Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active.
Review

Three waves changes, new variant strains, and vaccination effect against COVID-19 pandemic

Rehan M. El-Shabasy\textsuperscript{a,\textsuperscript{*,1}}, Mohamed A. Nayel\textsuperscript{b}, Mohamed M. Taher\textsuperscript{c,\textsuperscript{*}}, Rehab Abdelmonem\textsuperscript{d}, Kamel R. Shoueir\textsuperscript{e,\textsuperscript{f}}, El Refaie Kenawy\textsuperscript{g}

\textsuperscript{a} Department of Chemistry, Faculty of Science, Menoufa University, 32512 Shebin El-Kom, Egypt
\textsuperscript{b} Department of Animal Medicine and Infectious Diseases, Faculty of Veterinary Medicine, University of Sadat City, Sadat City 32897, Menoufa, Egypt
\textsuperscript{c} Department of Chemistry, Faculty of Science, Cairo University, 12613 Giza, Egypt
\textsuperscript{d} Department of Industrial Pharmacy, Faculty of Pharmacy, Misr University for Science & Technology, 6th October, Egypt
\textsuperscript{e} Institute of Nanoscience & Nanotechnology, Kafrelsheikh University, 33516 Kafrelsheikh, Egypt
\textsuperscript{f} Institut de Chimie et Procédés pour l’Energie, l’Environnement et la Santé (ICPEES), CNRS UMR 7515-Université de Strasbourg, 25 rue Becquerel, 67087 Strasbourg, France
\textsuperscript{g} Polymer Research Group, Chemistry Department, Faculty of Science, Tanta University, Tanta, Egypt

\textbf{ARTICLE INFO}

\textbf{Keywords:}
COVID-19
SARS-COV-2
Mutation
New variant strain
Vaccine
Reinfection

\textbf{ABSTRACT}

It has been more than one year since the first case of the coronaviruses was infected by COVID-19 in China. The world witnessed three waves of the corona virus till now, and more upcoming is expected, whereas several challenges are presented. Empirical data displayed that the features of the virus effects do vary between the three periods. The severity of the disease, differences in symptoms, attitudes of the people have been reported, although the comparative characteristics of the three waves still keep essentially indefinite. In contrast, the sense of danger toward the cries gradually decreases in most countries. This may be due to some factors, including the approved vaccines, introducing alternative plans from politicians to control and deal with the epidemic, and decreasing the mortality rates. However, the alarm voice started to rise again with the appearance of new variant strains with several mutations in the virus. Several more questions began to be asked without sufficient answers. Mutations in COVID-19 have introduced an extreme challenge in preventing and treating SARS-COV-2. The essential feature for mutations is producing new variants known by high tensmibility, disturbing the viral fitness, and enhancing the virus replication. One of the variants that has emerged recently is the Delta variant (B.1.617.2), which was firstly detected in India. In November 2021, a more ferocious mutant appeared in South Africa, also called omicron (B.1.1.529). These mutants grabbed world attention because of their higher transmissibility than the progenitor variants and spread rapidly. Several information about the virus are still confusing and remains secret. There are eight approved vaccines in the market; however, the investigation race about their effect against reinfection and their role against the new variants is still under investigation. Furthermore, this is the first time vaccinating against COVID-19, so the question remains: Will we need an annual dose of the corona vaccines, and the side effects don’t been observed till now?

1. Introduction

Since World War II, the COVID-19 pandemic has become the most critical global health emergency in this century and the greatest challenge for the human population [1]. In addition to being a global health calamity, coronaviruses pandemic have several critical effects in all fields of life, including; environmental, economic, social, political, and cultural. The first wave was so dangerous that almost no place on earth was saved from the impact of this epidemic, despite the differences in seasons; the southern hemisphere was affected later, but no less severely [2]. It is worth mentioning that the rate of cases and death in Africa, except in South Africa, was lower. This may be due to the low average age and Ebola disease some years ago, which helped to experience and decrease the problem [3]. The World Health Organization (WHO) is officially announced COVID-19 as a pandemic on 11 March 2020. This is because increasing the cases number dramatically outside China. On

\textsuperscript{*} Corresponding authors.
E-mail addresses: elshabasy1010@gmail.com, rehansh1010@yahoo.com (R.M. El-Shabasy), m.taher923@gmail.com (M.M. Taher).

https://doi.org/10.1016/j.ijbiomac.2022.01.118
Received 30 November 2021; Received in revised form 16 January 2022; Accepted 18 January 2022
Available online 22 January 2022
0141-8130/© 2022 Elsevier B.V. All rights reserved.
April 10, 2020, 1.5 million cases were confirmed in 184 countries and more than 92,000 deaths worldwide [1]. In February of this year, around 115 million cases became infected, and the death rate was estimated at more than 2.5 million worldwide [4]. Most individuals infected by COVID-19 were mild to moderate respiratory illness and recovered without specific treatment. However, older adults with medical complications are at a higher risk of severe prognosis [5,6].

On the other hand, environmental conditions could affect the present pandemic of COVID-19 [7,8]. Several studies have investigated the effect of temperature that plays a significant role in virus survival [1,8]. For instance, Chin and his group are recently reported that SARS-CoV-2 is very stable at 4 °C, but the sensitivity of the virus increased toward heat [9]. The survival time was reduced to 5 min as the incubation temperature rose to 70 °C. Epidemiological investigations have revealed the connection between COVID-19 and meteorological parameters; however, findings are controversial [10,11]. A published study by Xie and Zhu described that a 1 °C increase was connected with a 4.86% rise in the daily infected cases of COVID-19, when mean temperature (lag 0–14) was lower than 3 °C [11]. A positive relation between diurnal temperature range and daily deaths of COVID-19 and a negative correlation for relative humidity was recently reported [10].

In contrast, Yao and his colleges have reported that the transmission of COVID-19 did not show an association with temperature in different Chinese cities [12]. The viral mutations are also seasonable independent. It is more expected to happen in winter, spring, and autumn, whereas not preferable in summer [13]. The virus’s slow spreading in Summer may be related to the higher temperature, as established by a temporal multivariate time series model in study [14]. Preventive methods are the offered approach to avoid cases spreading because an epidemic will increase if we cannot control it very well. Protective strategies focus on careful control against infection and patient isolation, including suitable measures to be accepted through diagnosis and the introduction of effective clinical care to infected patients [15]. The current situation is that the coronavirus is spreading elsewhere while facing significant immune pressure from vaccines and naturally infected and recovered people [4,16].

2. Symptoms of COVID-19

With increasing numbers of infected cases, there is a simultaneous growing number of recovered patients. Not all people who have recovered from SARS-CoV-2 infection are free of symptoms. The symptoms also become changeable. Several reports have revealed bone and joint pain, continued fatigue, dizziness, insomnia, palpitations, and headaches. Other symptoms were appeared for irreversible pulmonary scarring and dysfunction, particularly in cases with severe pulmonary disease [17]. Several information about the viruses are still scare till now. The long-term investigations remain insufficient, and these patients’ outlook is still wholly missing, although several studies have addressed this issue [5]. The long-COVID refers to various symptoms affecting different organs reported by patients following COVID-19 infection [18]. For example, recent research displayed that half of cases (478 patients) after four months of COVID-19 hospitalization get one feature of long-term COVID at least [19]. Another report with 4182 instances of COVID-19 appeared that 13% of respondents self-reported long-COVID features, with some evidence for higher rates in women and older people [20]. The additional study followed 1733 patients hospitalized for COVID-19 for six months and found fatigue or muscle weakness in 63%, sleep difficulties in 26%, anxiety or depression in 23%, and lower rates of myalgia and headache [21]. It is expected that there will be many chronic consequences of COVID-19 beyond that will exceed the first wave of acute infections, and still, the coming time may reveal a lot of secrets. One long-term effect of COVID-19 that is becoming noticeably obvious is its influence on cognitive function and in patients with mild symptoms. Neurological symptoms appeared with one-third of COVID-19 patients, and there have been anecdotal accounts of ‘COVID-19 delirium’, manifesting as agitation, paranoid hallucinations, and confusion with a lot of hospitalized cases (<20%) [22,23]. Patients over 65 years old are considered one of the groups most susceptible to severe manifestations of COVID-19. Usually, they have mild cognitive impairment; therefore, the risk increased for developing delirium because of the underlying neurocognitive impairment [24,25]. Inflammation associated with COVID-19 developing blood-brain barrier, silent infarcts, coagulopathy, thrombosis that could increase neurological injury [26]. In addition to poor patient outcomes, the severe agitation associated with delirium in many COVID-19 patients creates difficulties for staff and compounds the stress of caring for these highly sick patients.

On the other hand, risk factors (Fig. 1) for severe SARS-COV-2 infection and increasing mortality rates comprise smoking, advanced age, and medical comorbidities, e.g., diabetes mellitus, hypertension, and obesity, the most common [27,28]. These risk factors showed a baseline neurocognitive frailty which enhanced the exposure to cognitive complications during inflammation cases [29]. Similar to perioperative neurocognitive disorders associated with surgery and anesthesia. Therefore, the maximum risk persons for intense SARS-COV-2 infection may also display the most fundamentally susceptible people for cognitive decline in the case of SARS-COV-2 inflammation.

3. Development of COVID-19 between three waves

Different countries worldwide have faced a three-wave pattern of reported cases; the first wave occurs in spring, the second period at the end of summer and autumn, and the third at the beginning of 2021. Many changes happened between the three waves of COVID-19 around the world, as tabulated in Table 1. The people's attitude toward the crisis was variable, and the sense of danger decreased gradually across the globe. This is may be due to some factors such as (i) vaccination availability, (ii) rate of death (iii) alternative plans from governments, as revealed in Fig. 2. For example, the first wave of infections in the USA started in March 2020, whereas the emergency was announced in the country [5]. The epidemic hit the area for the first time without any previous alarm, and the fighting tools were overwhelmed. Critically ill patients increased over the government's capacity, operating rooms were used as intensive care units, and temporary satellite hospitals were initiated to apply care for non-critical cases [30]. The country succeeded in flattening the curve after extreme mandatory precautions. After some months, different states started returning to their normal lives with ignorance of their stay-at-home restrictions. Hence the number of patients increased at a surprising rate, particularly in western and southern states [31]. Although epidemiologists still debate that the coronaviruses pandemic is far from over, the widespread administration of a COVID-19 vaccine helped remove the high-hearted fear of people [32].

3.1. 1st wave effect

The SARS-COV-2 epidemic was started in Wuhan, China, with the appearance of the first case in December 2019; however, the first wave was internationally detected in March 2020. The main causes of this wave are still secret; it looked like a ghost and fear from the unknown. The first wave was considered a disaster that heavily affected every field in life and represented a significant challenge in public health and disturbed social and economic activities globally [33]. For example, the economy was negatively affected worldwide; good production was almost stopped in all places, and the unemployment rate was highly increased. Another social problem appeared, e.g., violence in the family was increased significantly between children due to the extended stay at home [2].

3.2. 2nd wave effect

Owing to the low numbers of infected cases in the season summer
and also in that time mostly the infected young patients were exit from hospitals and became almost empty, some places of society were over thought the pandemic, disregarding the initial announcement for the coming second wave [2]. Like the other viruses, e.g., influenza blaze up seasonally, COVID-19 resumed in autumn as predictable. As several experts predicted, the second wave hit the countries with a much higher force than the first. The leading cause of robust second wave was not a relaxation of intervention, but rather the failure to enforce interventions as occurred in Europe [34].

During the second wave, politics were changed with different priorities than the 1st wave. In Germany, for example, the instructions were very sharp at the beginning of this pandemic, and the government managed very well in the 1st wave, but this position was changed quickly. This is may be due to some different factors and altered instructions from politicians. Now economic life has become the priority in the list, and businesses are open continuously rather than strictly decreasing contacts and thus infections and death rates [2]. In the second wave period, a complicated equation appeared as an alarm. Most countries are almost all filled to achieve the balance between successful medical care with a growing economy. During that wave, most countries were lost a lot of valuable time, and the number of infected cases dramatically increased [2]. However, seroconversions showed a lower effect than the 1st wave. This is because of the high restrictions of outdoor activities, obligatory face masks, and prevention of any people gathering. Such precautions caused hindrance for viral inoculation, whereas high inocula led to more severe SARS-CoV-2 infection [35].

| Subject          | 1st wave   | 2nd wave       | 3rd wave        |
|------------------|------------|----------------|-----------------|
| Time             | March 2020 | July 2020      | After Christmas |
| Precautionary measures | Extremely high (Obligatory) | Mandatory | Neglected from some people |
| Changes in symptoms | Fever, cough, sore throat, chest and muscle pain, confusion, dyspnoea, headache, anemia, and ageonia | - Communal cold, fever, pneumonia, dyspnoea and cough Sudden death | |
| Mortality rate   | Horrible rates | Increasing rates | Lower compared to prior waves |
| Economical effect| Completely stopped | Affected | Less affected |
| Vaccination      | Not available | Under investigation | Available |
| Re-infection     | Not appeared | Available | More common |
| Mutations        | Not present | Not appeared | Present new variants |
| Table 1 Comparison between the three waves of COVID-19. |

* Decreasing the rate of death
* Vaccine availability
* Resume the life normally
* Having alternative plans
* unkown epidemic
* Highly percautions
* Mandatory stay at home
* Increasing the rate of death
* Stop in all fields of the life
* The available information not sufficient
* No vaccine avialble

Fig. 2. Factors affecting the sense of danger toward coronaviruses crisis.
3.3. 3rd wave impact

The third wave of COVID-19 witnessed several changes worldwide and different features observed, such as increasing infected numbers due to home contacts. In addition, significant availability of speedy antigen tests helped in rapid diagnosis and then isolation, whereas severe cases and mortality rates were less than in prior waves. The most notable feature in this wave was the discovery of a new strain (B.1.1.7) that has higher potential transmissibility; hence, re-infections became more common. These several mutations that appeared within the virus may be the main cause of the third wave and the social activities and contacts between people without obligatory precautions. On the other hand, vaccination achieved positive results between patients, health care workers, and the elderly [36].

4. A new strain of COVID-19

Recent studies reported the high affinity of viruses to evade barriers for transmission, mainly when infections are still numerous [36]. Changes in coronaviruses are considered a perfect storm to allow immune escape mutants to emerge. This could initially occur with partial resistance but with possible superior resistance if coronavirus variants develop further [4], it is worthy of mentioning that the higher infections rates, the higher chances of mutations. This way will support the virus to survive and proliferate [36]. Virus evolutions will not be controlled when we approach herd immunity and do not keep restrictions [4].

Globally, with virus transmission, many SARS-COV-2 variants have recently appeared [37]. The different levels of genetic changes are basically due to some factors, including; the global absence of immunity against this new pathogen. In addition, mutation rates of the SARS-COV-2 that encode an enzyme with proof-reading function raise the fidelity of replication processes [38]. A recent study investigates gene sequences of COVID-19 to identify mutations. The results showed that 26,844 single mutations were followed in 203,346 human genomes of SARS-CoV-2, while the most common mutations involved S proteins and NSP3 [13]. By the end of December 2020, around 5000 mutations were recognized in the S protein. Virus mutations that lead to the spread of new emerging strains are distributed worldwide, particularly in the United Kingdom, e.g., lineage B.1.1.7 and variant 20I/501Y.V1. This variant is distributed due to several spike (S) protein changes, including deletion 145, N501Y, deletion 69–70, D614G, A570D, T716I, P681H, D1118H, and S982A [13]. The additional new strain discovered in South Africa (lineage B.1.351, variant 20H/501Y.V2) involved eight mutations through S protein; D80A, L18F, R246I, D215G, E484K, K417N, A701V, and N501Y. The variant 20 J/501Y.V3 and lineage P.1 has also spread in Brazil through three S protein mutations, N501Y, E484K, and K417N, in common with 20J/501Y.V2 [13]. The most significant character of all these variants is that they share N501Y mutation concerning the SARS-CoV-2 spike (S) protein which is considered the primary target of most COVID-19 vaccines. Mutations such as E484K, K417N, and N501Y in the S protein could disturb viral fitness and transmissibility (Fig. 3). However, the ongoing research on the influence of such variants on COVID-19 vaccines still lacks [4,13,39]; hence further and deep investigations are highly recommended.

Another study has revealed that global dispersal and the growing frequency of the COVID-19 spike protein variant D614G during the autumn of 2020 was evocative of a selective advantage, basically in increased transmissibility [40]. A blend of mutations in the B.1.1.7 strain makes it exceptionally infectious, while the variant of B.1.351 can evade antibodies owing to E484K [4] as indicated in Table 2. Lately, the appearance of N501Y mutants associated with super spreading events and outbreaks typically improved transmissibility. Stimulatingly, such variant was initially recognized in Denmark among the infected minks, while emergence in other parts proposes that it might increase independently from such animals [4]. This mutation could support SARS-COV-2 capability for binding with human receptor angiotensin-converting enzyme 2 (ACE2), therefore increasing the spread of the COVID-19 pandemic [41,42].

On the other hand, it is worthy of spotting the light on the new variants that recently appeared in South Africa; Omicron (B.1.529) and Delta in India (lineage B.1.617). The first case of the delta was announced in October 2020, and the variant has an advanced rate of transmission and infection compared with other previously known variants [43]. However, on 9 November 2021, omicron was discovered [44]. With some concern regarding the virus nature, there are no rules for control or ceiling of expectation as the time between the two variants is very short (almost one month), and this didn’t happen before. Additionally, omicron was distinguished by remarkable diffusion speed, whereas the transmission rate of this variant is much higher than the pre-existing variants because of the greater number of mutations [45]. The high number of mutations present in the S protein in the omicron variant may increase the virus’s ability to evade infection-blocking antibodies and other immune responses such as the T cell response. This observation agrees with preliminary evidence suggesting an increased risk of re-infection with omicron compared to different strains, but the information is still scarce [46]. The initial studies observed that the variant has several spike protein mutations, more than 30 mutations in the virus area, which encodes these proteins responsible for the entrance of the virus to human cells [43]. These mutations that happened in omicron are also identified in the previous strains of COVID-19, including alpha and delta variants [47]. However, several omicron mutations have not been previously appeared, which mandates further investigations to determine it. The number of mutations was twice the delta variant. Ten mutations were found on the RBD of the omicron variant, while there were only two in delta variants [48]. A recent study discussed that mutation group (H655Y, N679K, P681H) in the virus is very short (almost one month), and this didn’t happen before. Addi-

![Fig. 3. Examples of recent new strains and mutations with the most significant features of COVID-19.](image-url)
Vaccination effect against the new variants of COVID-19 and AstraZeneca’s ChAdOx1 novel coronavirus 2019 (nCoV-19) [56].

Despite a massive number of hospitalized and death cases, immunity ona, and Russia’s Sputnik vaccines [55], Janssen’s Ad26.COV2.S and vaccines were previously observed in alpha strain and also seen in omicron, and these mutations enhanced the infection rates [43].

On the other hand, there is currently no information suggesting that omicron symptoms differ from those of other strains [46]. Initially reported infections were among younger people (university students) who tend to have milder disease. Understanding of overall severity of disease associated with omicron will take several days, likely many weeks. A report from the African Medical Association displayed that omicron is seven times more contagious than the delta variant, but the reported cases and deaths in Africa have continued to decline, and the people infected by omicron did not show any serious aggravation in their condition [46].

Many secrets are still present regarding these newly emerging mutants, and most researchers are quite slow and passive in dealing with these potential emergencies. Subsequently, we highly recommended further and profound studies.

5. Vaccination effect against the new variants of COVID-19 and re-infection

Vaccination is considered an effective method applied to prevent several infectious diseases. Vaccines introduced to the body cause the immune system to respond to a foreign substance, producing specific antibodies to neutralize the pathogen. To be effective, vaccines must be able to stimulate the production of antibodies without causing the disease itself. The immune system then retains a memory of the foreign substance, so if the person is exposed to the virus again, the immune system can quickly respond to prevent illness.

Some vaccines, such as those for smallpox and polio, are made by inactivating the virus but leaving enough of it to cause a response from the immune system. Other vaccines use proteins or pieces of proteins that the virus makes to stimulate the immune system. These vaccines are called subunit vaccines.

Recent studies have shown that the current COVID-19 vaccines may not be as effective against the omicron variant as they are against the original strain of the virus. However, the severity of COVID-19 and the need for vaccination are high. Therefore, it is important to continue to research and develop new vaccines that can protect against future variants of the virus.

To the best of our knowledge, there are more than eight vaccines accepted for vaccination against COVID-19 among important categories worldwide focused on their investigations about anti-COVID agents but still the most important questions: Are these currently applied vaccines protect against COVID-19 re-infection, especially in the upcoming expected waves?

The protection afforded by COVID-19 infection is still unknown [59]. A number of recent studies have investigated the re-infection with phylogenetically distinct SARS-CoV-2 variants, but these are still rare [60–62]. Tests on infected patients during the pandemic of SARS in 2003 displayed that antibodies response from infection persists more than two years [63]. In contrast, infection with a mutual seasonal human coronaviruses does not convey permanent defense against re-infection, while re-infection within six months is not common [64]. The new variants that appeared in Brazil and South Africa increase worrying because these spike mutants decreased neutralizing antibodies, suggesting more re-infections and could render vaccines less effective [4]. In contrast, another study revealed that convalescent patients of COVID-19 and vaccinator persons can neutralize the variant of S01Y, and hence this enhanced protection of the current vaccines against B2B/S01Y.V1 strain [37]. This means that vaccines are needed in avoiding re-infections. Others have assessed parallel protection of earlier infection. Among the medical sector workers in the UK, the antibodies percent (91%) strongly reduced the risk of symptomatic re-infection in the next six months [65]. In association with the Moderna vaccine report, prior infection estimated 76% protectiveness in the placebo arm. The numbers were minimal (only one case of re-infection), and the confidence intervals were far-reaching [66]. Lately, a preclinical investigation established that the Ad26.COV2.S vaccine improved the presence of neutralizing antisera and induced defense against the COVID-19 spike variant (G614) [56].

A number of recent studies have investigated the reinfection with phylogenetically distinct SARS-CoV-2 variants, but these are still rare [60–62]. Tests on infected patients during the pandemic of SARS in 2003 displayed that antibodies response from infection persists more than two years [63]. In contrast, infection with a mutual seasonal human coronaviruses does not convey permanent defense against re-infection, while re-infection within six months is not common [64]. The new variants that appeared in Brazil and South Africa increase worrying because these spike mutants decreased neutralizing antibodies, suggesting more re-infections and could render vaccines less effective [4]. In contrast, another study revealed that convalescent patients of COVID-19 and vaccinator persons can neutralize the variant of S01Y, and hence this enhanced protection of the current vaccines against B2B/S01Y.V1 strain [37]. This means that vaccines are needed in avoiding re-infections. Others have assessed parallel protection of earlier infection. Among the medical sector workers in the UK, the antibodies percent (91%) strongly reduced the risk of symptomatic re-infection in the next six months [65]. In association with the Moderna vaccine report, prior infection estimated 76% protectiveness in the placebo arm. The numbers were minimal (only one case of re-infection), and the confidence intervals were far-reaching [66]. Lately, a preclinical investigation established that the Ad26.COV2.S vaccine improved the presence of neutralizing antisera and induced defense against the COVID-19 spike variant (G614) [56].

Prospective risk

* Higher transmission
* Moderate reduction in the neutralization efficiency of sera from convalescent cases or vaccines
* Developed disease severity

| Variants          | B.1.1.7   | B.1.351   | P.1       | B.1.1.529  |
|-------------------|-----------|-----------|-----------|------------|
| First detection   | September | October    | January   | November   |
| Detection country | United kingdom | South Africa | Brazil, Japan | South Africa |
| Mutations Number  | 7         | 9         | 12        | >30        |
| Defined in S protein | P681H, N501Y, A570D, D614G | A701V, L18F, N501Y, D80A, R246L, E484L, K417T, D614G, D215G, | K417T, L18F, R190S, P26S, T20N, D138Y, E484K, D614G, N501Y, V1176F, T1027I, H655Y | |
| Prospective risk  | * Higher transmission | * Reinflection rates are higher | * Reinflection rates are higher | * Reinflection rates are higher |
| Countries with reported cases | 82 | 40 | 19 | 52 |
| Sequenced countries | 64 | 35 | 14 | 87 |
infected cases could still have the virus for months, leading to conflict in actual reinfection rates. This phenomenon appeared because some omicron compared to those who were not immunized. A single dose of any vaccine was associated with a 35% reduction in the risk of hospitalization in symptomatic cases carrying the omicron variant, two doses with a 67% reduction up to 24 weeks after the second dose, and a 51% reduction 25 or more weeks after the second dose, when compared to people who had not received a vaccine [70]. In addition, A total of 6314 omicron patients enhanced the eligibility criteria, out of which 6312 were correlated with one delta case at least out of a total number of 8875 depending on gender, age, and onset date. Twenty-one (0.3%) were hospitalized, whereas zero cases were confirmed with death among matched omicron cases. This situation was compared by 116(2.2%) hospitalizations and seven (0.3%) deaths of reported delta cases [71]. The danger of hospitalization or death was 68% lower in patients with omicron infection compared to delta. After applying vaccination, the risk of hospitalization or death was 54% lower in omicron-infected cases. It is worth mentioning that the death rate is less, however increasing the number of infected persons. This may suggest the danger of the new variant in high transmissibility with increasing infection rate but with no dangerous effect and no more deaths.

On the other hand, a new generation of developed vaccines could be revealed in the future. For example, NVX-CoV2373 announced a new vaccine against SARS-COV-2, which displayed significant protection rates estimated by 95.6% and 85.6% against the original COVID-19 strain and some new variants (UK variant), respectively [72]. However, in South Africa, the same vaccine showed 60% effectiveness (phase II clinical trials) against replication of COVID-19 original strain and 49.4% when applied against the South African variant strain [73]. This is the first vaccine that displayed clinical efficiency against the original strain and the UK and South African [13]. In fact, all vaccines are newly manufactured and have been used quickly to counter the rapid development of the pandemic around the world. Some vaccines have also been developed to counter the genetic changes of the virus. Therefore, research on vaccines' effect and persistence is still few.

It is worthy of mentioning that some reports may overestimate the actual reinfection rates. This phenomenon appeared because some infected cases could still have the virus for months, leading to conflict in distinguishing between existence shedding and reinfection [74]. Protective effects of the currently applied vaccines against the several variants of SARS-COV-2 have been widely discussed; however, there is still a shortage of in-depth investigations and studies with an extensive sample size [13]. Additionally, the antibodies level decreased over time for all vaccines, and many reports which discussed the immunity to other coronaviruses [75–77] confirmed that the immunity started to lose within 1–3 years [61]. This is the first time vaccinating against COVID-19, so the question remains: Will we need an annual dose of the corona vaccines? In addition, Will be mandatory for children in the near future, especially with the spreading of the current fourth wave, which started in different countries?

Moreover, the side effects don't have been observed till now. Subsequently, we recommended further and deep investigation, especially after the new variant of concern, omicron, that witnesses an impressive number of mutations cause increasing in disease severity and significant transmission ability. The situation may get worse and get out of hand, especially when the sense of danger decreased and most people neglect precautionary measures with available vaccines.

6. Conclusion

The SARS-COV-2 pandemic is still causing considerable high mortality rates, introducing a significant burden on healthcare services worldwide and having profound economic and social consequences due to the different strategies implemented to control the virus. Three waves hit the world strongly till now, and more coming waves are highly expected. Although there are many recent reports about the new epidemic, there is still a state of mystery about the nature of this virus and its ability to transform. Vaccines are in a continuous race that highly requires controlling COVID-19 and protecting persons at high risk for complications. Vaccination showed significant activity against some new variants and played a positive role against reinfection. However, in the coming days my revealed further questions, especially with several virus mutations and extra waves as expected.

Author contributions

The manuscript was written through the contributions of all authors. All authors have approved the final version of the manuscript.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

References

[1] Y. Wu, W. Jing, J. Liu, Q. Ma, J. Yuan, Y. Wang, M. Du, M. Liu, Effects of temperature and humidity on the daily new cases and new deaths of COVID-19 in 166 countries, Science of the Total Environment 729 (2020) 1–7.
[2] H. Graichen, What is the difference between the first and the second/third wave of Covid-19? – German perspective, J. Orthop. 24 (2021) A1–A3, https://doi.org/10.1016/j.jor.2021.01.019.
[3] M.F. Chersich, G. Gray, L. Fairlie, Q. Eichbaum, S. Mayhew, B. Allwood, R. English, F. Sorge, S. Lucht, G. Simpson, M.M. Haghigi, M.D. Pham, H. Rees, Covid-19 in Africa: care and protection for frontline healthcare workers, GlobalHealth 16 (2020) 1–6, https://doi.org/10.1186/s12992-020-00574-3.
[4] F.-M.J. Soriani V, New SARS-CoV-2 variants challenge vaccines protection, AIDS Rev. (2021) 2–3, https://doi.org/10.24875/AIDSRev.M21000460.
[5] H.A. Baker, S.A. Safayinia, L.A. Evered, The ‘third wave’: impending cognitive and functional decline in COVID-19 survivors, Br. J. Anaesth. 126 (2021) 44–47, https://doi.org/10.1016/j.bja.2020.09.045.
[6] P.E. Litherland, S. Neumann, E. Tenison, K. Lloyd, T.J. Welsh, J.C.L. Rodrigues, J.P. Higgins, L. Scourfield, H. Christensen, Y.J. Haughton, E.J. Henderson, COVID-19 in older people: a rapid clinical review, Age Ageing 49 (2020) 501–515, https://doi.org/10.1093/ageing/aaaz093.
[7] J. Bransley, C. Heneghan, K.R. Mahanti, J.K. Aronson, Do weather conditions influence the transmission of the coronavirus (SARS-COV-2), Oxford COVID-19, Evid. Serv. 5 (2020), https://www.cebhm.com/do-weather-conditions-influence-the-transmission-of-the-coronavirus-sars-cov-2/.
[8] K.H. Chan, J.S.M. Peiris, S.Y. Lam, L.L.M. Poon, K.Y. Yuen, W.H. Seto, The effects of temperature and relative humidity on the viability of the SARS coronavirus, Aviol. 2011 (2011), https://doi.org/10.1155/2011/734690.
[9] A.W.H. Chin, J.T.S. Chu, M.R.A. Perera, K.P.Y. Hui, H.-L. Yen, M.C.W. Chan, M. Peiris, L.L.M. Poon, Stability of SARS-COV-2 in different environmental conditions, Lancet Microbe. 1 (2020), e10, https://doi.org/10.1016/s2666-5247(20)30003-3.
[10] Y. Ma, Y. Zhao, J. Liu, X. He, B. Wang, S. Fu, J. Yan, J. Niu, Effects of Temperature Variation and Humidity on the Death of COVID-19 in Wuhan, China, 2020, 10.3346/JKMS.2021.36.E124.
[11] J. Xie, Y. Zhu, Association between ambient temperature and COVID-19 infection in 122 cities from China, Sci. Total Environ. 724 (2020), 138201, https://doi.org/10.1016/j.scitotenv.2020.138201.
[12] Y. Yao, J. Pan, Z. Liu, X. Meng, W. Wang, H. Kan, W. Wang, in: No association of Variation and Humidity on the Death of COVID-19 in Wuhan, China, 2020, 10.3346/JKMS.2021.36.E124.
[13] Z. Jia, W. Gong, Will mutations in the spike protein of SARS-CoV-2 lead to the failure of COVID-19 vaccines? J. Korean Med. Sci. 26 (2021) 1–11, https://doi.org/10.3346/jkms.2021.36.e124.
[14] R. Rui, M. Tian, M. Tang, G.T. Ho, Analysis of the Spread of COVID-19 in the USA With a Spatio-Temporal Multivariate Time Series Model 2020, 2021.
[15] M. Cazzella, M. Rajnik, A. Camm, S.C. Deleoh, R.D. Napoli, Features Evaluation and Treatment Coronavirus ( COVID-19 ), 2020.
[16] B. Agerer, S. Immuno, B. Agerer, M. Kohlischke, V. Gudipati, L.F. Montano-gutierrez, A. Popa, J. Genger, L. Endler, D.M. Florian, V. Mühlgraber, M. Granzinger, S.W. Aberle, A. Huxa, L.E. Shaw, A. Lercher, R. Torralba-gombau.
[38] K. Rypdal, F.M. Bianchi, M. Rypdal, Intervention fatigue is the primary cause of V. Soriano, P. Ganado-Pinilla, M. Sanchez-Santos, F. G...V. Soriano, C. de Mendoza, F. Gomez-Gallego, O. Corral, P. Barreiro, Third wave of COVID-19 in Madrid, Spain, Int. J. Infect. Dis. 107 (2021) 212–214, https://doi.org/10.1016/j.ijid.2021.04.017.

R. Rathnasinghe, S. Janardhan, A. Cupic, M. Voysey, S.A.Costa Clemens, S.A. Madhi, L.Y. Weckx, P.M. Folegatti, P.K. Aley, Explained: What We Know so Far About the Omicron Variant of Covid-19, 2021, https://doi.org/10.35772/ghm.2021.01117.

A. Seth, M.S. Hsiang, J.M. Colford, A. Reingold, B.F. Arnold, A. Hubbard, J. Halloran, C. Cummings, R. Holstein, M. Prill, S. Nas, A.B. Perona-an-Oropeza, V.H. Borja-mez-Gallego, P. Barreiro, J. M. Luo, Z. Huang, S. Tu, Y. Zhao, L. Chen, D. Xu, Y. Li, W. Xie, D. Cai, L. Shanghai, F. Fan, X. Xu, D. Wang, Y. Yang, J. Wang, D. Zhang, B.cao, 6-month consequences of COVID-19 in patients discharged from hospital: a cohort study, Lancet 397 (2021) 220–232, https://doi.org/10.1016/S0140-6736(20)31266-6.

M. Voysey, S.A.Costa Clemens, S.A. Madhi, L.Y. Weckx, P.M. Folegatti, P.K. Aley, J. Anesi, The advisory committee on immunization practices monitor the fraction of people ever infected with COVID-19: an application to the United States, JAMA - J. Am. Med. Assoc. 325 (2021) e154-155, https://doi.org/10.1001/jama.2021.3331.

C.H. Sudre, B. Murray, T. Varsavsky, M.S. Graham, R.S. Penfold, R.C. Bowyer, J. C. Poveda, K. Klar, M. Antonelli, L. Canaviri, E. Motelnic, M. Modat, M. Jorge Cardoso, A. May, S. Ganesh, R. Davies, L.H. Nguyen, D.A. Drew, C.M. Astley, A. D. Joshi, J. Merino, N. Tsertelis, T. Fall, M.G. Eln, Duncan C. Menni, P.M. Williams, Frank’s, A. Chan, J. Wolf, S. Oursoin, T. Sertor, C.J. Stevens, Attributes and predictors of long COVID. Nat. Med. 27 (2021) 626–631, https://doi.org/10.1038/s41591-021-01292-y.

C. Huang, L. Wang, Y. Li, L. Ren, X. Gu, L. Kang, L. Guo, M. Liu, X. Zhou, J. Lao, Z. Huang, S. Tu, Y. Zhao, L. Chen, D. Xu, Y. Li, C. Li, L. Peng, Y. Li, W. Xie, D. Cai, L. Shanghai, F. Fan, X. Xu, D. Wang, Y. Yang, J. Wang, D. Zhang, B.cao, 6-month consequences of COVID-19 in patients discharged from hospital: a cohort study, Lancet 397 (2021) 220–232, https://doi.org/10.1016/S0140-6736(20)31266-6.

M.P. Guallar, R. Meiri, C. Martínez-Romero, L.C.F. Mulder, J. Luo, Z. Huang, S. Tu, Y. Zhao, L. Chen, D. Xu, Y. Li, W. Xie, D. Cai, L. Shanghai, F. Fan, X. Xu, D. Wang, Y. Yang, J. Wang, D. Zhang, B.cao, 6-month consequences of COVID-19 in patients discharged from hospital: a cohort study, Lancet 397 (2021) 220–232, https://doi.org/10.1016/S0140-6736(20)31266-6.

M. Voysey, S.A.Costa Clemens, S.A. Madhi, L.Y. Weckx, P.M. Folegatti, P.K. Aley, J. Anesi, The advisory committee on immunization practices monitor the fraction of people ever infected with COVID-19: an application to the United States, JAMA - J. Am. Med. Assoc. 325 (2021) e154-155, https://doi.org/10.1001/jama.2021.3331.

A. Meekins, C.D. McDowell, V. Balaraman, J.A. Richt, B.G. De Geest, L. Miorin, F. Krammer, V. Simon, A. García-Sastre, M. Schotsaert, The N501Y mutation in SARS-CoV-2 vaccines in nonhuman primates and humans, Sci. Adv. 7 (2021) eabe8065, (n.d.).
[61] R.L. Tillett, J.R. Sevinsky, P.D. Hartley, H. Kerwin, N. Crawford, A. Gorzalski, C. Lavender, S.C. Verma, C.C. Rossetto, D. Jackson, M.J. Farrell, S. Van Hooser, M. Pandori, Genomic evidence for SARS-coronavirus-2 strain confirmed by whole genome sequencing: One-Minute Summary PHO Reviewer’s Comments, (n.d.) 1–3.

[62] A.T. KK-WH IF-N 1-JP-C, AW H C-W M-T Ar., Review of ‘COVID-19 re-infection by a phylogenetically distinct SARS-coronavirus-2 strain confirmed by whole genome sequencing’ One-Minute Summary PHO Reviewer’s Comments, (n.d.) 1–3.

[63] L.P. Wu, N.C. Wang, Y.H. Chang, Y.T. Na, L.Y. Zhang, L. Zheng, T. Lan, L.F. Wang, G.D. Liang, Duration of antibody responses after severe acute respiratory syndrome, Emerg. Infect. Dis. 13 (2007) 1562–1564, https://doi.org/10.3201/eid1310.070576,

[64] A.W.D. Edridge, J. Kaczorowska, A.C.R. Hoste, M. Bakker, M. Klein, K. Loens, M. L.R. Baden, H.M. El Sahly, B. Essink, K. Kotloff, S. Frey, R. Novak, D. Diemert, S. J.E.M. van der Lubbe, S.K. Rosendahl Huber, A. Vijayan, L. Dekking, E. van Huizen, K.R.W. Emary, T. Golubchik, P.K. Aley, C.V. Ariani, B. Angus, S. Bibi, B. Blane, D.A.T. Cummings, A systematic review of antibody mediated immunity to SARS-CoV-2 infection in, Health Care Workers (2021) 533–540, https://doi.org/10.1056/NEJMa2034545,

[65] L.R. Baden, H.M. El Sahly, B. Esiinik, K. Korioff, S. Frey, R. Novak, D. Diemert, S. A. Specter, N. Roupahaed, C.B. Creech, J. McGettigan, S. Kheta, N. Segall, J. Solid, A. Bronz, C. Fierro, H. Schwartz, K. Neuzil, L. Corey, P. Gilbert, H. Janes, D. Pollmann, M. Marovich, J. Macusa, L. Polakowski, J. Ledgerwood, B.S. Graham, H. Bennett, R. Pajon, C. Knightly, B. Leav, W. Deng, H. Zhou, S. Han, M. Ivarsson, J. Miller, T. Zaks, Efficacy and safety of the mRNA-1273 SARS-CoV-2 vaccine, N. Engl. J. Med. 384 (2021) 403–416, https://doi.org/10.1056/NEJMoa2035389,

[66] J.E.M. van der Lubbe, B.S. Knudsen, M. Borrow, H. Ghebregziabher, B. Leav, W. Deng, H. Zhou, S. Han, M. Ivarsson, J. Miller, T. Zaks, Efficacy and safety of the mRNA-1273 SARS-CoV-2 vaccine, N. Engl. J. Med. 384 (2021) 403–416, https://doi.org/10.1056/NEJMoa2035389,

[67] J.E.M. van der Lubbe, S.K. Rosendahl Huber, A. Vijayan, L. Dekking, E. van Huizen, J. Vreugdenhil, Y. Choi, M.R.M. Baert, K. Feddes-de Boer, A. Izquierdo Gil, M. van Heerden, T.J. Dalebout, S.K. Myeni, M. Kikkert, E.J. Snijder, L. de Waal, K. J. Vreugdenhil, J.T.B.M. Tolboom, J. Serroyen, L. Muchene, L. van der Fits, L. Rutten, M. Pandori, Genomic evidence for reinfection with SARS-CoV-2: a case study, Lancet Infect. Dis. 21 (2021) 52–58, https://doi.org/10.1016/S1473-3099(20)30764-7,

[68] A.T. KK-WH IF-N 1-JP-C, AW H C-W M-T Ar., Review of ‘COVID-19 re-infection by a phylogenetically distinct SARS-coronavirus-2 strain confirmed by whole genome sequencing’ One-Minute Summary PHO Reviewer’s Comments, (n.d.) 1–3.

[69] F. Jebbink, A. Matser, C.M. Kinsella, P. Rueda, M. Ieven, H. Goossens, M. Prins, P. Saure, M. Dejls, L. van der Hoek, Seasonal coronavirus protective immunity is short-lasting, Nat. Med. 26 (2020) 1691–1693, https://doi.org/10.1038/s41591-020-1083-l,

[70] Public Health England, in: SARS-CoV-2 Variants of Concern and Variants Under Investigation in England 01, Sage, 2021, pp. 1–50.

[71] A.C. Uolloa, S.A. Buchan, N. Danceman, K.A. Brown, P.H. Ontario, C. Uolloa, K. A. Brown, Early estimates of SARS-CoV-2 omicron variant severity based on a matched cohort study, complete word count (excluding figures / captions): 634 corresponding author 116 (2021) 1–6.

[72] E. Callaway, S. Mallapaty, Novavax covid vaccine protects people against variants, Nature 590 (2021) 17,

[73] V. Shinde, S. Bhihka, Z. Hossain, M. Archary, Q. Ibhorat, L. Fairlie, U. Lalloo, M.S. L. Masilela, D. Moodley, S. Hanley, L. Fouche, C. Louw, M. Tameris, N. Singh, A. Goga, K. Dheca, C. Grobbelaar, G. Kruger, N. Carrim-Ganey, V. Baillie, T. de Oliveira, A. Lombard Koen, J.J. Lombarda, R. Mqgubisa, A.E. Bhhorat, G. Benadé, N. Lalloo, A. Pitts, P. L. Vollgraaff, A. Luabeysa, A. Isemil, F.G. Petrick, A. Oommen-Jose, S. Foulkes, K. Ahmed, A. Thombrayil, L. Fries, S. Cloney-Clark, M. Zhu, C. Bennett, G. Faust, J.S. Plested, A. Robertson, S. Neal, I. Cho, G. M. Glenn, F. Dubovsky, S.A. Madhi, Efficacy of NVX-CoV2373 Covid-19 vaccine against the B.1.521 variant, N. Engl. J. Med. 384 (2021) 1899–1909, https://doi.org/10.1056/NEJMoa2103055,

[74] M.M.S. BS, A.J.R.M. MBA, M.B.R.M. MPH, in: Reinfection Rates among Patients who Previously Tested Positive for COVID-19: a Retrospective Cohort Study, 2021, pp. 2–16.

[75] K.A. Callow, H.F. Parry, M. Sergeant, D.A. Tyrell, The time course of the immune response to experimental coronavirus infection of man, Epidemiol. Infect. 105 (1990) 435–446, https://doi.org/10.1017/S0950268800048019,

[76] S.C. Chang, J.T. Wang, L.M. Huang, Y.C. Chen, C.T. Fang, W.H. Sheng, J.L. Wang, C.J. Yu, P.C. Yang, Longitudinal analysis of severe acute respiratory syndrome (SARS)-coronavirus-specific antibody in SARS patients, Clin. Diagn. Lab. Immunol. 12 (2005) 1455–1457, https://doi.org/10.1128/CDLI.12.12.1455-1457.2005,

[77] A.T. Huang, B. Garcia-Carreras, M.D.T. Hitchings, B. Yang, L.C. Katzelnick, S. C. Chang, J.T. Wang, L.M. Huang, Y.C. Chen, C.T. Fang, W.H. Sheng, J.L. Wang, S.C. Chang, J.T. Wang, L.M. Huang, Y.C. Chen, C.T. Fang, W.H. Sheng, J.L. Wang, C.J. Yu, P.C. Yang, Longitudinal analysis of severe acute respiratory syndrome (SARS)-coronavirus-specific antibody in SARS patients, Clin. Diagn. Lab. Immunol. 12 (2005) 1455–1457, https://doi.org/10.1128/CDLI.12.12.1455-1457.2005,

[78] A.T. Huang, B. Garcia-Carreras, M.D.T. Hitchings, B. Yang, L.C. Katzelnick, S. C. Chang, J.T. Wang, L.M. Huang, Y.C. Chen, C.T. Fang, W.H. Sheng, J.L. Wang, S.C. Chang, J.T. Wang, L.M. Huang, Y.C. Chen, C.T. Fang, W.H. Sheng, J.L. Wang, C.J. Yu, P.C. Yang, Longitudinal analysis of severe acute respiratory syndrome (SARS)-coronavirus-specific antibody in SARS patients, Clin. Diagn. Lab. Immunol. 12 (2005) 1455–1457, https://doi.org/10.1128/CDLI.12.12.1455-1457.2005,