (TUI GROUP, 2020), and Booking Holdings recorded an 87% decrease in revenue from overnight stays, compared to the corresponding period of 2019 (Booking Holdings, 2020).

The tourism industry will recover from the third quarter of 2021 as predicted by the World Tourism Organization, while the OECD estimates that this will happen by 2022 and return to pre-epidemic levels, not before 2023 (OECD, 2020c).

5.6. World trade

In 2020, trade in goods and services was adversely affected by border closures and cross-border restrictions. In the first half, trade in products decreased by 14.3% compared to the same period in 2019, while the decline in trade in services reached 23%, well above the 9% recorded during the financial crisis of 2009. Impacts were higher in Europe (exports: -24.5%, imports: -19.3%) and North America (exports: -21.8%, imports: 14.5%), compared to Asia (exports: -6.1%, imports: -7.1%). However, not all categories of tradable products and services were equally affected. The sectors of the automotive industry (-70%) fuel and mining products (-38%), industrial (-19%), and agricultural (-5%) products showed large reductions.
In contrast, by September 2020, trade in telecommunications equipment products increased by 9% compared to the corresponding period of 2019, due to teleworking and tele-education, while the increase in PPE trade was spectacular (122% in May and 92% in the second quarter of 2020) (WTO, 2020). In September 2020, travel services (-68%), construction (-16%), and arts and entertainment (-14%) recorded a large decline, compared to the corresponding period of 2019, while they increased legal, administrative, accounting services, financial (+ 2%) and advertising services (+ 1%) (WTO, 2021).

5.7. Food security and nutrition

Severe disruptions occurred in food supply chains due to business closures, high production, and low demand, inadequate storage facilities, restrictions on basic foodstuffs (rice, wheat) from exporting countries, and high levels of contamination among workers. These disturbances affected food availability, prices, and product quality (Barrett, 2020). In Thailand, Vietnam, and the USA the price of rice increased by 32%, 25%, and 10% respectively, between February and April 2020 (Katsoras, 2020).

Imports of agricultural products (seeds, fertilizers) were also affected, and these products became rarer and more expensive in countries such as China and West Africa (Arouna, Soulier, Mendez del Villar, & Demont, 2020; Pu & Zhong, 2020).

In many poor countries, social protection programs (school meals*) stopped with the closure of schools. About 370 million children lost access to these programs, worsening the financial situation of poor households (Moseley & Battersby, 2020). Malnutrition is projected to increase, as an additional 83-132 million people will face food insecurity as a direct result of the pandemic, of which 38-80 million will come from poor countries dependent on food imports (Torero, 2020).

5.8. Education

The closure of schools and universities during the first wave of the pandemic affected 94% of the global student population (1.6 billion children and young people) in more than 190 countries worldwide. To ensure continuity in the educational process, distance home education was implemented. Pre-existing problems in education worsened and inequalities widened. According to UNICEF (2020) at least 463 million students worldwide did not have access to distance education without the necessary technological equipment, a suitable home learning environment, and the required parental support.

Similarly, higher education shifted to online teaching, with many foreign students returning to their homeland, thus losing many of the benefits of international mobility (OECD, 2020a). University institutions in Australia, Canada, the United Kingdom, and the United States that mostly rely on international students were severely affected, as tuition funds account for a large share of their revenue and is a major source of funding for academic research. (OECD, 2020d).

5.9. Society and psychology of citizens

The measures of social isolation, and closure of businesses and activities, affected people's social relationships and daily life, endangering their mental and emotional health. According to Fountoulakis et al. (2021), during the first lockdown in Greece in the period from April 11 to May 1, 2020, daily life was greatly affected, such as physical activity, exercise, appetite, sleep, increased need to strengthen relationships within in the family (increased need for communication> 40%, emotional support 24.16% and improvement in the quality of relationships 24.6%), while the conflicts mainly concerned families with children more vulnerable in confinement and with difficult management behavior.

Separation from loved ones, loss of freedom, uncertainty about the course of the disease (Cao et al., 2020; Li & Wang, 2020), restriction of movement, loss of social contacts, employment, income, transmission, psychological disorders (stress, depression, suicidal behaviors) (Kawohl & Nordt, 2020). Also, loneliness due to social isolation adversely affected the mental and physical health mainly of the elderly and people with disadvantaged socioeconomic status and physical and mental illness (Druss, 2020).

Increased psychological disorders were found in children, students, and healthcare staff. According to Orgiles et al. (2020) children aged 3 to 18 years from Italy and Spain had 76.6% difficulty concentrating, 52% boredom, 39% irritability, 38.8% anxiety and nervousness, and 31.3% feeling of loneliness. Anxiety symptoms developed in students in China due to the slowdown in their academic activities (Alvarez, Argente, & Lippi, 2020), their living in urban areas, the financial situation of their families, (Cao et al., 2020), and the infection of their relatives or friends.

Health personnel showed symptoms of depression (50%), anxiety (44.6%), insomnia (34%), and risk (71.5%) (Lai et al., 2020). Their interpersonal relationships were severed for fear of transmission to their
family and friends (Brooks et al., 2020) and many developed secondary traumatic stress disorder (Zaffina et al., 2014) when asked to choose which patients to be admitted to the Intensive Care Unit (Rana, Mukhtar, & Mukhtar, 2020; Roden-Foreman et al., 2017).

6. Therapeutic developments in the treatment of the pandemic

From the beginning of the pandemic, the scientific community used all the drugs in its quiver to treat patients with COVID-19 as they did not have effective vaccines to treat the disease. The US Food and Drug Administration (FDA) and the European Medicines Agency (EMA) have approved a variety of drugs (COVID19 Vaccine Tracker, 2022a; EMA, 2022) that have been tested to treat SARS-COV-2 infection, pneumonia, multiples organ dysfunction, and thrombolysis, such as antiviral, anti-inflammatory drugs used to treat other diseases (rheumatoid arthritis), anticoagulants and monoclonal antibodies Table 7.

<table>
<thead>
<tr>
<th>FDA Drug</th>
<th>FDA Type</th>
<th>EMA Drug</th>
<th>EMA Type</th>
</tr>
</thead>
<tbody>
<tr>
<td>Veklury (remdesivir)</td>
<td>antiviral</td>
<td>Veklury (remdesivir)</td>
<td>antiviral</td>
</tr>
<tr>
<td>Evusheld (ticagrelor and cilgavimab)</td>
<td>monoclonal</td>
<td>Kineret (anakinra)</td>
<td>anti-inflammatory</td>
</tr>
<tr>
<td>RoActemra (taciluzumab)</td>
<td>monoclonal</td>
<td>RoActemra (taciluzumab)</td>
<td>monoclonal</td>
</tr>
<tr>
<td>Sotrovimab</td>
<td>monoclonal</td>
<td>Regkirona(regdanvimab)</td>
<td>monoclonal</td>
</tr>
<tr>
<td>REGEN-COV (casirivimab/imdevimab)</td>
<td>monoclonal</td>
<td>Ronapreve (casirivimab/imdevimab)</td>
<td>monoclonal</td>
</tr>
<tr>
<td>Bamirnavimab and Etesevimab</td>
<td>monoclonal</td>
<td>Xevudy (sotrovimab)</td>
<td>monoclonal</td>
</tr>
<tr>
<td>Ohumanant (Baricitinib)</td>
<td>anti-inflammatory</td>
<td></td>
<td></td>
</tr>
<tr>
<td>COVID-19 convalescent plasma</td>
<td>recovery plasma</td>
<td></td>
<td></td>
</tr>
<tr>
<td>REGOCIT</td>
<td>substitution solution for (CRRT)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Paxlovid (nirmatrelvir, ritonavir)</td>
<td>PAXLOVID</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Molnupiravir</td>
<td>antiviral</td>
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</table>

Table 7. Medicines that have received emergency approval

Research for new treatments resulted in the development of specialized vaccines less than a year after the pandemic broke out, some of which were approved by international and national health organizations. On the 14th of January 14, 2022, there are 174 candidate vaccines, 557 ongoing vaccines in trials, 33 approved vaccines, and vaccination campaigns in many countries around the world (COVID19 Vaccine Tracker, 2022a).

Depending on the mechanism they use to activate the human immune system, vaccines are divided into two categories: A. Complex virus vaccines, and B. Whole virus vaccines. Table 8.

Table 8. Categories of vaccines and mechanism of human immunization
Of the 33 approved vaccines, 13 belong to the protein subunit vaccine class, 10 to an inactivated virus, 6 to the non-replicating virus vector, 3 to mRNA, and 1 to DNA. Three vaccines have been approved and administered in the United States, while there are five approved vaccines in the European Union. Pfizer-BioNTech (BNT162b2), Moderna (mRNA-1273), and Johnson & Johnson (Ad26.COV2.S) approved vaccines from the US Food and Drug Administration are available in the United States.

In the countries of the European Union, in addition to the above three vaccines, the vaccines AstraZeneca (AZD1222) and Novovax (Nuvaxovid, Covovax, NVX-CoV23) are also administered, which are also approved by the European Medicines Agency (EMA) (EMA, 2022; FDA, 2022).

In addition to Europe and America, other countries have developed vaccines against COVID-19 and have launched systematic vaccination campaigns (COVID19 Vaccine Tracker, 2022a), such as China, Japan, Taiwan, India, Iran, Russia, Turkey, Kazakhstan, Cuba, and Australia. These vaccines are also given in other countries such as Arab, Asian, South American, and some European countries.

Developed and some developing countries have started systematic vaccinations and have vaccinated hundreds of thousands or even millions of citizens since the end of December 2020.

In the Americas, Europe, Asia, and Oceania by July 7, 2022, rates of fully vaccinated citizens exceed 60% (South America: 76%, Asia: 70.78%, Europe: 65.87%, North America: 63.98%, Oceania: 62.43%) and of those who have received at least one dose are over 65% (South America: 84.7%, Asia: 75.98%, Europe: 68.82%, North America: 73.05%, Oceania: 65.16%) (Our World in Data, 2022) Figure 8.
In the African continent, the picture is disappointing, as the percentage of fully vaccinated is only 19.53% and those who have received one dose at 25.08% (Our World in Data, 2022). Nonetheless even this very low vaccination coverage does not represent all countries on the continent. There are countries with rates of fully vaccinated citizens below 10% (Congo: 2.4%, Madagascar: 4.2%, Cameroon: 4.5%, Mali: 6.3%, Burkina Faso: 7%, Somalia: 9, 9%, Sudan: 9.9%) and countries with rates exceeding 50% (Morocco: 63.2%, Rwanda: 64.9%, Botswana: 58.4%, Tunisia: 52%) (COVID19 Vaccine Tracker, 2022b).

The picture is bleak as the People's Vaccine Alliance report of October 2021 shows that rich countries are accumulating large stocks of doses and failing to meet their commitments to poor countries, while pharmaceutical companies are exploiting their monopolies to gain enormous wealth (Malpani & Maitland, 2021).

On one hand the report shows that the G7 and the Group of Europe (EU plus Iceland and Norway) have delivered 261 million of the promised 1.8 billion tranches to low- and middle-income countries, while more than 10 million have been disbursed to other high-income countries, such as Canada, and the United Kingdom, which received approximately 1.5 million doses of vaccine from COVAX in 2021.

On the other hand, these two countries have broken their promises, as the United Kingdom has delivered only 9.6% and Canada 8% of the promised installments. Similarly, other countries are failing to meet their commitments, such as Germany delivering only 12% of the promised 100 million installments, France only 9% of the 120 million installments, and the US 16% of the 1.1 billion installments (WTO, 2021). Pharmaceutical companies also promote vaccines in rich countries and do not meet their commitments to COVAX and the Vaccine Acquisition Trust (VAT)\(^v\).

In 2021, four major pharmaceutical companies (AstraZeneca, Pfizer / BioNTech, Moderna, and Johnson, and Johnson) provided a total of 47 times more installments in high-income countries than in low-income countries. Rich countries representing only 16% of the world’s population bought about 49% of these companies vaccines, while in poor countries through COVAX and AVAT mechanisms not even 50% of the agreed doses were delivered. Table 9.

Table 9. Agreed Supply and deliveries of doses of vaccines by [G7, AstraZeneca, Pfizer/BioNTech, and Moderna, to COVAX and AVAT (for 2021)]

<table>
<thead>
<tr>
<th>Pharmaceutical Company /Vaccine</th>
<th>COVAX</th>
<th>AVAT</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Agreed supply</td>
<td>delivery</td>
</tr>
<tr>
<td>J&amp;J (Ad26COV21)</td>
<td>20000000</td>
<td>0%</td>
</tr>
<tr>
<td>Oxford and AstraZeneca (AZD1222)</td>
<td>72000000</td>
<td>14%</td>
</tr>
<tr>
<td>Pfizer/BioNTech (BNT162b2)</td>
<td>4000000</td>
<td>39%</td>
</tr>
<tr>
<td>Moderna (mRNA-1273)</td>
<td>3400000</td>
<td>0%</td>
</tr>
</tbody>
</table>

Vaccination of the world population is two-speed. Rich countries already promote third- or fourth-dose vaccines (Israel) in their populations and vaccinations in adolescents and children, whereas in poor countries, due to lack of vaccines, high-risk population groups (e.g., elderly, and public health workers) have not been vaccinated (WHO, 2021). In the Americas, Asia, Europe, and Oceania the percentage of the population that has received a booster dose of vaccines exceeds 28%, (Asia: 28.98%, North America: 37.26%, Oceania 38.16%, Europe 41.30%, South America: 48.48%), while in Africa it reaches only 2.26% (Our World in Data, 2022).

As the pandemic spreads to every corner of the globe, equitable access to vaccines and treatments for all of humanity is the only answer to the global health crisis. The more vaccinated the world’s population is, the less likely the virus is to spread, mutate and make existing vaccines less effective (Williams & Burgers, 2021).

7. Discussion-Conclusions

Humans’ relationship with pandemics follows a two-way path. Humans, through their activities, disturb the environment and natural ecosystems and come into direct contact with wild animals that carry pathogens, resulting in their infection with zoonoses and the occurrence of epidemics-pandemics. The effects of pandemics have as the final recipient the human and his activities.
"We are invading the tropical forests, the wild nature where so many species of plants and animals live and with them so many viruses unknown to us. We cut down the trees, kill the animals or capture them and send them to the markets. We disrupt ecosystems and drive viruses out of their natural hosts, making them search for new ones. Well, we usually become their new home," characteristically states (Quammen, 2012), in his book "Spillover: Animal Infections and the Next Pandemic."

The COVID-19 pandemic is a prime example of this two-way relationship. Human activities (trade, sale and consumption of wild animals) in the open market of the city of Wuhan in China, probably contributed to the creation of a favorable environment for the transmission of the SARS-CoV-2 from bats (natural host) to humans, probably through a wild animal-intermediate host (pangolin, raccoon dog, red fox, mink, snake etc.). The virus was easily transmitted to the human population through respiratory droplets and human contact and spread across the globe through aviation, travel, and trade, causing the COVID-19 pandemic. The pandemic since its inception has caused enormous impacts on public health, health systems, the global economy, transportation, trade, tourism, education, and human societies.

The global medical and scientific community since the beginning of the pandemic used all the therapeutic means at its disposal to relieve symptoms and treat patients, developing safe and effective vaccines and continuing research to produce new vaccines and drugs for the control and mitigation of the pandemic.

The ideal scenario would be the eradication of the pandemic from every corner of the planet as happened in the past with the eradication of smallpox and poliomyelitis. However, this scenario is not certain to be repeated in the case of the pandemic, as shown by the behavior of the virus to date (mutations, the emergence of new variants, causing successive waves of infections). Humanity will have to learn to coexist with a controlled virus as it does with other viruses (flu virus). By increasing the vaccination coverage of the global population, the chances of mitigating the dynamics of the virus and its ability to mutate into something more powerful, spread, and affect the effectiveness of vaccines are enhanced. The stake is high and will only be won if country leaders, pharmaceutical companies, and international and national health organizations work together to ensure fair and equal access to vaccines and treatments for all people.

In addition to this bet, the pandemic brought to the surface other important issues that humanity needs to face, such as the protection of the natural environment, the elimination of illegal wildlife trade, and the prevention of zoonotic diseases. The answer to these questions can only be given if bold decisions are taken in the context of global cooperation by adopting policies to preserve and restore natural ecosystems, impose strict control regulations on the movement, trade, and consumption of wild animals, and develop modern zoonotic surveillance systems.

The benefit of these decisions would be very large, comparing the estimated cost of prevention policies ($22 - 31.2 billion) with the economic impact of the pandemic ($16 trillion by the end of 2021) if vaccines will prove effective in pandemic control. The costs of prevention strategies could be reduced even further ($17.7-26.9 billion) if the benefits of reducing deforestation are accounted for (IPES, 2020).

Today many financial possibilities exist, although the question is whether the resources will be distributed in interventions not only for the recovery and resilience of national economies but also in prevention policies, which would significantly reduce the risk of a new pandemic.

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Notes

1 Paguma lavrata: a species of muskrat found in the jungles of SE Asia, in the Indian subcontinent, as well as on tropical islands around the Asian continent such as Taiwan and Sumatra

2 Pangolin: Mammal of the order of Scales. There are 8 species of pangolins in Asia and Africa that are threatened with extinction due to poaching of meat and their scales intended for medical use. According to the UN Report on Wildlife Crime, it is estimated that 71% of the confiscated pangolin shells from 2007 to 2018 were destined for China. In 2019, Nigeria became the main export point of pangolins and Vietnam the primary destination (UNODC, 2020)

3 The El Niño (ENSO) Phenomenon during which the surface waters of the central and eastern Pacific Oceans are characterized by unusually warm temperatures. The effects of El Niño are characterized by excessive rainfall along the coasts of Central and South America and increased drought in parts of the western Pacific that often leads to catastrophic fires such as in Australia. El Niño occurs about once every 20 years, but it is estimated that the number of such events could double (one every 10 years) if the global average temperature increases up to 1.5°C (Wang, Zhang, Cole, & Chavez, 1999)

4 the E, M, N and S genes encoding structural proteins showed 100, 98.6, 97.8, and 90.7% amino acid identity

5 High similarity (85.5-92.4%) of the viral sequence between pangolin-nCoV and SARS-CoV-2

6 In Denmark, from June to November 2020, a total of 644 cases of COVID-19 were detected in people associated with mink farms, while 12 of them detected a mutation in the mink-associated SARS-CoV-2 virus (cluster 5). The Danish government has killed 17 million minks in 1,000 farms across the country to prevent the spread of the disease among farms, as well as the possibility of a new pandemic from mink to humans due to the virus mutation

7 There are 90 farms in Greece with 1 million minks. After the appearance of SARS-CoV-2 cases on a farm in Kaloneri, Kozani, 2,500 mink were killed

8 the reproduction rate (RO) is estimated to be between 2 and 3 (Omer, Yildirim, & Forman, 2020)

9 in the early days of the epidemic in China, the average "R0" index was 5.7 ranging from 3.8 to 8.9 (Sanche et al., 2020)

10 Temperature rises 1 °C: 3.08% new cases and -1.19% new deaths. Relative humidity increases 1%: 0.85% new cases and 0.51% new deaths

11 the daily cases decreased by 36% -57% with an increase of the average daily temperature (+ 1 °C) and by 11% -22% with an increase of the relative humidity (+ 1%)

12 an increase of 1 mg / m3 of particulate matter (PM2.5) in the atmosphere was associated with an 8% increase in COVID-19 mortality rates

13 Cost of rescue policies: about 7.5 trillion dollars (14% of GDP) in advanced countries and 150 billion dollars (3.6% of GDP) in emerging market economies (IMF, 2020)

14 in the poorest countries, school meals are the only daily food received by children of families living in extreme poverty as well as health programs (vaccinations, dehydration, and iron supplements)

15 G7: Informal group consisting of the 7 seven most industrialized countries in the world (USA, Canada, Japan, France, Germany, Italy, and the United Kingdom). The G7 meets annually on issues such as global economic governance, international security, and energy policy

16 COVAX: Global Initiative, coordinated by the Global Alliance for Vaccines and Immunization (GAVI), the Coalition for Epidemic Preparedness Innovations (CEPI), and the World Health Organization (WHO), to ensure equitable access to vaccines for all COVID countries, 19 regardless of income. 92 low-and low-middle-income countries will have access to vaccines through the Financial Instrument (AMC)

17 AVAT: Central Vaccine Market Agent of the African Union (AU) the Member States and secures funding to vaccinate at least 70% of the Union’s population. Key partners of AVAT are the African Union Centers for Disease
Control and Prevention (Africa CDC), the African Export-Import Bank (Afreximbank), and the United Nations Economic Commission for Africa (ECA).