Recent advances in therapeutic applications of neutralizing antibodies for virus infections: an overview

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Abstract
Antibodies are considered as an excellent foundation to neutralize pathogens and as highly specific therapeutic agents. Antibodies are generated in response to a vaccine but little use as immunotherapy to combat virus infections. A new generation of broadly cross-reactive and highly potent antibodies has led to a unique chance for them to be used as a medical intervention. Neutralizing antibodies (monoclonal and polyclonal antibodies) are desirable for pharmaceutical products because of their ability to target specific epitopes with their variable domains by precise neutralization mechanisms. The isolation of neutralizing antiviral antibodies has been achieved by Phage displayed antibody libraries, transgenic mice, B cell approaches, and hybridoma technology. Antibody engineering technologies have led to efficacy improvements, to further boost antibody in vivo activities. “Although neutralizing antiviral antibodies have some limitations that hinder their full development as therapeutic agents, the potential for prevention and treatment of infections, including a range of viruses (HIV, Ebola, MERS-COV, CHIKV, SARS-CoV, and SARS-CoV2), are being actively pursued in human clinical trials.”

Keywords Humoral immune response · Epitope · Neutralizing antibodies · Monoclonal · Polyclonal · Phage display · Glycoprotein

Introduction
An antibody is a protective protein produced by the immune system in response to the detection of foreign material known as antigen [1]. In 1890 Emil von Behring and Shibasaburo Kitasato discovered antibodies as protective antitoxins in the blood of animals exposed to diphtheria or tetanus toxin [2]. To date, antitoxins are still in use as a preventive and treatment option for some infections [3]. Antibodies have been a significant research subject due to their crucial role in adaptive immunity and their wide range of specificities [4]. Although vaccines aim to induce immune response in host cells to produce antibodies, the high rate of genetic mutation in viral strains increases the virus’ possibility of reactivation (due to loss of antibody specificity), which can lead to ineffective vaccination as in the case of HIV-1 [5, 6]. Although vaccines aim to induce immune response in host cells to produce antibodies, the high rate of genetic mutation in viral strains increases the virus’ possibility of reactivation (due to loss of antibody specificity), which can lead to ineffective vaccination as in the case of HIV-1 [5, 6].

Antibodies have been used for the development of vaccines with little use as biological therapy to combat viral infections [7, 8]. A new generation of broadly cross-reactive and highly
potent antibodies has led to a unique chance for intervention [9]. Antibodies are desirable for pharmaceutical products because of their ability to target specific epitopes with their variable domains.

An antiviral antibody is a unique research discovery used by virologists to better understand the exact mechanisms of binding and entering of viruses into hosts and their antiviral immunity. Based on the mechanisms of action, antiviral antibodies can be categorized into neutralizing and non-neutralizing antibodies. Neutralizing antibodies act by binding to a specific epitope on the viral envelope and lead to neutralization of the virus infection by several mechanisms which have been mentioned in details in this review, notably neutralizing antibodies have shown great advances in clinical trials targeting many viruses. In non-neutralizing antibodies, the antibody bind to an area on the surface of the virus without neutralizing the infection or total removal of viral titer. As the antibody does not block infection, the whole virus-antibody complex enters the cell via endocytosis. Then other non-neutralizing humoral effector functions perform as antibody-dependent cell-mediated cytotoxicity (ADCC) or antibody-mediated phagocytosis, aggregation, and even immune activation [10–13].

The versatility of therapeutic antibodies to combat cancer, autoimmune diseases and passive antibody therapies for infectious diseases have been reviewed in the past [8, 14, 15]; however, this review article mainly focuses on recent advancements in the use of neutralizing antibodies as an established therapeutic antiviral treatment in humans including how antibodies neutralize viruses, their clinical applications, isolation of potent and broadly neutralizing antiviral antibodies, and the limitations of neutralizing antiviral antibodies.

Neutralizing antibodies (NAbs)

A neutralizing antibody (NAb) is an antibody that protects the host cell from pathogens by neutralizing or inhibiting its biological effect. It blocks interactions of the viral envelope with the host cell receptor or inhibits the release of the viral genome. NAbs that emerge under different constraints are sentinels that provide a better understanding of the humoral immune responses, as well as great insight for the future development of immunotherapeutics [16].

NAbs are the foundation for successful disease prevention; nevertheless, for critical chronic infections such as HCV and HIV, the use of potent and broad NAbs from recombinant vaccines is incomplete [17, 18]. NAbs can inhibit multiple strains of a particular virus; next to vaccination, neutralizing antibodies gives the best protection against viral infections rather than anti-retroviral therapy ART [19, 20].

Monoclonal antibodies (mAbs)

Monoclonal antibody (mAb) has one type of immunoglobulin with defined specificity and a single isotype [21]. These types of antibodies are used clinically against particular antigens and exhibit therapeutic effects via the antigen-binding fragment (Fab).

Recently, in the Europe and the USA, about 80 monoclonal antibodies have been approved and several hundred of them in clinical trials. In the last decade and a large number of antibodies have been engineered, including novel antibody-like scaffolds, antibody-drug conjugates, and bispecific antibodies [22]. Recently, developed isolation technology produces humanized mAbs from immune and non-immune sources, and that has made distinctive progress in using antibodies for treatment [23].

Antiviral mAb has been considered as a new immunotherapy treatment to eradicate the outbreak of chronic infections [9]. Antiviral mAbs have characteristic features which include:

a) Ability to target specific epitopes
b) Highly specific
c) High potency
d) Minimal side effect in clinical applications, which enables and encourages large scale production

Polyclonal antibodies

Polyclonal antibodies are a diverse group of antibodies that target numerous epitopes on the viral envelope during the infection and provide vigorous neutralizing activity [24]. The ability of polyclonal antibodies to target multiple epitopes enables them to trigger a range of effector functions such as steric hindrance (prevent antibody- virus complex from attaching the host surface), aggregation (binding of antibodies to many viruses lead to clearance them from circulation), opsonization (activation of phagocytes cells), and activation of the complement system [25]. There are several advantages of polyclonal antibodies over monoclonal antibodies; low cost and a short production period, they are highly stable and with high affinity; besides, polyclonal antibodies are less sensitive to antigen changes than monoclonal antibodies. In contrast, due to their lack of specificity and a high degree of cross-reactivity of polyclonal antibodies, it has limited their use in the clinic [26].

Techniques for isolation of neutralizing antiviral antibodies

During viral infection in humans, there is a natural immune response toward the virus, which activates innate and humoral
immune cells to produce B cells. These B cells then produce antibodies to attack the virus.

Various approved strategies have been used to isolate and purify antibodies with specific characteristics and proper function.

**Phage displayed antibody libraries**

Phage display is a modern technique of using bacteriophage (viruses that infect bacteria) to express a unique protein variant (such as antibody fragments).

In phage display, a gene encoding a protein of interest is inserted into a phage, causing the phage to display the protein on the outside while containing the gene with the protein on the inside, resulting in a connection between genotype and phenotype [27].

Phage display begins with the amplification of the heavy and light chain gene of the target antibody and the construction of a recombinant phage display plasmid called phagemid. Once bacteria are co-infected with phagemids and helper phages, the phagemid will be replicated, translated, and assembled into infective phages with the target monoclonal antibody fragment displayed on the surface.

Phage display is further used to screen and select the antibody with unique properties, that is, phage-antibody library technology [28]. This process begins with antibody library preparations, followed by ligation of the variable heavy and variable light PCR products into phagemids. Analysis of the clone’s monoclonal antibodies is then performed (Fig. 1a).

A vast antibody library and efficient selection are needed to isolate specific monoclonal antibody from the cloned immunoglobulin repertoire. The library is screened for phage binding to an antigen through a surface monoclonal antibody biotechnology, called biopanning. Multiple rounds of phage binding to antigen, washing, elution, and propagation of the phage binders are to be done. During each round, specific binders are selected out from the pool by washing away non-binders and selectively eluting the binding phage clones. After 3 or 4 rounds, the highly particular binding of phage clones through the surface map is characteristic for directed selection on the immobilized antigen (Fig. 1a). This technique has been applied for isolating neutralizing antibodies that have the potential for a range of viruses (severe acute respiratory syndrome (SARS), Ebola virus, yellow fever virus, hepatitis C virus, measles virus, rabies virus, and influenza virus [24, 27]). Moreover, using yeast phage display technology has led to producing novel mAb against HIV-1 [29].

**Transgenic mice technique**

This technique requires the cloning of human mAb-expressing B cells. A human mini-immunoglobulin gene locus is knocked-in into a transgenic mouse (Fig. 1b). The human mAb-expressing B cells are then isolated from the immunized transgenic mouse and cloned [30]. The process begins with injecting the desired antigen into mice; until the mice develop an immune response, then, the isolated B-lymphocytes from mice’s spleen are fused with a myeloma cell line, which leads to producing immortalized B-cell myeloma hybridomas. The antibodies that are produced are then screened for the desired mAb [31].

Antiviral antibodies against rabies and acute respiratory syndrome coronavirus (SARS-CoV) have been successfully isolated by using this technology [32, 33]. Furthermore, the isolation of (5F10 and 8B10) mAbs showed broad neutralization against CHIKV infection in vitro [34], and the 20 heterosubtypic mAbs showed promising results in neutralizing influenza virus [35].

**Single B cell technique**

This isolation technique can be performed either in an antigen-selective manner from peripheral blood or in a random way [36]. Depending on the transformation method, Memory B cells collected from the blood of patients recuperating from viral infection are transformed with Epstein-Barr virus (EBV) and polyclonal memory B cell-activating elements, such as irradiated mononuclear cells and CpG oligonucleotides (Fig. 1c). These activating elements are TLR4 (Toll-like receptor 4) ligands with immense transformation efficiency [37]. Production of NAbs can be from B memory cells and plasma of non-human primate (NHP) [38–40]. For example, NAbs that have been produced from memory B cells are isolated from persons recovering from dengue and H1N1 influenza infections [41–43].

**Hybridoma technology and humanization**

The hybridoma is the most conventional technique for producing therapeutic antibodies from non-human sources. However, a further process called humanization is required.

In the mid-1970s, the discovery of hybridoma technology by César-Milstein’s group led to the isolation of monoclonal antibodies from mice that had distinct properties and neutralization capacity [44].

Humanization occurred by grafting complementarity determining regions (CDRs) that determine the binding site of the antibody. The appropriate framework region of the variable domains of a rodent donor mAb is grafted onto acceptor human antibody frameworks (Fig. 1d). It can reserve the specificity and binding affinity of the mAb with a low risk of immunogenicity.

Smart antibody-screening methods and proper selection of human donors lead to advanced isolation of rare, broadly (NAbs) for several viruses.
Mode of viral neutralization

Viral neutralization is a process by which antibodies prevent viral infection [45, 46]. Enveloped viruses enter the host cell by attachment to the host cell surface receptor. In contrast, non-enveloped viruses enter via lysis of the host cell membrane or making pore-like structures in the membrane. The fusion of virus (enveloped/non-enveloped) with host cell membrane needs particular conformational changes in the viral protein that can be triggered by low pH in endosomes [47].
Antibodies selectively bind to specific epitopes on the target antigen either by their variable or constant regions [48, 49] (Table 1). There are various mechanisms by which therapeutic antiviral antibodies function to neutralize viruses [24, 27]. Antibodies neutralize viruses either at the virus entry point or the post binding stage.

### Inhibition of virus entry into host cells

Viruses are neutralized in two ways at the stage. The antibody variable domain binds to epitopes on the viral glycoprotein envelope (Fig. 2a), or antibody through the interaction of fab region binds to FcγRs receptor or coreceptor on the host cell and leads to inhibit viral entry. Or post binding inhibition of antibody-virus complex as (c) antibodies can bind to a non-binding region in the virus envelope lead to inhibit the conformational change to allow membrane fusion. (d) for certain viruses that need low endosomal PH for conformational change, antibodies bind to viral inside the endosome lead to inhibit the change in PH to achieve the membrane fusion, and antibodies can inhibit the release of the viral virion

### Table 1 Targets for antibodies on enveloped viruses

| Glycoprotein                  | HIV | Ebola | Influenza | CHIKV | MERS-CoV |
|-------------------------------|-----|-------|-----------|-------|---------|
| Viral spike                   | Envelope (env) | Glycoprotein | Hemagglutinin (HA) | Glycoprotein | spike |
| Receptor binding domain       | Gp120 | GP1   | HA1       | FcγR  | RBD     |
| Fusion domain                 | Gp41 | GP2   | HA2       | Clq, FcγR | DPP4   |

This table shows the epitopes on glycoprotein envelope viral spike, the binding domain region required for entering the host cell, and the fusion binding domain that required in the fusion of the virus into the host cell. All of these can be a good target for antibodies to neutralize a group of viruses.

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**Fig. 2** Modes of viral neutralization. Antibodies neutralize viruses by several mechanisms, either inhibition of virus entry into host cells as (a) Antibodies bind to an epitope in the viral glycoprotein envelope, lead to inhibit attachment to host cells. (b) Antibodies through the fab region can bind to host cell receptors or coreceptors (have FcγR) lead to inhibit viral entry. Or post binding inhibition of antibody-virus complex as (c) antibodies can bind to a non-binding region in the virus envelope lead to inhibit the conformational change to allow membrane fusion. (d) for certain viruses that need low endosomal PH for conformational change, antibodies bind to viral inside the endosome lead to inhibit the change in PH to achieve the membrane fusion, and antibodies can inhibit the release of the viral virion
surface membrane (Fig. 2b). Recently the development of multi-antigen region binding to therapeutic antibodies enhances their neutralization potency [41].

**Post binding inhibition of antibody-virus complex**

The post binding inhibition of viruses occurs in three forms. In some viruses, binding of the antibody to the non-binding region on the virus envelope inhibits conformational changes required for membrane fusion [50], as illustrated in Fig. 2c. In the case of viruses that require low pH in endosomes for conformational changes to trigger viral membrane fusion, the binding of the antibody to the virus inhibits the conformational changes necessary for membrane fusion (Fig. 2d). Also, the antibodies can inhibit viral production by detecting the virion in order not to assemble its genetic material with that of the host cell. All of these mechanisms may not happen with a single virus type. It varies from virus to virus, depending on the viral glycoprotein structure. Perhaps significant neutralization effects of antiviral antibodies are mostly attained by blocking viral entry into the host. In some instances, viruses tend to escape neutralizing antibody responses. However, the Fc fragments of monoclonal antibodies can bind to the immune cells to perform further neutralization activities, such as clearing of circulatory viruses, activate cytotoxic killing or phagocytosis of the infected cell, and even induce activation of host immune responses in vivo (Fig. 3). These further activities happen via antibody-dependent cellular phagocytosis (ADCP) and antibody-dependent cellular cytotoxicity (ADCC). It includes the interaction of the Fc (fragment crystallizable domain) of an antibody with its Fcγ receptors (FcγRs) present on immune cell surfaces [6] [14, 51]. Studies on the functional and molecular properties of monoclonal antiviral antibodies give

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**Fig. 3 The antiviral mechanism through antibody FC fragment.** The Fc fragments of antibodies bind to the immune cells to perform neutralization activities. (a) Binding of complement to antibody activates complement-mediated virolysis and activation of phagocytosis. (b) Binding of antibodies to a phagocyte (Fcγr) via antibody-dependent cellular phagocytosis (ADCP) opsonizes the infected cell. (c) Antibodies bind to natural killer (NK) cells via antibody-dependent cellular cytotoxicity (ADCC) to lyse the infected cells.
excellent prospects to recognize epitopes and target antigens as a fundamental strategy for the development of human vaccines and novel immunotherapy [52].

**Synergistic effects of neutralizing antiviral antibodies**

Presently, there are no approved combinational antiviral antibodies for the prevention or control of many viral infections in humans. On the contrary, there are several prospective combinational therapies for various kinds of viral infections undergoing trials (Table 2). The potency of therapeutic antibodies can be enhanced by multiple antibody engineering technologies. Some of these technologies include the combination of two or more mAbs, the synergy between antiviral drugs and mAbs, and the vector-based delivery of antibodies.

A combination of mAbs shows high neutralizing activity more than that of individual mAbs. Research on Hepatitis B virus and Respiratory syncytial virus shows the neutralization potential of mAb cocktails (a combination of two or more mAbs) [53]. The synergistic ability of antiviral drugs with mAbs has allowed researchers to explore modes of treatment and routes for administration in certain viruses such as influenza [54]. As in the case of CHIKV virus, using a combination of CTLA4-Ig and antiviral antibody therapy can control severe CHIKV disease and arthritis and might be a great candidate for testing in humans [55].

The vector-based delivery of genes encoding antibody molecules is another synergy that increases neutralization activity. Transgenic mice gained full protection against HCV viral infection when treated with Adeno-associated viral vectors in combination with three bNAbs (AR4A, AR3A, and AR3B) [56].

Drugs can be coupled to antibodies via chemically stable linkers or by direct chemical bonding. Immunological and pharmacological principles determine the effective delivery of antibody-conjugated antiviral drugs. The antibody should be highly specific for a target site, and an adequate number of drug molecules should reach the target to exert their pharmacological effects. Regardless of the mode of coupling, drug-antibody complexes are internalized either by passive diffusion or by some active transport mechanism. Complexes internalized in pinocytic vesicles, ultimately, are digested by lysosomal enzymes that cleave the pharmacologically active drug from the antibody. The second mechanism of cell delivery is by the extracellular release of the drug on the surface of specific target cells.

**Clinical applications of neutralizing antiviral antibodies**

Anti-retroviral therapies (ARTs) are mostly used for the treatment and prevention of viral infections. Nevertheless, discontinuous administration of ARTs can lead to rebound viremia in patients, consequently making the use of ARTs a lifelong treatment. Currently, research suggests that the use of antiviral antibodies is the best alternative to substitute ART because they have the ability to clear circulatory viruses and boost the immune system to prevent viral infections [57, 58].

Unlike the use of antibodies in the treatment of cancer and autoimmune diseases, palivizumab is the only approved antiviral antibody for treating RSV [59].

NAbS show great response toward acute viral infections such as chikungunya virus, Measles virus, and poliovirus due to no evidence of viral escape from the antibody response. While in the chronic infections such as influenza, Ebola, HIV and Lassa virus; The expression of conserved epitopes wherein the neutralization of viruses occurs [15, 60, 61], remains insignificant, due to high mutation in some viral strains [62, 63].

Here, the recent updates on the development of some antiviral antibodies and their successful applications to fight viral infections are going to be discussed.

**NAbS against HIV**

In 1989, CD4-Fc-fusion was the first antibody used to target HIV. The effectiveness of CD4-Fc-fusion proteins has been widely investigated since its use. Currently, a couple of successful treatments have emerged, and it is on the verge of commercialization [64] with enhanced efficacy and safety. These promising results have broadened and encouraged research into the development of vaccines and immunotherapy [65].

Since a competent and safe HIV-1 vaccine is presently not forthcoming, new advances are required to avoid HIV infection [66]. Isolation of bNAbS with high levels of HIV-1 neutralizing activity from HIV-infected individuals is a further advancement to prevent HIV transfection [67-69]. The HIV-1 glycoprotein envelope (ENV) is the main target for NAbS. The ENV can be present on the viral surface in two forms, open or closed conformations with multiple epitopes such as V2 region, V3 region, CD4 binding site, MPER, and CD20 [70]. Certain groups of HIV-1-infected patients produce a vast and potent HIV-1 neutralizing antibodies (bNAbS) that, when administered in other hosts like mice or macaques, produce excellent results regarding treatment or prevention of infection. Moreover, many bNAbS typically fail to neutralize coexisting autologous viruses due to antibody-mediated selection against sensitive viral strains [71-73].

For instance, PGDM1400 and CAP256-VRC26.25, which are V2-specific antibodies, have exhibited impeccable neutralization against HIV-1 [74]. Moreover, a high neutralization effect is shown against nonoverlapping sites on the HIV-1 envelope as a CD4-binding site [56] and (V3) region is targeted by BG18 antibody [61] [77]. Besides, there have been promising outcomes of neutralizing
antibodies targeting the CD4 binding site and V3 region in preclinical studies, and they showed excellent results protecting rhesus macaques against simian-human immunodeficiency virus (SHIV) [78, 79].

The activation of bNAbS toward the membrane-proximal external region (MPER) has exhibited excellent neutralization breadth. The MPER is a highly conserved region within the type-1 HIV-1 glycoprotein envelope (ENV), which is naturally targeted by bNAbS and aids in membrane fusion [80].

Furthermore, the neutralization breadth of bNAbS happens when Fc domain of the antibody that binds to FcγRs of host cell contributes substantially to block viral entry, suppress viremia, and enhance the protection of host cell in vivo. These results expose the crucial role of Fc effector in the activity of HIV-1 bNAbS in vivo and provide strategies for developing bNAbS with improved efficacy [81, 82]. Most of the newly discovered HIV mAbS is tenfold more potent than the previous ones. PG9, PG16, and VRC01 bNAbS are 100-fold more potent than previous HIV mAbS [70, 71].

Among many studies that have been reported, promising clinical outcomes using antibodies with broad and potent activity in neutralizing HIV [83, 84] such as the anti-CD20 mAb, Rituximab that induces killing of the infected cell, which leads to reduce the pool of HIV-expressing cells when combined with latency reversal agents [85].

**NAbs against Ebola virus (EBOV)**

Ebola virus is a Filoviridae that causes severe hemorrhagic fever in humans with enormous public health concerns due to its high transmission and virulence [86]. Since its discovery, it has infected people from various parts of the world, with recent outbreaks in parts of West Africa. Ebola virus is currently one of the most deadly infectious agents in both non-human primates (NHPs) and humans [87]. The main target for the NAbs is EBOV is their glycoprotein (GP), and the need to eradicate this deadly virus in West Africa led to the development of the Ebola vaccine, which is currently under clinical trials. Two approaches were adopted to generate a vaccine:

(i) A combination of modified vaccinia virus Ankara (MVA-BN-Filo) and recombinant chimpanzee adenovirus (rChAd/EBOV) vaccine induce the immune system toward viruses, and 100% of macaques were protected against the Ebola infection following vaccination [88].

(ii) The expression of EBOV GP using a vesicular stomatitis virus also activates the production of antibodies. Here, non-human primates were protected against the virus after vaccination [89].

Among antiviral therapies in use presently to target filoviruses, NAbs seem to demonstrate the highest efficacy as a protective agent [90]. Study shows that the combination of three NAbs (1H3, 2G4, and 4G7) directly targeting the GP of the Ebola virus in cynomolgus macaques leads to complete survival with no visible side effects or perhaps full recovery [91, 92].

Unfortunately, until now, there is no licensed vaccine or any biological treatments available for EBOV in human use. However, in ongoing trials, mAbS targeting EBOV show great supportive care [93–95]. Recombinant vesicular stomatitis virus–Zaire Ebola virus (rVSV-ZEBOV) is an advanced Ebola virus vaccine candidate that was used as supportive care to fight the outbreak of Ebola virus disease in the Democratic Republic of the Congo [96].

ZMapp has been given to participants in a randomized control trial during the 2014 west Africa EBOV outbreak [97], while mAb114 is a monoclonal antibody that has been isolated from human survivor and successful targets (conserve epitopes in receptor-binding domain region of Ebola virus glycoprotein, which has shown promising results in the treatment of an Ebola outbreak). In the first human phase 1 study, mAb114 found to be safe and well-tolerated, which has been approved to use for an expanded access protocol of patients infected with the Ebola virus by the Democratic Republic of the Congo [98].

**NAbs against Chikungunya virus (CHIKV)**

Chikungunya virus is a type of mosquito-transmitted alphavirus that can cause acute or chronic polyarthritis and foot swelling [99]. CHIKV has genetically conserved envelope GPs (E1, E2, and E3). In acidified endosomes, low pH triggers conformational change, which allows the envelope GPs to engage in membrane fusion [100].

Antibodies block the entry of CHIKV to host cells by binding to Fcγ receptors of the host cell or binding to desired epitopes in CHIKV GPs. Previous trials in mice strain lacking Fcγ receptors with engineered antibody confirmed the therapeutic effects of antibodies on the expression of Fcγ receptors [101]. Using anti-CHIKV mAbS showed that the Fc region is critical for CHIKV infection during the acute phase of the disease and required for optimal protection against foot swelling. Combination therapy of two anti-CHIKV neutralizing mAbS [CHK-166 (anti-E1) and CHK-152 (anti-E2)] improved survival compared with single therapy of mAbS [102]. MAbS (5 M16, CHK-152) prevent the activation of membrane fusion of CHIKV through inhibition of conformational changes [103, 104]. Ultrapotent neutralizing human monoclonal antibodies (mAbS) have been isolated from the B cells of a survivor of natural CHIKV infection and encoded into mRNA vector. It has shown biologically significant levels in vivo after infusion of mRNAs in lipid nanoparticles [105].
NAbs against Middle East respiratory syndrome coronavirus (MERS-CoV)

>Middle east respiratory syndrome coronavirus is a type of zoonotic viral pathogen that can cause fatal acute respiratory infections [106].

receptor-binding domain (RBD) of envelope glycoprotein binds to the cell surface receptor dipeptidyl peptidase 4 (DPP4), which enables it to target host cells [107]. Using a combination of neutralizing antibodies like MERS-27 and MERS-4 synergistically increases the efficacy of neutralization against pseudotyped MERS-CoV [108]. This process reduces the viral infection in the lungs (90) [109]. Up to date, there is no licensed and approved treatment for MERS-CoV infections in humans. On the contrary, NAbs such as M336, LCA60, REGN3051, 311B-N1, and REGN3048 target MERS-CoV in transgenic mice and have exhibited tremendous preventive and therapeutic efficacy [110–112].

NAds against severe acute respiratory syndrome (SARS)

Severe acute respiratory syndrome (SARS) belongs to the Coronavirus genus in the Coronavirus family with positive-sense RNA genomes. It is a type of zoonotic viral pathogen that causes acute respiratory infections [107, 113, 114].

Spike (S) protein of (SARS) is the main target for antibodies’ neutralization [115, 116], which has two subunits N-terminal domain (NTD) and C-terminal domain (C-domain). Depends on the virus subtype, either NTD or C domain can serve as the receptor-binding domain (RBD) [117]. Recent studies confirmed that human angiotensin-converting enzyme 2 (hACE2) is the main target receptor for the treatment of SARS infection [118].

80R immunoglobulin G1 (IgG1), a human mAb acts against (SARS-CoV) spike (S) protein, has led to inhibition of viral entry in vitro and was investigated in vivo in a mouse model which showed promising results as immunoprophylaxis agent [119]. Moreover, a combination of human mAbs (CR3014 and CR3022) have potentially neutralized SARS-CoV by targeting different epitopes on the receptor-binding domain of (S)protein [120].

To date, there is no licensed and approved treatment for SARS-CoV infections in humans. While recent studies confirmed (LCA60 mAb) inhibits binding to (hACE2) receptor, it has the potential to be active against SARS-CoV [121].

NAds against severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)

Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is a type of zoonotic pathogen belong Coronaviridae family (non-segmented positive-sense RNA viruses) has led to cause a newly emerged Coronavirus disease 2019 (COVID-19) with mild to moderate symptoms of respiratory disease [122–124].

The spike glycoprotein (S) of coronavirus (CoVs) is the main target for antibody neutralization, which has two subunits N-terminal domain (NTD) and C-terminal domain (C-domain). Depends on the virus subtype, either NTD or C domain can serve as the receptor-binding domain (RBD) [117] [125]. In the case of MERS-CoV and SARS-CoV subtype viruses, the receptors are human dipeptidyl peptidase 4 (hDPP4) [126] and human angiotensin-converting enzyme 2 (hACE2) [118], respectively. Recent studies confirmed SARS-CoV2 uses the same cell entry receptor—angiotensin-converting enzyme II (ACE2)—as SARS-CoV [127, 128]; therefore, RBD in the S1 subunit of S protein contains the dominant neutralizing epitopes for inducing neutralizing antibodies [117, 129]. Moreover, the neutralizing mAbs against SARS-CoV RBD can be evaluated for significant efficacy against SARS-CoV-2 [130]. Neutralizing antibodies have potential activity against the receptor-binding domain (RBD) in spike protein, or that can bind to the ACE2 receptor could show great results toward SARS-CoV-2 [131, 132].

Neutralizing mAbs against pro-inflammatory cytokines (IL-1, IL-17, IL-6) may show significant effect against SARS-CoV2 [133], as one clinical trial (ChiCTR2000029765), using tocilizumab (mAb), reported quick control of COVID-19 symptoms in 21 patients in Anhui, China [134].

COVID-19 infection has led to a global outbreak with morbidity and mortality. Although new investment in biomedical and pharmaceutical researches under ongoing development, using antiviral drug repurposing and other supportive therapies have shown promising results [135]. Up to date, either effective vaccines or neutralizing antibodies have been approved to target SARS-CoV-2. Currently, polyclonal antibodies isolated from recovered SARS-CoV-2-infected patients have been used to treat SARS-CoV-2 infection, but no specific neutralizing mAbs for SARS-CoV-2 have been reported [136]. Markedly, studies confirm RBD-specific human neutralizing mAb for SARS-CoV, CR3022 could bind to SARS-CoV-2 RBD with high affinity [137, 138].

Furthermore, cocktails of RBD-specific antibodies and other regions in the S protein may further improve the potency of nAbs against SARS-CoV-2 and its escape-mutant strains. As well, scientists have tried human sera from convalescent patients to treat COVID-19 [136]. Recent research shows that the N terminal domain (NTD) of the S protein could also be a promising target for therapeutic mAbs against COVID-19 [139].

Limitation of neutralizing antiviral antibodies

Although mAbs might have remarkable therapeutic effects, their strong influence on public health would be attained by the complete eradication of viruses, which can be achieved by the development of vaccines that can produce antibodies to
target specific epitopes on viruses [140–142]. Neutralizing antibodies (NAbs) are the foundation for controlling many infections and prevention as the most established vaccines. However, in the case of some viruses that can cause chronic infections such as HIV or HCV, the production of broad NAbs from recombinant antibody vaccines has still been elusive [17].

The development of mAbs as anti-infectious agents has been delayed due to scientific challenges, clinical trials, and market challenges. These challenges include viral variability and antigenic escape, shortage of treatable viral infections, high cost for production of recombinant antibodies, and competition with effective antiviral vaccines entering the market that could rapidly replace the need for passively infused mAbs. Another potential obstacle is the relatively short duration of viral illnesses, which makes them less attractive commercial targets than diseases such as cancer or chronic inflammatory or autoimmune diseases that require longer mAb treatment cycles. Also, it is expected that antiviral mAbs will not only be used but in combination with other drugs that inhibit post-entry steps of the virus life cycle [24, 30].

The outstanding ability to separate and purify antigen and quickly determine the structural and functional effect of antibodies to target specific epitopes on viral envelope GP has dramatically increased our knowledge of the site of action of viruses.

Through many mechanisms, antibodies can protect host cells from viral infections and, moreover, alleviate the clinical symptoms and attenuate viral progression. Antiviral antibodies mentioned in this review confirm their great potential as preventive and therapeutic agents against many viruses. Using various antibodies characterization approaches discussed here might lead to the more efficient production of neutralizing antibodies to continue providing crucial insights into how we can finally develop immunotherapeutic against pathogens that have thus far evaded our efforts. More precisely, the combination therapy of antiviral antibodies would expect to have broad and potent neutralizing activity against wild-type viruses. (Tables 1 and 2).

**Conclusion**

For successful infection control, virus-mediated antibody responses can promote long life surveillance and better protection from life-threatening infections.

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Compliance with ethical standards

Conflict of interest All authors declare no competing interests.

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