## SARS-CoV-2: REALITY AND CHALLENGES IN TREATMENT AND VACCINE DEVELOPMENT, WHERE DO WE STAND?

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**Abstract:** The newly emerged SARS-CoV-2 has been a threat to the health of humans throughout the world. It causes mild to severe respiratory illness, mainly characterized by fever, cough, loss of taste and smell sensation and other respiratory symptoms. As this disease has been causing unsolicited deaths, antiviral treatment and vaccines are being continuously developed and clinically tested for their effectiveness against SARS-CoV-2. A wide variety of treatment options and vaccine candidates were observed to be tested in different stages of development in different durations. Although progress has been recorded from time to time, a proper treatment and vaccine acceptable for universal use has yet to be developed due to certain constraints. This review unfolds the prevailing treatment and vaccine developing them in a short period of time.

Keywords: SARS-CoV-2, treatment, vaccine, potential, challenges.

#### Introduction

The coronavirus disease 19, COVID-19 or severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), is a rapidly emerging infectious viral disease and considered a health emergency worldwide. The outbreak first emerged in Wuhan, China in 2019, and was suspected to have developed from bat viruses, with the most common clinical features being fever and other respiratory illnesses and most severely death (Mackenzie & Smith, 2020). SARS-CoV-2 rapidly spread across geographical ranges, by human to human transmission through respiratory droplets or fluids from coughing and sneezing of infected humans (Riou & Althaus, 2020, Unhale et al., 2020). Due to effective mutation of the virus itself, the virus is known to be extremely contagious and can infect several people at a time (Unhale et al., 2020). At the time of writing, there are more than five million SARS-CoV-2 cases and more than 300,000 deaths worldwide (World Health Organization, 2020a). These large numbers had forced many countries to implement complete and partial lockdowns, triggering economic recessions and interrupting daily work-related human activities.

To fight SARS-CoV-2, efforts to introduce effective preventive and therapeutic approaches have been taking place. Antiviral treatments inhibit the reproduction of viruses. However, antiviral treatment needs to be constantly upgraded, as viruses mutates and develop resistance towards existing drug (The National Academies, 2020). In contrast, vaccines function by imitating an infection in the body to develop immunity that produce antibodies and T-lymphocytes that will help prevent further spread of the infection (Centres for Disease Control and Prevention, 2018). In the current health scenario, treatment and vaccine development are needed immediately for recovery and prevention, respectively. In this case, several inhibitors have been highlighted to mainly inhibit replication and transcription of the SARS-CoV-2 virus through drug repurposing and usage of potential drugs (Bhardwaj et al., 2020a, Senger et al., 2020). However, all strategies are still at basic levels of development, making it time-consuming and costly. As of 28th May 2020, there are more than 2,751 globally active on-going projects on combatting SARS-CoV-2 in 132 countries, estimated to involve

about 277 billion dollars (World Bank, 2020). While treatments and vaccines are under progressive development, aggressive isolation is the best temporary prevention measure that can be taken to reduce transmission of SARS-CoV-2.

This review comprises the recent advances in therapeutic and prevention efforts by researchers worldwide. This review will be valuable to provide sufficient knowledge and insights on the molecular information of SARS-CoV-2, current status and challenges faced in treatment and vaccine development against the novel coronavirus.

#### Development of Treatments Targeting Structural and Non-structural Proteins of SARS-CoV-2

Drugs that are used in the treatment of SARS-CoV-2 infected patients normally work by targeting protein structures of SARS-CoV-2. SARS-CoV-2 is known to exist as a single stranded RNA, belonging to the betaCoVs category, round or elliptic, with a diameter of 60 - 140 nm and often seen in a pleomorphic form (Cascella et al., 2020). The genetic fragments of this virus express structural and non-structural proteins. Major structural protein targets include the spike (S) protein, membrane (M) protein, envelope (E) protein, and the nucleocapsid (N) protein along with other structural proteins such as hemagglutinin esterase (HE) glycoprotein (Hilgenfeld, 2014). Non-structural proteins, on the other hand, arise from the processing of polyproteins (PP1a and PP1ab) (McBride et al., 2014). The S protein is a structural protein that protrudes from the viral surface of the coronavirus, making the virus appear crownlike (Walls et al., 2020, Coutard et al., 2020). S protein immensely important in aiding the process of viral entry through the angiosteninconverting enzyme 2 (ACE2) host receptor. The development of many antiviral drugs is targeting this mechanism due to its promise as a drug target point (Liu et al., 2020). In regard to S protein, Nafamostat, also used as a treatment for pancreatitis, is currently under clinical

investigation to function as a therapeutic agent against SARS-CoV-2 (Wondmkun *et al.*, 2020).

The N protein is made up of three intrinsically disordered regions (IDR), consisting of the arm, central linker and tail domains (Chang et al., 2016). N protein is mainly responsible for preventing the proliferation of viral offspring, making it another highly regarded drug target after the S protein (Satarker et al., 2020). The M protein had been indicated to be present in two forms, one short and another long, where the long form is usually known to build S proteins (Neuman et al., 2011). The M protein together with the N protein is usually responsible for the maintenance of the viral envelope and also functions in the formation and release of viruslike particles (Schoeman et al., 2019). Among all other proteins present, this protein is basically present in the most generous amount (EA et al., 2019). Next, the E protein is comparatively the smallest protein, consisting of a hydrophobic domain and a charged cytoplasmic tail (Kuo et al., 2007, Venkatagopalan et al., 2015). The E protein focuses on viral morphogenesis, during viral assembly particularly and budding formation, as well as is responsible for determining the virulence factor of a virus (Venkatagopalan et al., 2015). Several antivirals, antibacterial, antitumor and anti-inflammatory drugs have been tested against SARS-CoV-2 as they are known to have a relatively good level of compatibility to the protein targets of N, M and E, potentially known to enable viral replication (Wondmkun et al., 2020). Finally, although not heavily studied, hemagglutinin esterase (HE) glycoprotein has the ability to act as lectins or also known as receptor destroying enzymes (Zeng et al., 2008). However, the effectiveness of targeting this protein is disputed as it has several issues related to toxicity and specificity when used as drug targets (Frick et al., 2006).

Polyproteins, on the other hand, appear as non-structural proteins that are usually produced by open reading frame (ORF)1 $\beta$ (Gildenhuys, 2020). Each polyprotein plays different roles and functions through separate actions of the binding process (Yoshimoto *et*  al., 2020). Among all existing polyproteins, the four most considered targets for drug discovery are the 3-chymotrypsin-like protease (3CLpro), RNA-dependent RNA polymerase (RdRp), papain-like protease (PLpro), and helicase, due to the presence of a clear enzyme active site and important biological functions (Subissi et al., 2014). At this point, several repurposed drugs had been reported to be under preclinical and clinical studies, targeting polyproteins, to test their effectiveness against SARS-CoV-2 (Wondmkun et al., 2020). These proteins have also been reported to be the most focussed upon, currently and in future, for the development of therapeutics against SARS-CoV-2 (Wondmkun et al., 2020). Collectively, all protein targets discussed have great potential to aid in the process of drug development against SARS-CoV-2, however, more trials are still required to identify the best outcome, either from a repurposed or a newly developed drug.

#### Potential Antiviral Treatment for SARS-CoV-2

#### Potential Drugs: Classification and Mechanisms of Action

Many individuals infected with SARS-CoV-2 have died due to the absence of antiviral drugs. Hence, a globally acceptable drug must be identified while waiting for a permanent preventive solution in the form of vaccine. Solidarity clinical trials (testing of drugs on an international scale) are reported to have been initiated by the World Health Organization (WHO) to test the effectiveness of potential drugs among patients from multiple countries (World Health Organization, 2020b). The drugs recommended by WHO had also been included as a part of the European Discovery Project in their initial clinical trials (Eynde, 2020b). The attempt of a single randomized trials had been done on drugs such as Remdesivir, Lopinavir/Ritonavir, Lopinavir/Ritonavir with Interferon beta-1 $\beta$  (IFN- $\beta$ ) and Hydroxychloroquine/Chloroquine (HCQ/CQ) as treatment options (World Health Organization, 2020b).

Malaysia had been reported to be one of the countries to be included in this clinical trial (New Straits Times, 2020). The trial in Malaysia will be aided by the Ministry of Health (MOH), Institute for Clinical Research (ICR) and Clinical Research Malaysia (CRM) (World Health Organization, 2020c). These drugs can be divided into several inhibitor categories based on their mechanism of actions. Remdesivir is observed to target viral replication while Lopinavir/Ritonavir are seen to target viral proteases. HCQ/CQ, on the other hand, work by targeting the viral entry of SARS-CoV-2 virus for inhibition. These drugs have distinctive functions as inhibitors, where the main goal is to eliminate the virus from the human body at cellular level. Examples of other drug options that are FDA approved are included for further information. The general classification of the SARS-CoV-2 drugs can be seen in Table 1.

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Strategy/Inhibitor Categories	Example of Drugs	Target	Antiviral Strategy	Mechanism of Action	Reference
Polymerase inhibitor	Remdesivir* (investigational)	ADA D	Viral	Interferes and hinders viral replication	Sanders <i>et al.</i> , 2020 Wang <i>et al.</i> , 2020a Martinez, 2020
(nucleoside analogue)	Favipiravir (still under investigation)	dama	replication	Reduces the reproduction of RdRp by binding to it	Jeong <i>et al.</i> , 2020 Cai <i>et al.</i> , 2020
Protease inhibitor	Lopinavir/Ritonavir*	Mbro	Viral	May inhibit SARS-CoV-2 protease and interfere with	Harrison <i>et al.</i> , 2020 Sanders <i>et al.</i> , 2020 Lim <i>et al</i> , 2020
	Darunavir	- 1	proteases	virus replication	Jeong <i>et al.</i> , 2020 Dong <i>et al.</i> , 2020
Antimalaria, antiviral &	Chloroquine*		-	Interferes with the glycosylation process	Wang <i>et al.</i> , 2020a Gautret <i>et al.</i> , 2020
antiinflammatory	Hydroxycloroquine*	VITUS/CEII IUSION	viral entry	involving cellular receptors of SARS-CoV-2 and increases endosomal pH	Gao <i>et al.</i> , 2020a Yao <i>et al.</i> , 2020
Antiviral & antinflammatory	Cepharanthie, selamectin, and mefloquine hydrochloride (still under investigation)	Virus/cell fusion	Viral entry	Showed significant reduction of cytopathic effects related to SARS- CoV-2, successfully decreasing the viral load	Jeong <i>et al.</i> , 2020 Fan <i>et al.</i> , 2020a
Transmembrane protease, serine 2 (TMPRSS2) inhibitor	Camostat mesylate	Transmembrane Serine Protease 2 (TMPRSS2)	Viral entry	Interferes and hinders viral entry	Hoffmann <i>et al.</i> , 2020 Sanders <i>et al.</i> , 2020
Antiparasite	Ivermectin	Non-structural 3 helicase (NS3 helicase)	Viral replication	Inhibits replication of SARS-CoV-2, in vitro	Sharun <i>et al</i> ., 2020 Caly <i>et al</i> ., 2020
Rheumatoid Arthritis (RA) drug, AP2-associated protein kinase 1 (AAK1) inhibitor	Baricitinib	Cytokines	Viral entry	Interferes viral entry through the inhibition of endocytosis regulators	Zhang <i>et al.</i> , 2020 Richardson <i>et al.</i> , 2020
*representing main drugs di	scussed in this manuscript				

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These drugs act as inhibitors at different stages of viral penetration. Focussing on the main drugs discussed, HCQ/CQ can act in many modes of action to treat SARS-CoV-2, at an intracellular level. At its most initial stage, HCQ/ CQ drugs are able to hinder ACE2 terminal glycosylation, which inhibits the entry of SARS-CoV-2 virus (Vincent et al., 2005). Next, HCQ/ CQ can disable viral replication by providing a non-suitable condition (increase in endosomal pH) needed for host-virus cell infusion through endocytosis and annulment of virus-endosome fusion (Sharma et al., 2020, Eynde, 2020). Apart of that, HCQ/CQ are also capable of restricting post-translational modification of proteins that may lead to the inhibition of the SARS-CoV-2 virion (Uzunova et al., 2020). In addition, HCQ/ CQ may also be able to prevent SARS-CoV-2 intracellular polymerization and lower the level of cytokines in SARS-CoV-2 infected patients (Sharma et al., 2020). Inhibition by blocking the sialic acid receptors is also another mode of action proposed through the use of HCQ/CQ (Satarker et al., 2020). Contrarily, Remdesivir functions by targeting the activity of viral RdRp while the drugs Lopinavir/Ritonavir may result in the production of immature virus particles, however, more studies on Lopinavir/Ritonavir may be needed to confirm its effectiveness in preventing the infection of SARS-CoV-2 virus in humans (Gordon et al., 2020, Weller et al., 2001). The schematic representation of mechanism of action for all treatment drugs discussed can be seen in Figure 1 and Figure 2.



Figure 1: Schematic representation of mechanism of action for Remdesivir and Lopinavir/Ritonavir against SARS-CoV-2 (Uzunova *et al.*, 2020)



Figure 2: Schematic representation of mechanism of action for chloroquine/hydroxychloroquine against SARS-CoV-2 (Adapted from Cortegiani *et al.*, 2020)

#### **Current Status of Potential Drugs under Trials**

Currently, more than 200 drugs and drug combinations are being tested on patients and results based on the best form of evidence has been recorded (US National Library of Medicine, 2020a). Among the vast number of drug trials, information on the drugs prioritized by WHO in the treatment of SARS-CoV-2 have been focussed in this review. The Remdesivir is a repurposed therapeutic antiviral drug that has been recommended by experts through a wide range of analysis. Remdesivir has been observed to be potentially effective to inhibit SARS-CoV-2 with almost no or minimum adverse effects (Wang et al., 2020b, Holshue et al., 2020, Choy et al., 2020). Several successful attempts on inhibiting Middle East respiratory syndrome (MERS-CoV) and severe acute respiratory syndrome coronavirus (SARS-CoV) shows the potential of remdesivir to be effective on SARS-CoV-2 as well (Sheahan et al., 2020, Wang et

*al.*, 2020b, Agostini *et al.*, 2018). Remdesivir was also nominated as a preferential drug for clinical treatment to be tested as a trusted antiviral therapy option (Alexander *et al.*, 2020). Although assumed potentially successful, Remdesivir has to be further tested in terms of safety and efficacy before declaring it an official cure. The first clinical trial of Remdesivir was reported a failure, however, a final conclusion regarding the effectiveness of this drug had not been made due to arising debates among researchers on its trial procedures (New Straits Times, 2020).

The therapy of Lopinavir/Ritonavir is also being considered as reduced mortality, shortened ICU and hospital stay were seen during trials, and chest CT scans showed the minimum of promise for it to be used in treating patients with SARS-CoV-2 (Deng *et al.*, 2020, Dorward J. & Gbinigie K. 2020, Ampel., 2020). A combination of the therapy with Arbidol had also been suggested to yield better results (Deng *et al.*, 2020). This drug, however, was perceived as a negatively inclined treatment as it also produced insignificant results with almost no difference even in primary outcomes (Cao *et al.*, 2020a, Cao *et al.*, 2020b). It was further suggested that Lopinavir/Ritonavir was to be used within ten days of symptoms appearance (Yan *et al.*, 2020). It was also reported that gastrointestinal adverse effects were observed when the drug was used on patients (Cao *et al.*, 2020b). Hence, the status of Lopinavir/Ritonavir as a potential successful treatment is also still debatable.

The drug combination of Lopinavir/ Ritonavir with IFN- $\alpha$  and Lopinavir/Ritonavir with IFN- $\beta$ are used to treat relapsing multiple sclerosis and previous SARS-CoV and MERS-CoV respectively: a. No studies were however reported to have been done on these drug combinations.

Being an antimalarial drug, many studies have shown positive viral efficacy and safety during trials on Chloroquine (Cheepsattayakorn A. & Cheepsattayakorn R, 2020, Gao*etal.*, 2020b, Wang *et al.*, 2020b, Colson *et al.*, 2020). The use of Chloroquine was nevertheless questioned for its credibility of treating critical ICU patients of SARS-CoV-2 based on least convincing clinical data, antiviral abilities and trials (Taccone *et al.*, 2020). Apart from that, studies have shown improvement in the condition of SARS-CoV-2 patients administered with hydroxychloroquine (Chen *et al.*, 2020, Lane *et al.*, 2020, Tang *et al.*, 2020). However, a study in China opposed the usage of hydroxychloroquine as it might not help in curing SARS-CoV-2 patients and may cause more severe adverse effects in return (Frellick, 2020). Thus, the credibility of hydroxychloroquine as a therapeutic drug is also still unknown until a proper trial is done.

Nevertheless, several other studies also recommended a range of drugs for the treatment of SARS-CoV-2. The first review on the therapeutics used on SARS-CoV-2 patients suggested that studies on corticosteroid therapy was the most focused followed by studies on Lopinavir and then Oseltamivir (Tobaiqy et al., 2020). Other drugs for treatment include the antiviral EIDD-2801, Japan flu drug, Tocilizumab, and Losartan (LiveScience, 2020a). All the drugs tested on SARS-CoV-2 are still in trial stages and have produced many pros and cons to be considered for efficient use. A combination of antiviral drugs as treatment, such as anticoagulant therapy with heparin and statins with angiotensin II type I receptor blockers (ARB) had also been suggested (Tang et al., 2020, Fedson et al., 2020). No treatment has been finalized to be able to treat SARS-CoV-2 patients. The summary of information on prioritized drugs can be seen in Table 2.

	Table 2: Recomm	ended drugs f	or the treatment of SARS-CoV-2 (V	Vorld Health Organization, 2020)
Recommended Drug	Description	Previous Usage	Recommended Dosage for SARS-CoV-2	Status of Clinical Development of SARS-CoV-2
			CT NCT04252664: 200 mg loading dose on day one is	<u>Wang M. et al. (2020b)</u> Remdesivir was found effective at a post virus entry stage and inhibited SARS-CoV-2 infection in Vero E6 cells.
Remdesivir	Nucleoside inhibitor	Ebola	given, followed by 100 mg iv once-daily maintenance doses for nine days. CT NCT04257656: 200 mg	<u>Holshue, M. L. <i>et al.</i> (2020)</u> No adverse events and improved patient's condition were seen to have occurred after administration in patients in the United States.
			loading dose on day one is given, followed by 100 mg iv once-daily maintenance doses for nine days.	Choy <i>et al.</i> (2020) Combinational therapy of 6.25 $\mu$ M remdesivir and 0.195 $\mu$ M emetine may achieve 64.9% inhibition in the SARS-CoV-2 viral field.
				<u>Deng <i>et al.</i> 2020</u> Improved chest CT scan in patients after seven days of administration. The combination of Lopinavir and Ritonavir with Arbidol would produce better results for SARS-CoV-2.
Lopinavir/ Ritonavir	Protease inhibitor	HIV infection	500mg once, twice a day, 2 weeks	Dorward J. & Gbinigie K. (2020) A small clinical improvement in terms of reduced mortality of 28 days, shortened ICU stays and shortened time from hospital by one day.
				<u>Cao et al. (2020a)</u> Lower 28 days mortality rate observed compared to pre-trial, however not significant.
Lopinavir/	Lopinavir and Ritonavir	SADS &	IFN-α 5 million unit or activation does in 2 mL starila	Clinical trial for SARS-CoV-2:
Ritonavir with IFN-β	inhibitors combination protease inhibitor and host	MERS	water via nebulization twice a day	U.S.National Library of Medicine, 2020b Phase IV at https://clinicaltrials.gov/ct2/show/NCT04343768

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				Clinical trial for SARS-CoV-2:
Lopinavir/ Ritonavir with IFN-β	Lopinavir and Ritonavir are antiretroviral protease inhibitors combination protease inhibitor and host	Relapsing multiple sclerosis	For MERS: Lopinavir/ Ritonavir 400mg $+100$ mg / ml twice daily for 14 days and IFN- $\beta$ 0.25 mg subcutaneous every alternate day for 14 days	<u>U.S. National Library of Medicine, 2020b</u> Phase IV at https://clinicaltrials.gov/ct2/show/NCT04343768
				Cheepsattayakorn A. & Cheepsattayakorn R. (2020) Chloroquine was found to block SARS-CoV-2 infection at low-micromolar concentration, with a halfmaximal effective concentration (EC50) of 1.13 μM and a halfcytotoxic concentration (CC50) greater than 100 μM.
Chloroquine	Antimalarial agent, hemepolymerase inhibitor	Malaria prophylaxis and treatment	Adult dose of 500 mg twice per day for no more than 10 days	<ul> <li><u>Gao et al. (2020b)</u></li> <li>Chloroquine phosphate has shown apparent efficacy and acceptable safety against SARS-CoV-2.</li> <li>Recommended to be included in the next version of the Guidelines for the Prevention, Diagnosis, and Treatment of Pneumonia caused by SARS-CoV-2 by the National Health Commission of China.</li> </ul>
				<u>Wang <i>et al.</i> (2020b)</u> Chloroquine has effective antiviral activities (at both pre and post entry stages) along with immune modulating activity. Usage: 6.90 μM in Vero E6 cells.
				Colson <i>et al.</i> (2020) Chloroquine administration showed a significant reduction in the length of stay in hospitals and improvement in the evolution of SARS-CoV-2 pneumonia with a proper concentration of dosage.
				Chen <i>et al.</i> (2020) The use of Hydroxychloroquine significantly reduced body temperature recovery and cough emission time. Improvement on pneumonia condition was also observed in patients in the study.

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<u>Lane et al. (2020)</u> Only short-term Hydroxychloroquine is recommended while the combination with azithromycin may cause severe adverse effects such as heart failure and cardiovascular mortality.	<u>Tang. (2020)</u> More alleviations of clinical symptoms in patients without any antiviral treatment was observed with slightly increased adverse effects, with no increase of any serious adverse effects.	Mitja (2020) Approved doses are able to possibly prevent the infection of SARS-CoV-2 and mitigate viral shedding. Positive preliminary outcomes have been obtained.
	400mg per day for 5 days/200mg thrice daily	
	Malaria, lupus and rheumatoid arthritis	
	Antimalarial agent, hemepolymerase inhibitor	
	Hydro- xychloroquine	

### Alternative Treatment Methods against SARS-CoV-2

Recently, researchers have proposed convalescent plasma therapy for patients severelv affected by SARS-CoV-2. Convalescent plasma therapy was observed to increase or maintain neutralizing antibodies to work against the SARS-CoV-2 virus, reduce clinical symptoms in about three days along with a positive but varying rate of absorption of lung lesions seen through radiological examination (Duan et al., 2020). Additionally, this therapy showed no distinct adverse effects and also showed reactions that give rise to the elimination of SARS-CoV-2 virus in patients tested with throat swab (Ye et al., 2020). Several patients at different levels of infection were also observed to improve after treatment with this therapy, leading to them being discharged from hospital (Shen et al., 2020). Early administration of this therapy has been said to promote higher effectiveness level in eliminating the virus (El-Aziz & Stockand, 2020). This discovery can be considered to have the most potential so far, however, more clinical trials involving a variety of patients can determine the ultimate outcome.

Apart from that, research on bioactive molecules extracted from natural molecules and products, as well as natural lead compounds, have also shown positive contribution as potential inhibitors against the SARS-CoV-2 virus. Findings on bioactive molecules function as an alternative to the limited therapeutics available against SARS-CoV-2 and might be proven effective following further clinical trials and testing. These bioactive molecules are known to work against the main protease (Mpro), one of the drug targets of SARS-CoV-2. Out of 65 potential bioactive molecules, Oolonghomobisflavan-A, extracted from the tea plant (Camellia sinensis) had been highlighted as a potential lead molecule that can inhibit the Mpro of the SARS-CoV-2 virus due to its good number of hydrogen bonds and higher Mechanic/Poisson-Boltzmann Surface Area (MM-PBSA) binding energy with Mpro (Bhardwaj et al., 2020b). In another study, six

out of ten polyphenols, namely broussochalcone A, papyriflavonol A, 3'-(3-methylbut-2-enyl)-3',4',7-trihydroxyflavane, broussoflavan A, kazinol F and kazinol J, extracted from *Broussonetia papyrifera* have exhibited stronger binding ability with Mpro, thus recommended as potential inhibitors against SARS-CoV-2 (Ghosh *et al.*, 2020).

Similarly, using the in-silico approach on 15 flavonoid based phytochemicals, researchers have suggested three polyphenols, namely, rutin (readily available as a drug), isorhamnetin-3-O-β-D, calendoflaside, derived from Calendula officinalis, to be effective inhibitors of the Mpro through their assessment on binding affinity, stability and flexibility of the phytochemicals with the Mpro of SARS-CoV-2 virus (Das et al., 2020). Furthermore, polyphenols, such as EGCG, theaflavin-3'-O-gallate (TF2a), theaflavin-3'gallate (TF2b), theaflavin 3,3'-digallate (TF3), which have been widely used for therapeutic effects for several diseases, have been tested effective based on their strong ability to bind to the RNA-dependent RNA polymerase (RdRp) of SARS-CoV-2 and aid in the inhibition of RdRp (Singh et al., 2020). These findings have been further confirmed by the MM-PBSA calculation and through the analysis of druggability, hence found effective to be used as a therapy against the SARS-CoV-2 virus (Singh et al., 2020). As alternatives, the compounds highlighted above can be potentially successful to combat the SARS-CoV-2 virus, though many further trials would be required to properly test the antiviral properties of polyphenols against SARS CoV-2.

#### Factors Influencing the Effectiveness of Antiviral Treatment

An antiviral drug usually works by aiming for molecular targets in every possible stage of a virus life cycle. Although antiviral treatment has been developed since the 1980s to overcome HIV infection, many more new and advanced drugs are still in the process of development against coronaviruses, dengue, filoviruses and many others (Ryu, 2017, Richman & Nathanson, 2016). An effective antiviral treatment will usually display high rates of safety in the treatment used, low cost and adequate supply of the particular drug (Mitja, 2020). Currently, development of antiviral treatment is being studied for the newly emerged SARS-CoV-2 viral disease, and there are many factors that can influence the effectiveness of a particular antiviral treatment, hence, influencing the rate of availability of the antiviral treatment needed.

The most important factor that influences the effectiveness of an antiviral treatment against SARS-CoV-2 is the safety of the treatment towards patients and the absence of adverse effects post-treatment. Based on suggested antiviral drugs listed above, the trials involving Remdesivir, were related to cause constipation, nausea, diarrhea, vomiting, poor appetite and increased levels of liver enzyme (Fan et al., 2020b). Ribavarin and Lopinavir/ Ritonavir showed adverse effects, such as, reduction of haemoglobin, which causes respiratory distress and gastrointestinal adverse effects, namely, nausea, vomiting and diarrhoea respectively (Martinez, 2020, Jean et al., 2020). Chloroquine was observed to cause effects on hepatic, hematologic and renal systems with a major shortage of antimalarial armamentaria and was not recommended for use, while hydroxychloroquine was not recommended to those with allergy to hydroxychloroquine, pregnant and breastfeeding mothers (Cortegiani et al., 2020b, Gautret et al, 2020). The adverse effects reported currently influence the effectiveness of the antiviral drugs for administration, hindering proper utilization of one or more drugs as the cure for patients infected with SARS-CoV-2. This in return also slows down the process of finalizing the antiviral drug to treat SARS-CoV-2.

The massive demand for antiviral drugs that have been repurposed for the use against SARS-CoV-2 serves as a challenge as complaints have been filed against the shortage of Remdesivir supply in the United States (Ledford, 2020). The distribution capacity of this antiviral drug to every country is also doubted (Bai, 2020). While potential antiviral drugs are already being used by researchers worldwide for clinical trials against SARS-CoV-2, the use of the same drugs to treat existing diseases other than SARS-CoV-2 are still ongoing. This affects the supply of antiviral drugs needed and demand is yet to be met. Apart from that, patients with cerebrovascular diseases and diabetes have been reported to be the ones who usually die even with treatment (Yang *et al.*, 2020). Hence it is also difficult to propose an antiviral drug that is suitable for both readily healthy and ill patients infected with SARS-CoV-2.

Likewise, other factors such as the time lag between diagnosis and antiviral treatment prescribed to treat SARS-CoV-2 patients using repurposed drugs can also affect the effectiveness of the antiviral treatment (Smith et al., 2020). The viral kinetic profile for each virus type varies, and in the case of SARS-CoV-2, it is suggested that the peak is estimated to occur eight to 10 days after infection (Smith et al., 2020). When treatment is not initiated accordingly, the drug may no longer work to eliminate the virus. For example, an influenza infection will have to be treated within three days of its first appearance of symptoms in order to work effectively, where any prolonged duration might reduce the efficiency of the drug used (Aoki et al., 2003). There is a chance of more than 50% of efficacy if treatment is initiated prior to symptoms onset, however, this might also not work on patients with severe illness (Goncalves et al., 2020). This can only be achieved when patients report for swab testing right away upon recognition of symptoms, which might be challenging if the public fail to identify symptoms correctly or intentionally stay away from being tested.

The optimization of dosage used to treat SARS-CoV-2 patients using repurposed drugs plays a role in hindering the effectiveness of a drug as well. For this, it is important to take note and understand the potency of drugs before establishing them to patients (Smith *et al.*, 2020). Hydroxychloroquine was found to be more potent than Chloroquine, suggesting different dosage requirements to be followed as compared to Chloroquine for effective treatment

and recovery (Yao *et al.*, 2020). Similar effort should be taken on the available drugs every now and then to ensure optimized dosage when treating patients with SARS-CoV-2 for better chances of recovery.

The emergence of SARS-CoV-2 virus is extremely new and requires more studies to be carried out in order to completely understand the virus. Thus, the unknown aspects that are yet to be discovered can hinder the process of introducing a proper antiviral treatment to the public as in the demanded period. The development of a treatment against SARS-CoV-2 can be based on the previously identified SARS-CoV virus, however, the differences in some molecular aspects and the extent of its effects towards humans is questionable (Ceccarelli, 2020). Discovering an ultimate treatment might be challenging and cannot be developed too soon; efforts to combat SARS-CoV-2 must be ongoing.

#### Potential Vaccines Development for SARS-CoV-2

Vaccines are being developed to prevent the spread of SARS-CoV-2 to healthier populations across the globe. An international effort using randomized trials of potential vaccine candidates has been introduced to aid an effective, reliable and appropriate development of vaccines with a global standard (World Health Organization, 2020b). Vaccines are also in the process of development partially based on the responses to Ebola and SARS (UNICEF, 2020). Clinical trials on potential candidates of vaccine from different types of vaccine platforms are carried out in several countries around the world. Currently, a total of six candidate vaccines from four vaccine platforms were reported to have entered clinical evaluation while an increased number of 77 candidate vaccines were reported to be in the preclinical evaluation (World Health Organization, 2020d). Trials using new candidate vaccines are emerging rapidly, entering the clinical and preclinical stages respectively. Candidate vaccine platforms consist of non-replicating viral vectors, deoxyribonucleic acid (DNA), ribonucleic

acid (RNA), inactivated, live attenuated virus, protein subunits, replicating viral vectors, viruslike particles (VLP) and also unknown platforms (World Health Organization, 2020d). A total of 83 vaccine candidates nominated from the vaccine platforms listed above had been used for the vaccine development trials. Candidate vaccines from protein subunits platforms were observed to be used the most while live attenuated platforms were least used for trials among all other platforms as seen in Figure 3. Candidate vaccines from non-replicating viral vectors and RNA vaccine platforms were equally nominated for trials. Candidate vaccines among inactivated, VLP and unknown platforms are seen to be used in small numbers probably due to its least suitable properties to prevent the replication of SARS-CoV-2 virus in humans. Candidate vaccines from the remaining DNA and replicating viral vectors platforms are seen to be nominated for trials in moderation, ranging from 10% to 12% of total percentage of usage.

Among all the vaccine candidates used, the six candidate vaccines that have been observed to

enter clinical evaluation stage are from the nonreplicating viral vector, DNA, inactivated and RNA vaccine platforms as seen in Table 3. All candidate vaccines tested in this clinical stage were tested for the development of vaccines for several other diseases except for the inactivated vaccine candidate that has not been tested for any other previously occurred viral diseases. The candidate vaccines are most likely to have varying advantages and tested for a promising outcome. For instance, platforms of mRNA and DNA were reported to offer flexibility such as antigen manipulation and potential speed, whereas viral vector platforms usually have higher levels of protein expression along with durable stability and highly induced immune responses (Le at al., 2020). Adjuvants are also recorded to be aiding in the enhancement of immunogenicity in which, vaccine developers such as GlaxoSmithKline have been working on, to be used along with vaccines developed by researchers (Le at al., 2020). All reported candidate vaccines are evaluated for safety and immunogenicity along with tolerability and persistence.



Figure 3: The percentage of vaccine platforms used for the development of a potential universal vaccine against SARS-CoV-2 (World Health Organization, 2020)

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Non-replicating vectors are usually vaccine platforms that are known to be safe with a long history of usage for gene therapy and are present in multiple serotypes and chimeric forms, where Adenoviruses are considered the most widely utilized vectors for vaccine development (Robert-Guroff, 2007). As such, candidate vaccines mainly used for development against SARS-CoV-2 under the non-replicating vectors platform are the Adenovirus Type 5 Vector and chAdOx1. Adenovirus Type 5 Vector is made from a weakened common cold adenovirus that normally affects chimpanzees, while chAdOx1 is an adenovirus that infects humans, with longer lasting immune responses (U.S Department of Health and Human Services, 2020, LiveScience, 2020b). Currently, both vaccine candidates have been showing satisfying results through the development of neutralizing antibodies and T-cells responses in the majority of their participants with mild adverse reactions. As a result of successful trials from Phase I and Phase II studies, both vaccine candidates have been allowed to proceed to Phase III of clinical stage evaluation (University of Oxford, 2020, Bar-Zeev et al., 2020, Zhu et al., 2020, Folegatti et al., 2020).

The inactivated vaccine platforms on the other hand, are dead pathogen versions as opposed to live, weakened virus vaccines which may not necessarily be as protective as live vaccines and need booster shots in between for continued protection against SARS-CoV-2 (U.S Department of Health and Human Services, 2020). Both vaccines using the inactivated vaccine platform had been also displaying promising results as both vaccines highlighted the development of antibodies against SARS-CoV-2 with no serious adverse effects (LiveScience, 2020b). The inactivated vaccine candidate trial was reported to be further carried out on children and the vaccine had also been claimed to be used by the employees of the developer while the inactivated vaccine with alum as placebo had already begun their Phase III clinical trials in Abu Dhabi (Liu et al., 2020, Barrington et al., 2020).

In contrast to the traditional vaccine development as above, the RNA vaccine platform consists of a first-hand quick and easily manufactured technology based vaccine candidate made up of a genetic material known as messenger RNA (mRNA) that triggers the immune system \ when naturally exposed to the virus, similar to the traditional vaccine candidates above (LiveScience, 2020b). Although the use of these vaccines was earlier opposed as it was considered unsafe and ineffective, the mRNA vaccine candidate had showed positive results and minimal adverse effects when used against SARS-CoV-2 through the development of a strong immune response when tested in rhesus macaque monkeys (Pardi et al., 2018, Jackson et al., 2020). This result, however, varied from the outcome produced through chAdOx1, where rhesus macaque monkeys were only observed to be immune against pneumonia and not SARS-CoV-2 (Corbett et al., 2020). The DNA vaccine platform, known to be the only nucleicacid based vaccine, has notable advantages such as being inexpensive, providing longer immunogenicity, easy development and storage (Alarcon et al., 1999, Sasaki et al., 2003, Khan et al., 2009, Khan, 2013). This vaccine has been reported to produce positive results in its Phase I clinical trial, where antibodies detected displayed negative binding to S protein with the ACE2 receptor of SARS-CoV-2 and will begin its second and third phase of trials once regulatory concurrence is obtained (Smith et al., 2020, Inovio Pharmaceuticals, 2020).

The developers of these vaccines are also seen to be more varied through the involvement small-scale, institutional of developers. Additionally, five out of six candidate vaccines are observed to be used among healthy adults aged 18 to approximately 60 years old. Candidate inactivated virus vaccine was observed to target a wider age range; reportedly from as young as six years old and above. Based on the data given, the candidate Adenovirus Type 5 Vector vaccine has also been in Phase 2 clinical evaluation, that typically focuses on gathering preliminary data on the efficacy of the vaccine on people with underlying medical conditions while the rest

	Table 3: Candidate vacc	ines in clinical e	valuation stage (Worl	ld Health Organizatio	ו & National Library of Medicine א מלו	e, 2020)
Vaccine Platform	Candidate Vaccine	Clinical Evaluation Stage	Study Type/ Allocation	Study Sample/ Sample Size	Developer	Previous Use for Diseases
Non-	Adenovirus Type 5 Vector	Phase II	Interventional/ randomized	Healthy adults aged above 18 years/108 participants	Beijing Institute of Biotechnology	Ebola
reprictating viral vector	chAdOx1	Phase I/II	Interventional/ randomized	Healthy adults aged 18-55 years/1,090 participants	University of Oxford	MERS, Influenza, TB, Chikungunya, Zika, MenB, plague
	Inactivated	Phase I	Interventional/ randomized	Healthy subjects aged 6 years and above/45,000 participants	Beijing Institute of Biological Products	NA
Inactivated	Inactivated + alum	Phase I/II	Interventional/ randomized	Healthy adults aged 18-59 years/600 participants	Sinovac Research and Development Co., Ltd	SARS
DNA	DNA plasmid vaccine (INO-4800) followed by Electroporation	Phase I	Interventional/ non-randomized	Healthy adults aged 18-50 years/120 participants	Inovio Pharmaceuticals	Lassa, Nipah, HIV, Filovirus, HPV, Cancer indications, Zika, Hepatitis
RNA	LNP – encapsulated mRNA	Phase I	Interventional/ non-randomized	Healthy adults aged 18-99 years/120 participants	National Institute of Allergy and Infectious Diseases (NIAID)	Multiple candidates
*NA = Not Avail	lable					

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of them are still in Phase 1 for safety testing among healthy individuals. The development of candidate vaccines to prevent SARS-CoV-2 is still under process as the candidates would have to fulfil safety and efficacy measures and should be able to cover patients under different conditions and health complications.

The development of a vaccine is time consuming and would require much more research to produce a universal vaccine for the entire human population. Nevertheless, it was reported that a genetic engineering method using DNA and RNA that hold instructions for faster vaccine productions compared to the conventional techniques was used (Schmidt et al., 2020). However, the efficiency of this method developed during the MERS outbreak is still unclear. Recently, a pilot study involving a purified inactivated SARS-CoV-2 virus vaccine candidate (PiCoVacc) tested on animal models resulted in complete and partial protection against SARS-CoV-2 (Gao et al., 2020b). Today, as positive as it sounds, there is still no evidence that the same result can be expected from humans, where the vaccine would need more trials to confirm similar or better results without imposing any risk of adverse effects. Hence, in the meantime, the public are advised to adhere to proper prevention measures to stay safe from the infection of SARS-CoV-2.

# Challenges and Time Constraints in Vaccine Development

Although vaccine development being is accelerated, there are many challenges in developing an effective vaccine for everyone. These challenges include improper understanding of immune responses to infection, mediocre marketing abilities especially in poorer countries, insufficient financial facilities, safety and efficacy factors, ineffective patent protection and regulation and the absence of proper health coverage worldwide (Oyston & Robinson, 2012, Kaufmann et al., 2014, Possas et al., 2019). Hence, the need to strategize on the development of improved next-generation vaccines is highly important to overcome existing pandemics (Wei *et al.*, 2020). In addition to the challenges faced, the lack of trust in vaccine safety and the adverse effects that may be caused by vaccination have also hindered active manufacturing and distribution of these vaccines, restricting access to vaccines in many countries (Geoghegan *et al.*, 2020).

As such, vaccine development against SARS-CoV-2 virus is also burdened by several circumstances that require proper consideration before declaring the vaccine to the public. One such example may lie within the antigenic target and vaccine related disease aggravation (Nick et al., 2020). Based on research on SARS-CoV, this is directly associated with concerns over antibody-dependent enhancement (ADE) from the phenomenon of enhanced pathogenesis through the activation of macrophages via the antibody's Fc domain, where ADEs are known to accumulate over time, with increased severity of risk with age (Nick et al., 2020, Jianqing, 2020). Since the SARS-CoV-2 is known to have close homology with SARS-CoV, this aspect can be expected to be a challenge as well (Chan et al., 2020). Apart from that, the development of a vaccine might also engender the enhanced respiratory disease exacerbation (ERD) as reported in the formaldehyde-inactivated vaccine development against respiratory syncytial virus (FI-RSV), if implemented for SARS-CoV-2 (Nick et al., 2020). Vaccine's testing using animal models could also help in enhancing immunity against SARS-CoV-2, however, the unknown immune response and adverse effects that may be caused is the underlying challenge as human physiology and characteristics of immune response is different and more complex than animals (Saif, 2020). It is also observed to be difficult to gather all aspects and elements needed for more intense studies using animal models as only limited laboratories are used to isolate live strains of SARS-CoV-2 (Jianqing, 2020).

Altogether, clinical safety and efficacy of a vaccine towards human health was considered the most difficult challenge to overcome in developing trusted vaccines (Calina *et al.*,

2020). A suitable candidate vaccine to overcome SARS-CoV-2 virus with minimum or almost no severe adverse effects is still difficult. For example, a review that was based on the experience reported from pioneer Chinese experts in handling the virus, suggested that the use of inactivated virus vaccine candidates might leave immunopathological enhancement effects and need long-term and careful deployment (Huaiyu, 2020). This could be used by anti vaxxers to strengthen their beliefs, leaving healthcare professionals with a greater challenge to influencethe public to get vaccinated. This is because not everyone vaccinated can truly be immune to this disease due to the varying genetic and immunological aspects (Colgrove, 2007). Ample of clinical trials and laboratory analysis would be required to produce a vaccine that will be safe for every human being, globally. This would in return, consume more time to develop a proper vaccine. Factors such as age and the time frame or duration of vaccine response can also impact the efficacy of the vaccine (Lord, 2013, Boda et al., 2018).

Apart from that, other important factors, such as the technology used for vaccine development and capacity for large-scale production of vaccines are also challenges (Calina et al., 2020). Technology may have to be updated thus, requiring more trial and error. Furthermore, in a situation where if a vaccine is properly developed, the lack of commerciality in SARS-CoV-2 vaccines would leave manufacturers exhausted because of largescale new vaccine production at the expense of previously produced vaccines needed for other diseases. (Calina et al., 2020). Hence, the development and manufacturing of a vaccine is not easy and needs thorough research and trials due to the long list of challenges. More time to make concrete findings is needed from initial development to the final stage, involving implementation policies.

#### Conclusion

Several combinations of treatment and vaccine approaches are constantly being tested by researchers. Health authorities and researchers dependent on currently available are still therapeutic efforts to help cure infected patients as an effective vaccine is yet to be finalized. However, the efficiency of a treatment varies according to the state of health of an individual, while vaccine development usually takes many approved trials at several levels before rolled out to the public, making it more challenging with the limited time available. This is because SARS-CoV-2 is a novel viral disease that needs more research to be done to effectively overcome the pandemic. While treatment and vaccine development are ongoing, it is important to practice prevention guidelines, such as physical distancing, frequent washing of hands, use of sanitizers, restraining movements such as travelling and avoiding unnecessary close contact. Although challenging, this new norm must be adapted to limit the spread of the disease and if possible, to completely overcome SARS-CoV-2 in the near future.

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