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Review

COVID-19 vaccine: A recent update in pipeline vaccines, their design and development strategies

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A R T I C L E I N F O

A B S T R A C T

Coronavirus Disease 2019 named as COVID-19 imposing a huge burden on public health as well as global economies, is caused by a new strain of betacoronavirus named as SARS-CoV-2. The high transmission rate of the virus has resulted in current havoc which highlights the need for a fast and effective approach either to prevent or treat the deadly infection. Development of vaccines can be the most prominent approach to prevent the virus to cause COVID-19 and hence will play a vital role in controlling the spread of the virus and reducing mortality. The virus uses its spike proteins for entering into the host by interacting with a specific receptor called angiotensin converting enzyme-2 (ACE2) present on the surface of alveolar cells in the lungs. Researchers all over the world are targeting the spike protein for the development of potential vaccines. Here, we discuss the immunopathological basis of vaccine designing that can be approached for vaccine development against SARS-CoV-2 infection and different platforms that are being used for vaccine development. We believe this review will increase our understanding of the vaccine designing against SARS-CoV-2 and subsequently contribute to the control of SARS-CoV-2 infections. Also, it gives an insight into the current status of vaccine development and associated outcomes reported at different phases of trial.

1. Introduction

By the end of year 2019, Wuhan city of China had witnessed several cases of pneumonia like conditions, which later on found to be caused by 2019-novel coronavirus (2019-nCoV) (Zhou et al., 2020b). Later, on February 11, 2020, the novel coronavirus was termed as severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) by the international virus classification commission and on the same day the disease caused by SARS-CoV-2 was termed as Coronavirus Disease 2019 (COVID-19) by World Health Organization (WHO) (Li et al., 2020a,b). WHO had declared COVID-19 as a global health emergency on January 30, 2020 and characterized it pandemic on 11 March 2020. The case fatality ratio (CFR, defined as the proportion of individuals diagnosed with a disease who die from that disease) was reported as 1–3% and it was shown to be higher in elderly patients especially men (Spychalski et al., 2020). CFR is therefore a measure of severity of infection among detected cases, which is found to be varying among the WHO defined region (summarized in Table 1). The probable reason could be the region specific immune response of individuals towards the infection. Since the southeast Asian countries are well immune to SARS and MERS infection, the fatality

ratio is comparatively lesser than those in WHO defined American and European region. This potentially fatal virus is known to transmit quickly and have spread across 216 countries and territories affecting more than 45 million people worldwide (WHO, 2020). This came within the 11 months after the first case have reported in Wuhan. Given the severity of the disease, there is an urgent need of therapeutics or vaccines to treat or prevent the spread of SARS-CoV-2.

Currently, no potential drugs are available to treat COVID-19. Only strategies followed worldwide to combat COVID-19 is to relieve the patient's symptoms. There are many therapeutic approaches being tried all over the world to tackle SARS-CoV-2 infections. One approach is to understand the structure, life cycle and pathogenesis of the virus, which ultimately facilitates a deeper understanding of various targets to be approached for devising a specific drug for SARS-CoV-2. Another approach like repurposing of existing drugs, such as those for influenza, hepatitis B, hepatitis C, and filoviruses, was considered to allow more rapid development (Li and De Clercq, 2020). Various therapeutic treatments like antiviral drugs (nucleotide analogue) (Cao et al., 2020; Holshue et al., 2020; Wang et al., 2020), antimalarial drugs (those inhibits viral cell entry and its replication) (Gao et al., 2020; Gautret et al.,

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Table 1
Case fatality rate across the world as per WHO.

WHO Region	Confirmed Cases	Deaths	CFR
Americas	19,737,794	625,973	3.17%
Europe	9,664,042	270,972	2.8%
South East Asia	8,969,707	140,827	1.5%
Eastern Mediterranean	2,955,552	75,133	2.5%
Africa	1,298,315	29,277	2.2%
Western Pacific	715,300	15,314	2.1%

2020), immunomodulators (Bian et al., 2020) and cell and plasma-based therapy (Leng et al., 2020) are currently being used worldwide. There are number of clinical trials ongoing all over the world in search of a specific treatment for COVID-19. Despite the various approaches there are number of cases where patients once declared clinically recovered from COVID-19 have suffered reinfection (An et al., 2020; Xing et al., 2020; Yuan et al., 2020) and these shortcomings of current approaches highlight the necessity of vaccines against SARS-CoV-2. For the development of a vaccine against SARS-CoV-2, numbers of approaches are being tried all over the world like targeting S protein of the virus and generating antibodies against this. Besides this various other vaccines are designed against the virus, including RNA and DNA based vaccines and these are described further in details.

Scientists and researchers all over the world are trying to understand this novel virus, SARS-CoV-2, to develop specific treatments for controlling and preventing this potentially fatal disease. Immune system plays pivotal role in the pathogenesis of SARS-CoV-2 infection, and to develop understanding of the immune response and the underlying mechanism is very important in developing an effective vaccine. In this review, we discuss different targets that can be approached for vaccine design and development against SARS-CoV-2 infections. Also, we discuss different platforms that are currently being used all over the world for the development of vaccine candidates.

2. Key features of human coronaviruses

Coronaviruses are positive-sense single-stranded viruses with RNA as genetic material and they are named so because of their crown like shape obtained by the presence of surface glycoproteins (Cui et al., 2019). These viruses mainly cause infection in birds and mammals. In humans, these generally cause mild respiratory infections with common cold like symptoms. However, in the past two decades, three of the human coronavirus of zoonotic origin including SARS-CoV, MERS-CoV and SARS-CoV-2 have resulted in lethal endemics in 2002, 2012 and 2019 respectively. All of these three viruses belong to the genus Beta coronavirus within the family Coronaviridae (Ahmed et al., 2020). The comparative characteristics are tabulated in Table 2. Studies have shown that SARS-CoV-2 is most likely originated in bats but some intermediate host might be there responsible for amplifying the virus (Zhou et al., 2020b). Another study has suggested pangolins

Table 2
Comparative characteristics of zoonotic beta cornoviruses lethal to humans.

Characteristics	SARS-CoV-2	MERS-CoV	SARS-CoV
Date first detected	December 2019	June 2012	November 2002
Potential Reservoir Host	Bat	Dromedary Camels (WHO)	Bat
Possible Intermediate host	Pangolin	Camel	Palm civets
Target cellular receptor	Angiotensin converting enzyme 2 (ACE2)	Dipeptidyl peptidase 4 (DPP4)	Angiotensin converting enzyme 2 (ACE2)
Incubation Period	2–14 days	2–10 days	2–10 days
Speed of spread	High	Low	Moderate

as a potential amplifying host for SARS-CoV-2 (Zhang et al., 2020a, 2020b). The similarity between existing SARS-CoV and novel SARS-CoV-2 helped in revealing the entry mechanism of virus into host. Studies have shown that just like SARS-CoV, SARS-CoV-2 used angiotensin-converting enzyme-2 (ACE2) receptor for entering into hosts ranging from bats to humans including all the animals that express ACE2 receptor (Letko et al., 2020; Wan et al., 2020). The swift genomic sequencing of SARS-CoV-2 remains helpful in determining its genetic similarity with MERS-CoV and SARS-CoV (Lu et al., 2020). This has been helpful in identifying the genomic regions of interest that can be targeted therapeutically for prevention as well as treatment.

3. SARS-CoV-2 structure and targets for vaccine development

SARS-CoV-2 is a spherical shaped virus with genome size of 30 kb. The two third of the genome of SARS-CoV-2 shows expression of the replicase and the other part of genome codes for structural and accessory proteins (Adnan et al., 2020). The four types of structural proteins are spike protein (S), envelope protein (E), membrane glycoprotein (M) and nucleocapsid protein (N). The genetic sequence of SARS-CoV-2 was published on 11 January 2020 by Wu et al. (2020c). Studies have shown approximately 80% similarity in the RNA genome of SARS-CoV-2 and SARS-CoV yet notable variations are reported in the genome of both these viruses as well as with MERS-CoV genome as shown in Fig. 1A. These differences include the absence of 8a protein and difference in the number of amino acids that codes for 8b and 3c protein in SARS-CoV-2 (Wu et al., 2020a) (Fig. 1A). It is also reported that genome of SARS-CoV-2 has undergone genetic recombination with SARS-related CoV and it might be a combination of SARS-CoV of bat and unknown beta-CoV (Zhang et al., 2020). This new strain of SARS-CoV-2 can easily recognize the evolving human ACE2 receptor, and therefore its S protein possesses efficient transduction efficiency resulting in the fast transmission of SARS-CoV-2 in humans (Walls et al., 2020). The S protein comprises of two functional subunits S1 and S2 subunits, S1 subunit being responsible for interaction with the host's ACE2 receptor through Receptor Binding Domain (RBD) and S2 subunit responsible for the fusion of viral and cellular membranes. The S protein also contains a functional polybasic (furin) cleavage site at the S1–S2 subunit boundary which led to the acquisition of three O-linked glycans around the site (Fig. 1A). Both the polybasic cleavage site and the three adjacent predicted O-linked glycans are unique to SARS-CoV-2 and were not previously seen in lineage betacoronaviruses (Andersen et al., 2020; Hu et al., 2020).

The continuous evolution of the novel coronavirus poses its ability to quickly transmit among population (Zhang et al., 2020). The similarity between the two viruses can be helpful in vaccine development against SARS-CoV-2 with the help of previously researched protective immune response against SARS-CoV. Studies have demonstrated that S protein is a prime target for the designing and developing vaccines (Ahmed et al., 2020). It is responsible for the attachment of virus to the host's cell surface receptor (most likely ACE2). Just as in SARS-CoV, antibodies targeting the S protein of SARS-CoV-2 can interfere with the virus resulting in neutralization of the infection caused by the virus (Liu et al., 2017). Hence, many vaccines developing platforms containing target antigen commonly target the S protein (Fig. 1B).

4. COVID-19 immunopathology and vaccine designing

The immunopathology of COVID-19 explains the reaction of immune system against novel coronavirus infection. SARS-CoV-2 attacks hosts' immune system leading to an uncontrolled inflammatory responses, lymphocytpenia, high cytokine levels and increased antibodies. The most prominent immuno invasion processes observed in patients were: Depletion of lymphocytes, increased neutrophils, cytokine storm and antibody dependent enhancement (Yang et al., 2020a). The SARS-CoV-2
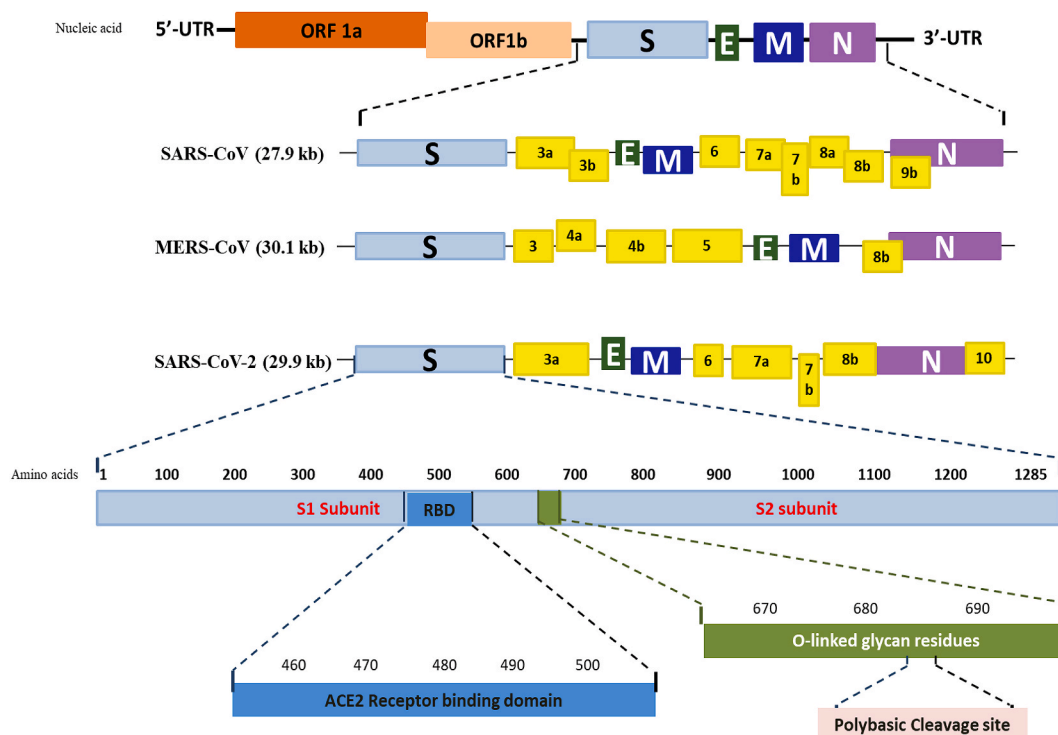
Fig. 1a. Schematic diagram of genomic organization or open-reading frames (ORFs) of SARS-CoV, MERS-CoV and SARS-CoV-2. Variations in accessory proteins (yellow) in all the three strains of betacoronvirus organized within structural proteins, Spike (light blue, S), envelope (green, E), membrane (dark blue, M), nucleocapsid (purple, N) and non-structural proteins expressed in ORF 1a (brown) and ORF 1b (peach) are indicated. Untranslated regions at the N and C-terminals are represented respectively as 5′-UTR and 3′-UTR; kb indicates kilo base pairs. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)
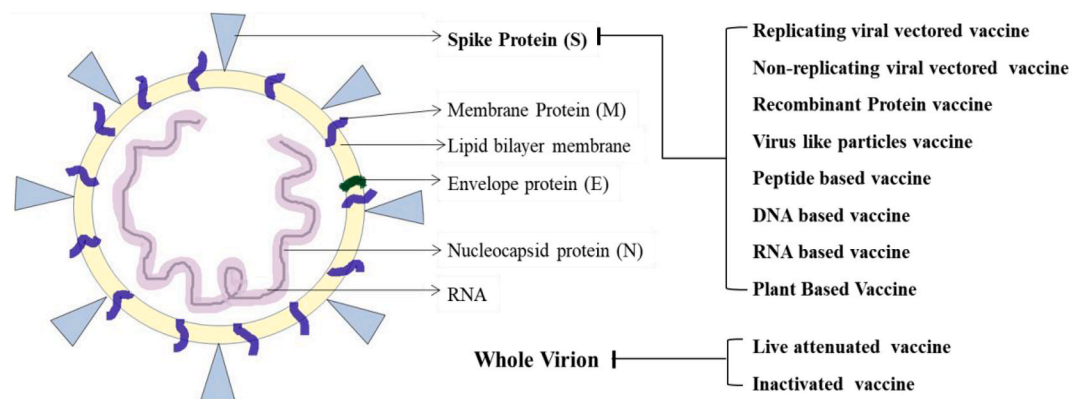

Fig. 1b. Schematic structure of SARS-CoV-2 with its key structural proteins as target antigens for various vaccines production platforms. S protein is the major target antigen for most of the platforms except the conventional ones (Live attenuated and Inactivated vaccines) where the whole virion or the subunit of it is used to develop vaccines.

virion attacks ACE2 receptors on type 2 alveolar cells and release the viral genome (RNA) into the cytoplasm of host cell, the RNA further gets translated into structural proteins and then begins to replicate (Perlman and Netland, 2009). The viral particles subsequently undergo transformation in the endoplasmic reticulum-Golgi intermediate compartment and released entrapped into vesicles, which on fusion with plasma membrane release the viral particles (De Wit et al., 2016). Simultaneously, while virus enters the cells, a part of it as antigen is presented on the surface of antigen presentation cells (APC), which is crucial for the body's immune response. Antigen presentation on B and T cells ultimately leads to the production of the typical pattern of IgM and IgG antibodies. Researchers have found that these IgM antibodies lasts until

week 12, while the IgG antibodies remain long lasting and can provide prolonged protection (Li et al., 2003). These type of virus-specific neutralizing IgG antibodies play major role by regulating the infection at later phase and prevents reinfection in the future (De Wit et al., 2016). In general, T cells are the major immune cells those fight against viral infections in body either by production of virus specific antibodies (CD4+ helper type T cells mediated response) or by killing the virus-infected cells (CD8+ cytotoxic type T cells mediated response) (Mubarak et al., 2019). However, the report of COVID-19 patients shows marked reduction in CD4+ and CD8+ T cells population, whereas the high proportions of HLA-DR (CD4 3.47%) and CD38 (CD8 39.4%) double positive fractions indicates the hyperactivated status of T cells

imparting severe immune injury (Xu et al., 2020). There could be the several possible mechanisms involved in the depletion and dysfunctioning of T lymphocytes induced by SARS-CoV-2 such as, (1) SARS-CoV-2 can directly infect T cells and macrophages by binding to the surface markers (CD 26 and CD 147) (García, 2020) or the ACE-2 receptors present on their surface (Dandekar and Perlman, 2005) as well as SARS-CoV-2 induce spleen atrophy and lymph node necrosis thereby destroying lymphatic organs (Cao, 2020; Li et al., 2020a,b; Tan et al., 2020); (2) another mechanism is cytokine storm induced by SARS-CoV-2, the increased level of cytokines like TNFα, IL-6, and IL-10 is inversely correlated with the decreased T cell population (Moon, 2020). The lymphocytopenia occurring in the patients further explains the increased neutrophil count in patients, since the depletion of T cells make patients more susceptible to microbial infections and thereby promoting the recruitment and activation of neutrophils in patients' blood (Deshmukh et al., 2014). The clinical outcomes and associated manifestations are briefly summarized in Table 3.

Apart from lymphocytopenia, cytokine storm is another major clinical outcome observed in patients with severe COVID-19. SARS-CoV-2 after invading the host, activates host's immune system first inducing innate immune response and then subsequently generating the adaptive immunity. The innate response is marked by the production of cytokines specifically Type-I and –III antiviral Interferons and different chemokines (Hur, 2019). Chemokines induce infiltration of more innate cells like monocytes, Natural Killer cells, dendritic cells, polymorphonuclear leukocytes, which produces more chemokines (García, 2020). On the other hand direct activation of T cells by the SARS-CoV-2 infection leads to the activation of pathogenic T helper 1 (Th1) cells that secrete GM-CSF and produce inflammatory cytokine IL-17 which recruits macrophages and neutrophils to the site of infection and stimulates IL-1β and IL-6 cytokine cascades, where IL-6 plays the major role in development of cytokine storm in COVID-19 patients (Wu and Yang, 2020; Zhou et al., 2020c). T cells ultimately led to the activation of B lymphocytes which induce humoral response by producing neutralizing antibodies. COVID-19 patients were found to have poor outcome even after antibody upregulation, this can be explained by the phenomenon Antibody Dependent Enhancement (ADE) (Ulrich et al., 2020). Various infections like Ebola, Dengue, MERS have shown the antibody dependent enhancement of viral infection where it was observed that the viral entry and replication is being enhanced by sub-neutralizing antibodies (Cao, 2020; Katzelnick et al., 2017; Roncati et al., 2020). However, the ADE mediated inflammatory response and its association with disease severity and progression requires further investigation (Yang et al., 2020a). The interconnection between all these processes collectively giving rise to lung injury and disease progression is briefly schematized in Fig. 2.

Studies on SARS-CoV patients have shown the presence of memory T cells after the four years from recovery and those memory T cells remain able to respond against the S protein of SARS-Cov (Fan et al., 2009; Tang et al., 2011). The similar results were reported in MERS coronavirus infection in mice (Zhao et al., 2014). The aftermath, strongly indicates the importance of T cells in controlling the previously existing SARS and MERS coronavirus infections; hence there is high possibility that it can also provide protection against novel coronavirus, SARS-CoV-2 infection. The epitopes identified for SARS and MERS coronaviruses mostly concentrate on structural proteins especially S protein, hence mapping these epitopes against SARS-CoV-2 can be beneficial in designing the vaccine against all three human coronaviruses in the future (Mubarak et al., 2019).

4.1. Role of bioinformatics in vaccine designing-

The remarkable scientific and technological advancements since last century have played a major role in the improvement of vaccines' potency as well as in speeding up their production. The fusion of computational technologies with the recombinant DNA technology, the possibility of accelerated and massive sequencing of complete genomes and the fast growth of biological and genomic information in database banks have made this possible (He et al., 2010; María et al., 2017). Reveres vaccinology, where one could identify the structure of the bacteria, virus, cancer cell or any pathogen or allergen capable of inducing immune response using the bioinformatics tools, possess various advantages over traditional vaccine production methodology like reduced time and cost of vaccine development (Rappuoli, 2000; Seib et al., 2012). A wide variety of softwares are available providing various functions ranging from genomic sequencing of an organism to finding out the possible immunogenic response associated with different proteins, thereby easing the vaccine development process. Also there are websites that with the help of 3D structure of protein, can predict conformational epitopes for B cells (María et al., 2017). NIAID Virus Pathogen Database and Analysis Resource (ViPR) and Immune Epitope Database (IEDB) helped in comparing epitopes of SARS COV 1 and SARS COV 2 which further found helpful in epitope mapping and targeting SARS-CoV-2 spike protein in order to develop various vaccines against the infection (Ahmed et al., 2020).

5. SARS-CoV-2 vaccine production platforms

The rapid transmission of SAR-CoV-2 infection across the world ignited extensive efforts toward the development of COVID-19 vaccines that can be used globally for ending the pandemic. The full length spike glycoprotein or RBD of virion are able prevent host and virus interaction

Table 3
Symptoms and Clinical outcomes of SARS-CoV-2 induced immunopathology in humans.

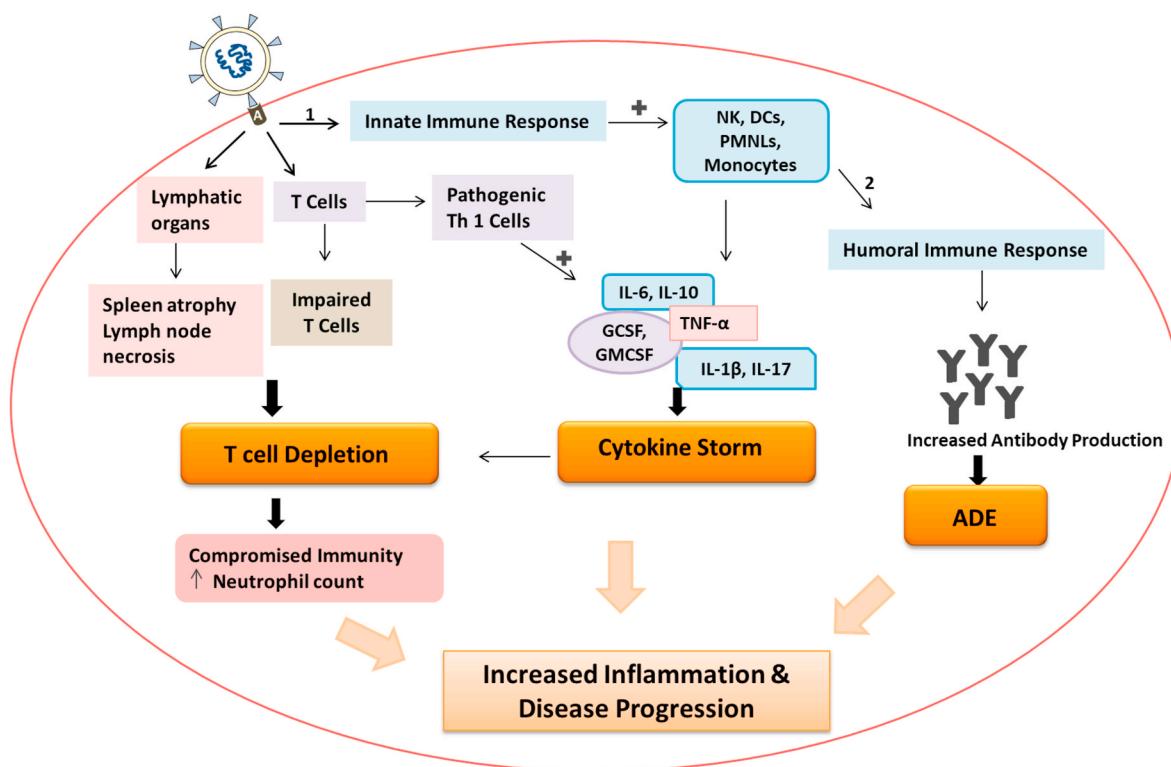| Organ System	Clinical Outcomes	Clinical Manifestations	References
Vascular System	Cytokine storm	Elevated IL-6, TNFα, IL-1β	(Huang et al., 2020; Rodrigues et al., 2020; Zhou et al., 2020a,b,c)
	Lymphocyte count	Elevated T helper 17 cells (TH17), plasma cells, CD8+ T cell activity, and decreased regulatory T cells	(Huang et al., 2020; Qin et al., 2020)
	Thrombocytopenia associated mortality	Reduced platelet to lymphocyte ratio	Lippi et al. (2020)
	ARDS	Elevated levels of ferritin	(Wu et al., 2020a,b,c; Zhou et al., 2020a,b,c)
	Vasculitis and Vascular Dysfunction	Elevated VEGF, IL-10, IL-8	(Becker, 2020; Huang et al., 2020; Zhou et al., 2020a,b,c)
Lungs	Pneumonia with ARDS and dyspnea	Elevated IL-6, TNFα, IL-1β, IL-10, IL-8	(Huang et al., 2020; Xu et al., 2020)
Kidneys	Proteinuria and hematuria	Elevated urea and creatine levels	(Cheng et al., 2020; Su et al., 2020; Yang et al., 2020a,b)
Liver	Steatosis and abnormal Liver function	Elevated AST, ALT, CRP, albumin levels	(Ling, 2020; Zhou et al., 2020a,b,c)
Heart	Acute myocardial injury and chronic CVS damage	Elevated CK and LDH levels	(Huang et al., 2020; Yang et al., 2020a,b)
Intestine	Microbial infection, diarrhea, severe acute ulcerative colitis	Reduced T cell and NK cell count (Lymphopenia)	(Carvalho et al., 2020; D'Amico et al., 2020; Yang et al., 2020a,b)
Brain	Encephalopathy, headache, ischemic stroke	Elevated CRP, D-dimer, ferritin levels	(Beyrouti et al., 2020; Huang et al., 2020)

Fig. 2. **Schematized representation of immunopathogenesis of SARS-CoV-2.** SARS-CoV-2 acting on ACE-2 receptor (A) giving rise to the cascade of pathological mechanisms including, Activating innate line of defense (1) and subsequently giving rise to Humoral response (2), leading to Antibody dependent enhancement (ADE), T cell depletion, Lymphatic organs damage and Cytokine storm.

by inducing neutralizing antibodies and hence is considered as most important vaccine target antigen (Prompetchara and Chutitorn Ketloy, 2020). Most of the COVID-19 vaccine development projects ongoing all over the world are using S protein as target antigen (Table 4).

In the last decade, there is the marked evolution of vaccine platforms including the development of nucleic acid based vaccine candidates, vectored vaccines and recombinant protein vaccines. SARS-CoV-2 vaccine candidates currently being developed worldwide and under evaluation are based on different vaccine platforms briefly detailed in Fig. 1B. Some of the platforms are already in use for different vaccines and are licensed for use by different regulatory bodies while some of them are still in trials. However, the understanding of these in the field of oncology has encouraged researchers to use next-generation approaches in order to increase the speed of development and manufacture.

5.1. Replicating and non replicating vectors

Based on their capability to replicate in the host cell, these can be replicating and non replicating recombinant viral vectors. These are evaluated as delivery of viral genome encoding the gene of interest. The longevity of the immune response generated by vaccine depends on the type of viral vector used. Most commonly used viral vector is Adenoviral (Ad) vector. The major advantage of this platform is the capability to induce both humoral and cellular immunity. Despite the complex production of viral vector based vaccines, these are known to induce strong immunological response (Geiben-Lynn et al., 2008; Rollier et al., 2011). However, sometimes these are not capable of inducing immunogenicity due to the presence of preexistent immunity (Sekaly, 2008). A recombinant replicating vaccine, Ervebo is licensed in the European Union as ebolavirus vaccine (Marzi et al., 2011). The adenoviral vector Ad5 being used for COVID-19 vaccine development is cost effective approach and

already been used for Ebola virus. One of the limitations of Ad5 vectors is their association with high prevalence in the human population and therefore an additional trial using chimpanzee derived adenoviruses (ChAd) is being conducted to combat preexisting immunity. Currently, there are 9 COVID-19 vaccine candidates developed using non replicating vector platform in clinical evaluation and 19 candidates in preclinical evaluation stage. Four COVID-19 vaccine candidates developed using replicating vector platform is in clinical evaluation stage and 18 candidates are in preclinical evaluation stage (World Health Organization, 2020).

5.2. Virus-like particles (VLPs)

These are protein multimers mimicking the structure of real virus but lacking genetic material and hence are non-infectious in nature. These can self assemble with more than one type of protein forming protein chimeras known as cVLPs (Roldão et al., 2010). VLPs act by stimulating antigen presenting cells mediated activation of B- and T-cell immune responses. These are also involved in CD8$^+$ cytotoxic T cell mediated killing of pathogenic cells. The immune system recognizes VLPs in the same way as it recognize original virus and thereby induce immune responses (Hemann et al., 2013). VLP formulations because of their poor immunogenicity require adjuvants in most of the cases. VLP-based vaccines are well established platform for prophylactic use. These are less time taking and production cost depends upon the expression system used which is comparatively low for bacterial system than the mammalian expression system. The licensed vaccines based on this platform are currently in use for human papillomavirus (HPV) (Zhao et al., 2013). Currently, there are two COVID-19 vaccine candidates developed as VLPs in clinical evaluation and 15 COVID-19 vaccine candidates in preclinical evaluation stage developed (World Health Organization, 2020).

Table 4

List of COVID-19 Vaccine candidates under clinical evaluation stage.

Vaccine Platform	Vaccine Name	Vaccine characteristics/Immunogen	Stage of Development[a]
Non Replicating Viral vector	ChAdOx1-S nCoV-19 vaccine	Adenovirus Type 5 vector that expresses S protein	Phase 3 NCT04516746 ISRCTN89951424 NCT04540393 CTRI/2020/08/027170
	Ad-5 vectored COVID-19 vaccine	Weak recombinant adenovirus carrying S protein expression	Phase 3 NCT04526990 NCT04540419
	Gam-COVID-Vac	Adeno-based vectored combined (rAd26-S + rAd5-S) expressing S protein	Phase 3 NCT04530396 NCT04564716
	Ad26.COV2.S	Adenovirus vectored vaccine	Phase 3 NCT04505722
	GRAd-COV2	Replication defective Gorilla Adenovirus that encodes for SARS-COV-2 Spike protein	Phase 1 NCT04528641
	hAd5-S-Fusion + N-ETSD vaccine	Human adenovirus serotype 5 (hAd5) vector with E1/E2b/E3 deletions expressing viral S fusion protein and nucleocapsid with an enhanced T-cell stimulation domain (ETSD).	Phase 1 NCT04591717
	Ad5-nCoV	Recombinant Novel Coronavirus Vaccine (Adenovirus Type 5 Vector)	Phase 1 NCT04552366
	VXA-CoV2-1	Ad5-vector based vaccine expressing a SARS-CoV-2 antigen and dsRNA adjuvant given as oral tablets	Phase 1 NCT04563702
	MVA-SARS-2-S	Modified Vaccinia Virus Ankara (MVA) vector expressing the SARS-CoV-2 spike protein (S)	Phase 1 NCT04569383
DNA	INO-4800	DNA plasmid encoding S protein with electroporation delivery mechanism	Phase 1/2 NCT04447781 NCT04336410
	AG0301-COVID19 AG0302-COVID19	DNA plasmid vaccine administered with adjuvant	Phase 1/2 NCT04463472 NCT04527081
	Novel Corona Virus-2019-nCov vaccine	DNA plasmid vaccine expressing S protein	Phase 1/2 CTRI/2020/07/026352
	GX-19	DNA vaccine expressing SARS-CoV-2 S-protein antigen	Phase 1/2 NCT04445389
RNA	BNT162b1	Prophylactic SARS-CoV-2 mRNA enclosed into lipid nanoparticle	Phase 3 NCT04368728
	mRNA 1273	Novel lipid nanoparticle (LNP)-encapsulated mRNA-based vaccine that encodes for a full-length spike (S) protein of SARS-CoV-2	Phase 3 NCT04470427
	CVnCoV mRNA vaccine	mRNA	Phase 2 NCT04515147
	ARCT-021 mRNA vaccine	Self-replicating mRNA encoding SARS-Cov-2 spike protein encapsulated in lipid nanoparticle	Phase 1/2 NCT04480957
	LNP-nCoVsaRNA vaccine	Purified mRNA which mimics the virus gene for a spike protein on its surface	Phase 1 ISRCTN17072692
	SARS-CoV-2 mRNA vaccine	mRNA vaccine	Phase 1 ChiCTR2000034112 ChiCTR2000039212
Inactivated	Adsorbed COVID-19 (inactivated) Vaccine	Inactivated SARS-CoV-2 with adjuvant alum	Phase 3 NCT04456595 669/UN6.KEP/EC/2020 NCT04582344
	Inactivated SARS-CoV-2 vaccine (Vero cell)	Inactivated SARS-CoV-2	Phase 3 ChiCTR2000034780 ChiCTR2000039000
	Inactivated SARS-CoV-2 vaccine (Vero cell)	Inactivated SARS-CoV-2	Phase 3 ChiCTR2000034780 NCT04560881
	Inactivated SARS-CoV-2 Vaccine	Inactivated SARS-CoV-2	Phase 1/2 NCT04470609
	QazCovid-in® - COVID-19 inactivated vaccine	Inactivated SARS-CoV-2	Phase 1/2 NCT04530357
	BBV152	Whole Virion Inactivated SARS-CoV-2	Phase 1/2 NCT04471519 CTRI/2020/09/027674
	Inactivated SARS-CoV-2 Vaccine (Vero Cells)	Inactivated SARS-CoV-2	Phase 1 ChiCTR2000038804
Replicating Viral Vector	TMV-083	Live-attenuated recombinant measles vaccine virus vector expressing a modified surface glycoprotein of SARS-CoV-2	Phase 1 NCT04497298
	V590	Replication-competent VSV delivering the SARS-CoV-2 Spike	Phase 1 NCT04569786
	DelNS1-2019-nCoV-RBD-OPT1	Intranasal flu-based-RBD	Phase 1 ChiCTR2000037782
	rVSV-SARS-CoV-2-S Vaccine	Replication-competent SARS-CoV-2 Spike protein	Phase 1 NCT04608305
Protein Subunit	SARS-CoV-2 rS	Full length recombinant SARS CoV-2 glycoprotein nanoparticle vaccine adjuvanted with Matrix M	Phase 3 2020-004123-16
	Recombinant new coronavirus vaccine (CHO cells)	Adjuvanted recombinant protein (RBD-Dimer)	Phase 2 NCT04466085
	KBP-COVID-19	RBD-based	Phase 1/2 NCT04473690
	Recombinant S protein	Recombinant S protein (baculovirus production) with different adjuvants	Phase 1/2 NCT04537208
	SCB-2019	Native like Trimeric subunit Spike Protein vaccine	Phase 1 NCT04405908
	COVAX-19 vaccine	Recombinant spike protein with Advax™ adjuvant	Phase 1 NCT04453852
	MF59 adjuvanted SARS-CoV-2 Sclamp vaccine	Molecular clamp stabilized Spike protein with MF59 adjuvant	Phase 1 ACTRN12620000674932p ISRCTN51232965
	MVC-COV1901	Spike protein with CpG 1018 and aluminum content as adjuvant	Phase 1 NCT04487210
	SOBERANA 01	RBD + Adjuvant	Phase 1 IFV/COR/04
	EpiVacCorona	Peptide antigens of SARS-CoV-2 proteins, conjugated to a carrier protein and adsorbed on an aluminum-containing adjuvant (aluminum hydroxide)	Phase 1 NCT04527575
	Recombinant SARS-CoV-2 vaccine (Sf9 cell)	RBD (baculovirus production expressed in Sf9 cells)	Phase 1 ChiCTR2000037518
	pVAC	SARS-CoV-2 HLA-DR peptides	Phase 1 NCT04546841
	UB-612	High-precision designer S1-RBD-protein based vaccine containing a Th/CTL epitope peptide pool, that could bind to human MHC-I and MHC-II to activate T cells	Phase 1 NCT04545749
Virus Like Particle	RBD-HBsAg VLPs		Phase 1/2 ACTRN12620000817943

(continued on next page)

Table 4 (*continued*)

Vaccine Platform	Vaccine Name	Vaccine characteristics/Immunogen	Stage of Development[a]
	Coronavirus-Like Particle COVID-19 Vaccine	RBD antigen is conjugated to the hepatitis B surface antigen that stimulates immune system to produce anti-RBD antibodies Plant-derived VLP unadjuvanted or adjuvanted with either CpG 1018 or AS03	Phase 1 NCT04450004

[a] Source: (Cochrane, 2020; National Library of Medicine, 2020; World Health Organization, 2020).

5.3. DNA platform

DNA-based vaccines were introduced two decades ago and these are non-infectious and non-replicating. These are easy to produce within a short duration and are stable and cost-effective at the same time. These confer long term immunogenicity to the host, however these remain hopeless when used in humans due to their poor immunogenic property. Also these are easily degraded by host enzymes and there is always the risk of its integration into host DNA. While it has shown efficacy in animal models, but yet to establish the potency in human trials (Kim , 2009). Currently, there are 4 COVID-19 vaccine candidate in clinical evaluation and 14 candidates in preclinical evaluation stage developed using DNA platform (World Health Organization, 2020).

5.4. RNA platform

After the failure of DNA based vaccines, researchers established the RNA-based vaccines which have several advantages over DNA based vaccines. The mRNA is directly injected into the host's cell, which undergo translation in the cytoplasm. Currently, there are two kinds of mRNA-based vaccines established: non-amplifying mRNA based vaccines and self-amplifying mRNA based vaccines (Rodríguez-Gascón et al., 2014). The self-amplifying mRNA based vaccines (SAM) capable of being rapidly developed are potentially feasible for epidemics or disease outbreaks as observed in Shanghai, where within 7 days of H7N9 influenza outbreak the vaccine developed by SAM technology was tested in mice (Hekele et al., 2013). The SAM vaccine technology is capable of swift and cost-effective vaccines production. These can be devised for both therapeutic as well as prophylactic purpose as shown by variety of preclinical and clinical projects while the poor immunogenicity remains their biggest limitation although it can induce antigen-specific T and B cell immune response. Currently, there are 6 COVID-19 vaccine candidates in clinical evaluation and 19 candidates in preclinical evaluation stage developed using RNA platform (World Health Organization, 2020).

5.5. Recombinant protein based vaccine

Highly purified recombinant proteins from different etiologic agents are most common candidates under investigation for vaccines. Different genes from virus particles encoding the antigenic determinant have been processed (cloned, expressed in various expression system and purified) as recombinant proteins and established as vaccines. For the antigens which undergo post-translational modifications, mammalian cells expression systems can used while the bacterial expression systems provide the ease of handling and high level expression (Nascimento and Leite, 2012). It is the most common of all the platforms used for production of vaccines since it possesses several advantages such as safety profiles and cost effective production but the adjuvants are required to produce long-lasting immune response. There are several examples of recombinant protein based vaccines being used in humans like hepatitis B vaccine (HBV) (Kumar and Meldgaard, 2018). Currently, there are 13 COVID-19 vaccine candidates in clinical evaluation and 55 COVID-19 vaccine candidates in preclinical evaluation stage developed as recombinant protein subunit (World Health Organization, 2020).

5.6. Live attenuated vaccines

The most common traditional method which involves manually weakened live pathogen which is no longer able to induce infection but able to induce immune response and hence mimic features of natural infection. It is capable of inducing both humoral and cellular immune response. Live attenuated vaccines on intranasal administration induces secretion of IgA and hence provide local mucosal immunity (Barría et al., 2013). These vaccines are popular to induce strong lifelong immune responses within two doses. These are easy to produce for some viruses but challenging for complex pathogens. Most common example is Oral Polio Vaccine (Kumar and Meldgaard, 2018). Currently, there are 3 COVID-19 vaccine candidates in preclinical evaluation stage developed using this platform (World Health Organization, 2020).

5.7. Inactivated vaccines

These are produced by completely inactivating or killing the pathogen, on injecting it to the host, they primarily induce protective antibodies against epitopes on hemagglutinin glycoprotein on surface of virus. The subunit formulations that primarily consist of a part of killed pathogen in spite of the whole pathogen are used more frequently than the other formulations because potential association of whole virus formulation with increased reactogenicity. These vaccines tend to produce a weaker immune response than live attenuated vaccines, thus adjuvants are required to provide an effective immune response (Kumar and Meldgaard, 2018). Inactivated Polio Vaccine is the classical example where the whole killed pathogen is provided and Diphtheria and Tetanus vaccination is the example of subunit formulation (Nascimento and Leite, 2012). Currently, there are 7 COVID-19 vaccine candidates in clinical evaluation and 12 candidates in preclinical evaluation stage developed using this platform (World Health Organization, 2020).

5.8. Plant based vaccines

There are numerous limitations of injectable vaccines including its high cost and maintenance, transportation, and requirements of trained medical personnel, making its availability more difficult specifically in the developing countries. On contrary, plant based vaccines being capable of producing edible vaccines can be easily available globally to the needy population. Various ongoing reports have investigated the development and stability of edible vaccines (Sartaj Sohrab et al., 2017). The another low cost alternative could be the plant based virus like particles (VLPs) used against H1N1 influenza. Medicago Inc. has described the production of VLPs in plants (Rosales-Mendoza et al., 2020). The specific proteins can be expressed into the desired plant with the help of agrobacterium mediated transformation and can be grown in the desired location. Antigens are released as bioencapsulated vaccine from the ingested recombinant plant which later on get absorbed by M cells in the intestine and pass to the macrophages and antigen presenting cells thereby activating immune system and induce production of specific antibodies and memory T cells against the antigen (Sartaj Sohrab et al., 2017). So far many of the plant based vaccine candidates have been tested and entered into clinical trials like the vaccine against swine flu, hepatitis B (Govea-Alonso et al., 2014). As per the news articles,

Medicago Inc. has started Phase I trial of its plant derived COVID 19 vaccine named, CoVLP following the single adjuvanted mediated positive immunogenic effects obtained in the preclinical testing. According to the enterprise, VLPs have a multi-modal mechanism of action that works differently to inactivated vaccines, by activating both antibody and cell-mediated responses. The adjuvants that will be used alongside the CoVLP vaccine candidate are GlaxoSmithKline's pandemic adjuvant technology and Dynanvax's CpG 1018 (Balfour, 2020).

6. Natural products as immunomodulators

Natural products mainly comprising of Ayurvedic medicines, and Traditional Chinese medicines (TCM) are used for decades as the regulator of the immune system. These compounds either alone or in combinations are used for centuries in both Ayurveda and alternative medicine. Studies have documented their role in the treatment or prevention of many viral infections including influenza, HIV, SARS, and MERS (Gyawali et al., 2020; Islam et al., 2020; Leung, 2007). These medicines boost the immune system and also participates in the innate and adaptive immune regulatory pathways (Wang et al., 2018). The use of natural products and TCM to prevent or treat COVID-19 is largely inspired by the treatment of SARS. Many trials are registered in the clinicaltrials.gov for investigating the role of traditional Chinese medicines and natural products (Nigella sativa, natural honey, curcumin, Boswellia, and vitamin C) in COVID-19 patients (National Library of Medicine, 2020). Compounds like *Withania somnifera* (Ashwagandha), *Tinospora cordifolia* (Guduchi), *Asparagus racemosus* (Shatavari), *Phylanthus embelica* (Amalaki), and *Glyceriza glabra* (Yashtimadhu) possess the potential immunomodulatory activity and hence they may be considered for prophylaxis and as an add-on treatment for COVID-19 (Balasubramani et al., 2011; Tillu et al., 2020). Rajkumar et al. have highlighted the possible role of Ayurveda in the treatment of COVID-19 through psychoneuroimmune pathways (Rajkumar, 2020). Apart from Ayurveda, medicines from traditional Siddha system has been found effective as immune system booster in COVID-19 patients. Kabasura kudineer, a herbal concoction regarded as immune booster and capable of combating viral infections found to be helpful in regaining the lost sense of smell in COVID 19 patients. Currently a clinical trial is ongoing on Kabasura kudineer and vitamin C-zinc supplementation in the management of mild COVID 19 patient (CTRI/2020/05/025215, 2020; Livemint, 2020; Soruban and Rajeev, 2020). TCM like Chinese Rhubarb extracts, Houttuynia cordata extract, hesperetin, Fructus Forsythiae, etc. have also demonstrated anti-inflammatory activity by inhibiting the cytokines like TNF-α, IL-1β, and IL-6 and anti-viral activity by inhibition of 3CLpro and RdRp protein (Yang et al., 2020a,b). Recently, a systematic survey by Lee et al. has reported the beneficial role of TCM in COVID-19 patients (Lee et al., 2020). TCM has been used as a treatment of COVID-19 in China, and Qingfei Paidu decoction has reported treating 214 confirmed cases. Yang et al. have reported that the rate of TCM treatment of COVID-19 in China was 87% and only 5% of patients have worsened clinical manifestations (Yang et al., 2020a,b). To date, more than 50 clinical trials have registered in China for the evaluation of TCM in COVID-19. Although shreds of evidence are available for the efficacy of TCM and natural products in COVID-19 patients, still the randomized clinical trials in larger populations are required to evaluate their efficacy and safety in COVID-19 patients.

7. Current status of vaccine candidates in pipeline

Currently, as of 29th October there are 201 vaccine candidates active globally, of which 45 are in clinical phase and 156 in preclinical phase. List of vaccine candidates under different platforms in different evaluation stages summarized in Table 4. The time required by these vaccine candidates to become available in the market depends on the success of all the phases of clinical trials. As discussed further in this section, 10 of the vaccine candidates have successfully passed phase 1 and phase 2 trials and found to have generated anti-SARS-CoV-2 immune responses and are safe to use, however phase 3 trials for the same are ongoing.

7.1. ChAdOx1 nCoV-19 vaccine

ChAdOx1 nCoV-19 vaccine (AZD1222) consists of the non replicating simian adenovirus vector ChAdOx1 containing the full-length structural spike protein of SARS-CoV-2 developed by University of Oxford and AstraZeneca. The preliminary report released by the research group shows that the candidate ChAdOx1 nCoV-19 vaccine given at a dose of 5×10^{10} viral particles was safe and tolerated, however a higher reactogenicity profile was observed than the control vaccine, meningococcal conjugate vaccine MenACWY. The reactogenicity was reduced on administration of 1 g paracetamol for the first 24 h after vaccination. 14 days after the first dose of vaccine the cellular responses were observed and on day 28 post vaccination the humoral responses to spike protein, i.e. the spike specific antibodies were found to be at peak. The second dose of vaccine administered after 28 days of first dose induced neutralizing antibodies in all the participants and the reactogenicity was also found to be reduced after the second dose. Overall, the two doses of vaccine was found to be sufficient to induce potent cellular and humoral immunogenicity in all the participants and no serious adverse reactions occurred. The study was conducted in the healthy individuals excluding all the potential participants with recent COVID-19-like symptoms or with a history of positive PCR test for SARS-CoV-2, however a small number of participants found to have high levels of neutralizing antibody at baseline which probably indicates prior asymptomatic infection (Folegatti et al., 2020).

In conclusion, ChAdOx1 nCoV-19 showed an acceptable safety profile, and sufficient immunogenicity and reduced reactogenecity when given with paracetamol. A single dose found to elicit both humoral and cellular responses against SARS-CoV-2 and booster dose required to enhance neutralizing antibody concentration. The preliminary results supported the large scale evaluation of vaccine in an ongoing phase 3 programme.

7.2. mRNA-1273 vaccine

The mRNA-1273 vaccine is a lipid nanoparticle encapsulated, nucleoside-modified messenger RNA (mRNA)-based vaccine developed by Moderna and NIAID. It encodes the spike glycoprotein (S-protein) present on the surface of SARS-CoV-2. The preliminary findings reported that it generate sufficient immunogenic response and induce vigorous binding antibody responses to both full-length spike protein and receptor-binding domain in all participants after the first vaccination. The immune responses were found to increase with time and increasing dose of the vaccine i.e. 25-μg, 100-μg and 250-μg. And high neutralizing antibody responses were elicited in a dose-dependent fashion. The two-dose vaccination schedule was finalized because of its low pseudovirus neutralizing activity even after the 2 weeks of first vaccination. No serious toxicity was reported after the two dose series of vaccination; systemic adverse events after the first vaccination reported were all graded mild to moderate in severity. Of the three doses evaluated 250 μg, 100 μg and 50 μg, the 100 μg dose elicits high neutralization responses and more favorable reactogenicity profile than that of the higher dose. The rapid and vigorous immunogenicity profile of the candidate vaccine is attributed to the presentation of naturally folded conformation of the S protein designed as antigen coupled with lipid nanoparticle delivery system and using modified mRNA to avoid early inactivation of interferon associated genes. The current report released by research group comprised the result of follow-up of participants through day 57, hence the durability of the immune responses and their characterization is not yet established. A phase 2 trial of mRNA-1273 in 600 healthy adults, evaluating doses of 50 μg and 100 μg is ongoing (ClinicalTrials.gov number, NCT04405076) and the phase 3 trial to evaluate 100 μg dose is expected to begin soon (Jackson et al., 2020).

7.3. Ad5-vectored COVID-19 vaccine

The candidate vaccine is a type of non replicating adenovirus type Ad5-vectored COVID-19 vaccine developed by CanSino Biological Inc. and Beijing Institute of Biotechnology. It is found to elicit both cellular and humoral response in 95% of the participants by the day 28 post vaccination in both doses, i.e. 1×10^{11} viral particles and 5×10^{10} viral particles. No serious adverse reactions were reported, those reported were all ranging from mild to moderate in severity and resolved within a short period of time maximum by 48 h of reaction onset. Earlier three doses were given to the participants, but higher proportions of grade 3 adverse reactions were reported in the higher dose of 1.5×10^{11} viral particles compared with the low dose (5×10^{10} viral particles) and middle dose (1×10^{11} viral particles). Further investigation revealed that the vaccine at 5×10^{10} viral particles had a better safety profile than the vaccine at 1×10^{11} viral particles and also shows the better immunogenicity profile. The preliminary report suggests the acceptable safety profile of Ad5-vectored COVID-19 vaccine. Apart from the results, the major obstacle that the researchers face with this vaccine candidate was the high pre-existing anti-Ad5 immunity in the participants with older age (55 years or older) due to which single dose of vaccine was not inducing high levels of humoral immune responses. Therefore, researchers proposed to use the additional dose of vaccine in the older age population to gain the required effects. They will be evaluating it in phase 2b stages of the ongoing trial. Moreover rest of the population showed better results (in terms of safety profile and adequate immunogenicity) with the single immunization of Ad-5 vectored COVID-19 vaccine at dose 5×10^{10} viral particles making it a potential candidate for emergency vaccination (Zhu et al., 2020). In conclusion, the results of the trial have shown acceptable results supporting the testing of candidate vaccine at dose 5×10^{10} viral particles further in phase 2b and phase 3 trial in healthy adults.

7.4. Inactivated COVID-19 vaccine

The candidate vaccine is an inactivated whole SARS-CoV-2 virus vaccine developed by the Wuhan Institute of Biological Products and Sinopharm. The whole virus pathogen was cultivated in vitro in cell lines and the infected cells further inactivated twice using β-propiolactone under specific conditions and further adsorbed to 0.5 mg alum. The phase 1 trial was conducted using three doses with 2.5 µg, 5 µg and 10 µg antigen protein content per dose and the results showed the inactivated vaccine candidate to have better safety profiles comparable to the other vaccines under different platforms and the neutralizing antibody titres induced were also adequate in all the three doses. No serious adverse event was reported, only mild to moderate self limiting reactions were reported and the incidence rate of adverse reactions associated with the current vaccine candidate was very low nearly 15% as compared to the other vaccines where the incidence rate is reported around 60% to even 100% in some studies. Phase 2 trial was conducted using a middle dose of vaccine, i.e. 5 µg antigen protein content and the results showed the vaccine candidate is able to effectively produce antibody titer with minimal adverse reactions reported. However researchers came to conclusion that booster dose is required to generate adequate immunogenic response, also the Phase 1 and Phase 2 trial indicated that with longer interval (21 days and 28 dya) between the first dose and subsequent booster dose generates higher antibody titers as compared to shorter interval group (14 day schedule) (Zhu et al., 2020). In conclusion, the interim analysis of the inactivated vaccine showed the acceptable safety and better immunogenic profile of the candidate vaccine supporting its longer term adverse events assessment in ongoing phase 3 trials.

7.5. BNT162b1 RNA vaccine

The candidate vaccine is the lipid nanoparticle encapsulated mRNA based SARS-CoV-2 vaccine named as BNT162b1 developed by Bio-NTech, Fosun Pharma and Pfizer. It consists of the nucleoside modified messenger RNA that encodes the receptor binding domain of the SARS-CoV-2 spike protein. Researchers have administered the BNT162b1 vaccine candidate at three different doses 10 µg, 30 µg and 100 µg and found that the vaccine candidate exhibit acceptable safety profile and elicited adequate antibody titer. Based on the tolerability profile, the participants receiving two of the doses 10 µg and 30 µg was given second booster dose and found to show sharper dose vs immunogenic responses while no data for immunogenic response was reported for the dose 100 µg till now since no second vaccination was provided to the participants to avoid the adverse reactions and the study is ongoing. Results indicated higher reactogenicity after the booster dose, however symptoms were limited to mild to moderate in severity which were resolved within few days of onset (Zhu et al., 2020). In conclusion, the RNA based vaccine candidate BNT162b1 showed acceptable safety profile and also found to produce adequate antibody titre after booster dose, hence it supports the long term assessment of the candidate vaccine for efficacy and tolerability in the ongoing phase 3 trials.

7.6. Gamaleya's Sputnik V

It is the type of non replicating adenovirus (rAd26-S + rAd5-S) type vaccine developed by Gamaleya Research Institute Epidemiology and Microbiology, Health Ministry of the Russian Federation. The vaccine consists of the two components, one with the recombinant adenovirus vector based on the human adenovirus type 26 and another with the adenovirus vector based on the human adenovirus type 5, both containing SARS-CoV-2 S protein gene. This adenovirus vector-based vaccine was registered by the Russian Ministry of Health on August 11 and became the first registered COVID-19 vaccine on the market. Prior to its registration and market launch, the vaccine had successfully cleared preclinical trial on various animals and also its safety and efficacy was assessed in humans in Phase 1 and 2 clinical trials. The first component, adenovirus vectored type 26 vaccine containing SARS-CoV-2 S protein gene was given as the first vaccination dose and another component, adenovirus vectored type 5 vaccine containing SARS-CoV-2 S protein gene was given as second vaccination dose to all the participants. No unwanted side effects were reported and the vaccine induced strong antibody and cellular immune response. Currently, the post registration Phase 3 trial is ongoing involving more than 40000 participants in Russia and Belarus and the plan to execute the Phase 3 trial of Sputnik V in various other countries like UAE, India, Venezuela, Egypt and Brazil is underway (Sputnik V, 2020).

7.7. Bharat Biotech's Covaxin

It is the India's first vaccine against COVID-19 and is an inactivated whole SARS-CoV-2 virus vaccine developed by Bharat BioTech, Indian Council of Medical Research (ICMR) and National Institute of Virology (NIV). It has successfully cleared Phase I/II human trial and is given permission for Phase III trial by DCGI (Drug Controller General of India). The vaccine trial Phase I/II has been conducted in 12 hospitals in different cities across country, in healthy volunteers with no co-morbidity conditions. The vaccine was found to generate robust immune responses, thereby preventing infection and disease in the primates upon high amount of exposure to live SARS-CoV-2 virus. As per the vaccine makers the phase III study would cover around 28,500 subjects aged 18 years and above, however the complete safety and immunogenicity data of the phase II trial is still not revealed (Ray, 2020; Sharad Sharma, 2020).

7.8. Cadila's ZyCoV-D vaccine

It is India's second vaccine, a type of DNA plasmid vaccine expressing SARS-CoV-2 S protein developed by Zydus Cadila Healthcare.

Currently, the phase II trial of the candidate vaccine is underway. The Phase I trial of vaccine was conducted on healthy volunteers and was found safe in all the participants (Ray, 2020; CTRI/2020/07/026352, 2020).

8. Concluding remarks

COVID-19 pandemic has imposed a huge financial and social burden to the world. With regard to its continuously evolving nature, it remains a challenging task to restrict the transmission of infection among the population. One of the prominent approaches for preventing SARS-CoV-2 infection is the development of vaccines against COVID-19. A great deal of work has done so far and till now 33 vaccine candidates have entered into a clinical trial and 143 are in the preclinical phase. Despite the extraordinary efforts of the firms and researchers across the globe for developing the COVID-19 vaccines, there is still a lot more to explore that could be helpful in developing more specific vaccine. Gathering of important information like immunization route, finding more target antigen(s) apart from targeting S protein as antigen only, can be helpful in eradicating the infection. The immunopathological basis of COVID-19 need to be explored more so that its immune evasion mechanism can be targeted and it will also provide deeper insights into vaccine designing strategies. The three main criteria which should be taken into consideration while developing a vaccine are: speed, scale up manufacturing and global access. The astonishing efforts by researchers across the globe in terms of scale and speed of vaccine development have fastened the vaccines development journey from bench to bedside within a few months only. The work done so far in this regard is very commendable as 5 of the candidates are already in Phase 3 clinical trial and hopefully will be available by the end of 2020. This represents a huge change in the conventional pathway of vaccine development which usually takes 5–10 years timescale. Along with speeding up the development process, it is equally important to evaluate the effectiveness and safety of vaccine at each step, and this has been the major hurdle for researchers in establishing the vaccine's efficacy so far. However, the phase 3 trial will play a major role in establishing the vaccine that is safe and effective in a large and diverse population. Finally, the animal models specific for COVID-19 should be established in order to assess vaccine efficacy. Also, the regulatory authorities should need to ensure the equal distribution of vaccines to all the affected areas.

Declaration of competing interest

The authors declare no conflict of interest.

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