

1 **The epidemiology, evolution, transmission, and therapeutics of COVID-19 Outbreak: an**
2 **update on the status**

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19 Abstracts:

20 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is an etiologic agent of the
21 respiratory disease in humans that is known as coronavirus disease 2019 (COVID19). The first
22 outbreak of the disease was initially documented in Wuhan, Hubei Province, China in late
23 December 2019 where people had experienced SARS pneumonia-like symptoms with unknown
24 etiology. Since then it has been observed that COVID-19 positive patients have been showing
25 mild to severe upper respiratory illness symptoms. The type of virus is known to make its
26 transfer from animals to humans and for the concerned virus; researchers have claimed its origin
27 from bat coronavirus at whole-genome level with a 96 % sequence identity. The COVID-19
28 virus is very contagious and communicable in nature and has been spread throughout the globe
29 since its first outbreak in China. On March 9, 2020, WHO declared it as a Pandemic, and within
30 a month it was already reported to have shown its presence in 213 countries and territories or
31 areas. As of April 29, 2020, this novel virus infected 3,218,183 people and caused 228,029
32 mortalities worldwide with a variable mortality rate from 3-13 % across the planet and also
33 varied by age and gender. Diagnosis of the disease is a key component in understanding and
34 controlling the spread of the virus and several techniques have been devised including RT-PCR,
35 ELISA, and sequencing-based approaches. To cure COVID-19 patients as of now we do not
36 have proven to be a safe and effective treatment. Therapeutic options currently under
37 investigation in various parts of the world. However, there are various effective therapeutic
38 targets to repurpose the present antiviral therapy for developing potential interventions against
39 SARS-CoV-2. Boosting the immune system can also help to prevent and spread of COVID-19
40 using various medication and exercises. In this review, our goal to summarize and discussed the
41 present scientific advancements to fight against this novel pandemic.

42 **Keywords:** COVID-19, Evolution of SARS-CoV-2, replication, emerging disease 2019 and
43 diagnostic tools

44

45 **Introduction:**

46 The first report of Severe Acute Respiratory Syndrome like pneumonia with unknown etiology
47 was observed in late December 2019 in Wuhan, China [1-3]. International Committee on
48 Taxonomy of Viruses (ICTV) named the agent responsible as Severe Acute Respiratory
49 Syndrome Coronavirus 2 (SARS-CoV-2). Later a number of cases of the same have surged and
50 WHO named this new disease as coronavirus disease 2019 [4]. The onset of the initial animal to
51 human transfer has been traced back to the wet market in Wuhan and shortly after that human to
52 human transmission was observed [5]. The researchers have traced its emergence to pangolins,
53 and more precisely to Bat (*Chiroptera*) with 96 % of the genome matched. The category of virus
54 is known to make an inter species transfer to finally find a way to attach to a host cell receptor in
55 humans. SARS-CoV-2 is the 7th coronavirus known to us that has made an infectious transfer
56 from animals to humans [5]. Previously, documented coronaviruses i.e. severe acute respiratory
57 syndrome coronavirus (SARS-CoV) and Middle East Respiratory Syndrome Coronavirus
58 (MERS-CoV) were some that have made similar transfers to humans and then spread within the
59 same species.

60 SARS-CoV-2 is a newly emerged member of a Coronaviridae with a genome size of ~30,000
61 bases. The size of its virion is roughly 70-90 nm in diameter [6]. Coronaviruses are made up of
62 four different structural proteins; spike (S), envelope (E), membrane (M), and nucleocapsid (N)
63 and 16 non structural proteins [7]. The previously known coronaviruses are large, positive-sense

64 and single-stranded RNA (+ssRNA) viruses that are classified into 4 different genera which are
65 alpha coronavirus, beta coronavirus, gamma coronavirus and delta coronavirus. Among these
66 alpha and beta coronavirus are known to infect humans. Infections of these viruses can be fatal
67 in some severe cases. SARS-CoV, MERS-CoV, and SARS-CoV-2 can cause severe upper
68 respiratory illness in humans while 4 other known coronaviruses HCoV 229E, NL63, OC43, and
69 HKU associate with mild symptoms in human [8-11] which globally account for 10-30 % of
70 upper respiratory tract infections specifically in adults. Coronaviruses have different ecological
71 niches with a wide diversity documented in bats [12-17]. Thus, having a higher probability of
72 SARS-CoV-2 evolving from Bat. Further we are addressing following questions:

- 73 A. What are the different modes of transmission of SARS-CoV-2?
- 74 B. How the SARS-CoV-2 originate and what its apparent replication mechanism?
- 75 C. What are the current effective diagnostic tools and therapeutic options are available to
76 control and cure the COVID-19?

77

78 **Transmission:**

79 Transmission of SARS-CoV-2 occurs through different modes; primarily through respiratory
80 droplets of infected person when exhaled, sneezed, or cough, viral particles come out in air with
81 coronavirus droplets. Respiratory droplets emissions are classified into “large” and “small”
82 droplets. Compared to small droplets, large droplets quickly settle down than it evaporate into
83 air, contaminating the nearest environment of the infected person. [18]. Van et al. [19]

84 documented that this novel virus can survive up to 3 hours in air at 65% relative humidity. Other
85 modes of SARS-CoV-2 spreading is through persons if comes in contact with the infected person
86 or with objects which are already used by infected person or he/she is already comes in contact
87 with virus contaminated surfaces and objects like doors knobs, water-tap, bathroom, utensils,
88 groceries stuffs, currency, clothes, papers, *etc.* As it is known that SARS-CoV-2 can survive on
89 plastic and stainless steel up to 72 hours, on copper less than 4 hour, on card board less than 24
90 hours therefore to prevent the virus spreading, avoid the contact with these materials for at least
91 four day or unless it is properly sanitize [19]. The warm climate could be a barrier of
92 transmission of the COVID-19 probably due to the quick loss of humidity in air and it might lead
93 to death of Coronavirus. There is no direct evidence but as per the disease confirmed case
94 indicating that may be good for those countries which have higher temperature like India,
95 Pakistan, Sri Lanka, Australia, Africa, Sri Lanka *etc* because the number of cases are lower in
96 these countries as of April 16, 2020. There are “n” number of possible modes of transmission
97 could be from fruit market, vegetable market, groceries stores, grains market *etc* because we do
98 not know if some people are asymptomatic or have very mild symptoms which are very hard to
99 predict about the COVID-19. As of the current death and worsening of the symptoms suggest
100 that those who are aged or might have some other associated disease such as asthma, diabetes,
101 heart patients or genetics of person *etc.*

102

103 **Pathogenicity:**

104 Virus get entry into the host cells possibly via fusion of spike protein with membrane or through
105 the interaction of spike RBD with either human ACE2 receptor or Glucose Regulated Protein 78

106 (GRP78) receptor present on the type II alveolar cells (AT2) [20-25]. However, SARS-CoV2
107 does not use other receptors that SARS-CoV uses for entry into cells. Once virion gets into the
108 respiratory system, it gets a chance to interact with lung type II alveolar cells receptor ACE2 and
109 is followed by its priming by TMPRSS2. Because lung AT2 cells strongly co-express ACE2 and
110 TMPRSS2 receptors [22, 24] which is the primary root for SARS-CoV-2 entry into host cells.
111 Hamming et al [26] reported the ACE2 abundantly expressed on the type-2 alveolar (AT-2)
112 epithelial cells and small intestine, which is most susceptible for infection of COVID-19. SARS-
113 CoV2 human infection proceeds through stage I to III. In stage I, patients will experience
114 asymptomatic or mild symptoms of dry cough, diarrhea and headache. Whereas stage II, the
115 patient feels shortness of breath with pneumonia-like symptoms, and at stage III, chronic
116 inflammatory response occurs *i.e.* high blood pressure, heart problems or chronic respiratory
117 conditions, cardiac failure [27].

118 The higher expression level of ACE2 is possibly associated with a higher rate of
119 mortality in the older age group or with people who have heart disease due to COVID-19
120 patients. Increased ACE2 expression at protein and transcriptional level has been observed in
121 patients with heart disease therefore such patients were considered as more vulnerable to virus
122 infection and may have severe effect of COVID-19. Therefore, mortalities observed in China
123 with COVID-19 could be linked to myocardial injury in patients [27-28]. Pericytes that have
124 ACE2 expression could be linked to the target of cardiac cells for SARS-CoV-2 entry. Pericytes
125 damage linked with virus infection which may further be associated with the capillary cells,
126 endothelial cell malfunction leads to microvascular dysfunction [29]. Acute respiratory illness
127 noticed in some of COVID-19 patients, like it was noticed with SARS-CoV and MERS-CoV
128 infected individual with signature of the pulmonary ground glass alteration on imaging [30].

129 Higher level of different cytokines; proinflammatory (IL-2), inflammatory (IFN- γ), and anti-
130 inflammatory (IL-6, IL-10) observed in the peripheral blood of critical cases compared to the
131 mild cases of COVID-19 [31]. However, the exact role of these cytokins is unknown in the
132 aspect of COVID-19. Different rates of mortality (3-13 %) was observed in different countries
133 which could be governed by different factors, such as immunological cross protection via some
134 earlier infection, presence of specific miRNA, hygiene, pollution or climate effect,, type of
135 population like percentage of old age group, culture, density, etc.

136

137 **Genome and replication:**

138 Coronaviruses are large, enveloped, (+)ssRNA that belong to Coronaviridae, order Nidovirales.
139 The previously known coronaviruses are classified into 4 different genera which are alpha
140 coronavirus, beta coronavirus, gamma coronavirus and delta coronavirus Among these alpha
141 and beta coronavirus are known to infect humans [32-33]. The first genome of SARS-CoV-2 was
142 discovered from the samples collected from bronchoalveolar lavage fluid from a patient from
143 Wuhan, China [34]. SARS-CoV-2 genome is +ssRNA, ~30 000bp long with 5'-cap and
144 3'-poly-A tail belonging to Coronaviridae family. The sequence of genes (5' to 3') in the viral
145 genome map is open reading frame 1a, (ORF1a), ORF1b, spike (S), envelope (E), membrane
146 (M) and nucleocapsid (N) [34]. SARS-CoV2 enters into the airway primary epithelial cells using
147 ACE2 receptors. Once it have in the cells it directly use its RNA genome as a template to
148 synthesize polyprotein 1a/1ab that encodes 16 non structural protein 1-16 (nsp1-16) to
149 established replication transcription complex in the double membrane vesicle [35] and after
150 wards ssRNA genome synthesized through replication transcription complex in discontinuous

151 manner. Roughly two-third of the entire genomes size covers the first open reading frames 1a/1b
152 ORFs (ORF1a/b), that encode 16 non-structural proteins. Other $\frac{1}{3}$ of the virus genome towards
153 the 3' end encodes spike, membrane, envelope, and nucleocapsid proteins [34].

154

155 **Evolution of SARS-CoV-2:**

156 The evolution of SARS-CoV-2 based on its phylogenetic and taxonomic nature has been
157 speculated from family of coronaviruses in host organisms with higher resemblance to SARS-
158 CoV as suggested by Coronavirus Study Group (CSG) of the International Committee on
159 Taxonomy of Viruses [36] SARS-CoV-2 whole genomic sequence share 96 % similarity with
160 Bat coronavirus (BatCoV RaTG13) source from *Rhinolophus affinis* from Yunnan Province in
161 China, indicating that SARS-CoV-2 has a very high probability of having its origin from bats
162 [37]. Primary genomic sequencing obtained from 5 different infected people with SARS-CoV-2
163 in China exhibited a 100 percent of sequence similarity. However, SARS-CoV 2 shares 96 %
164 and 91.02 % genomic sequence similarity with BatCoV RaTG13 and Pangolin-CoV respectively
165 [38].

166 Malayan pangolins (*Manis javanica*) coronavirus (Pangolin-CoV) shows 91.02% and
167 90.55% identity at the whole genome with SARS-CoV-2 and BatRaTG13 respectively [39].
168 Zhang et al. [39] study indicates that Pangolin-CoV is the common ancestor for Bat RaTG13 and
169 human SARS-CoV-2. Amino acid sequence analysis suggests that the S1 protein sequence of
170 Pangolin-CoV resembles more with SARS-CoV-2 as compared to Bat RaTG13. Interestingly 5
171 amino acid residues of Spike RBD that interact with human Angiotensin-converting enzyme 2

172 (ACE2) are conserved between SARS-CoV-2 and pangolin-CoV, but are not identical with
173 RaTG13 due to four amino acid mutations. The binding efficiency of Pangolin COV and SARS
174 CoV-2 spike RBD with ACE2 resembles that indicating the possibility of it being an
175 intermediate host. However, the difference in genetic sequence similarity of SARS-CoV-2 with
176 other coronaviruses could be a recombination of different coronaviruses, maybe via infecting a
177 common unknown host before coming in contact with humans. These developed chimeric
178 viruses simultaneously that can interact with various cells of the host and infect. Though
179 verification of this hypothesis warrants further studies. Concluded that there is any intermediate
180 host for viruses to infect humans.

181

182 **Diagnosis and Prevention:**

183 Diagnosis is an important part of controlling a disease, especially a communicable one and has
184 played its pivotal part in the containment of COVID-19. With proper diagnosis we can steer the
185 tactful implementation of control measures to limit the spread primarily by identifying and
186 isolating the affected individual. Proper diagnostic measures are very important as the symptoms
187 shown by infected people are very wide and aren't a true prognosis or measure of the infection.
188 Thus, validating the requirement of highly sensitive and specific laboratory diagnostic
189 assessments. Almost four months after the initial cases the genomic and proteomic components
190 of the pandemic have been discovered. However, the host response to the virus infection is yet to
191 be explored [40]. Various approaches are being followed to understand the COVID-19 infection
192 including a combination of computed tomography imaging, whole genome sequencing, and
193 electron microscopy [41]. Moreover, ELISA based antibody and antigen detection tests are being

194 developed as well. Although not yet been completely validated and still under research. This can
195 have a much precise outcome given genetic sequence similarity to the antibodies generated [42-
196 43]. The lack of diagnostic measures have paved the path to use RT-PCR as a gold standard
197 diagnostic measure and have shown reliable results worldwide [44]. Furthermore, RT-PCR
198 detection also previously exhibited high sensitivity and specificity for SARS-CoV and MERS-
199 CoV infection. Another commonly used nucleic acid based detection technology is high-
200 throughput sequencing of the entire viral genome, but it is an expensive and time consuming
201 detection method. To tackle these problems, CRISPR-based sensitive, rapid and potentially
202 portable diagnostic tests are in the development to facilitate rapid point-of-care testing and solves
203 worldwide testing shortages of COVID-19. [45]. In India, recently CRISPR-Cas9 technology
204 based paper-strip tool was developed and operates by converting the RNA into DNA then
205 amplification and finally utilizing the Cas9 complex to detect the genetic material of COVID-19.

206 The unavailability of clinically proven therapeutic agents is also a gigantic reason for
207 such a blanket spread of COVID-19. However, development of therapeutics and vaccines is
208 underway. There are various potential therapeutic targets to repurpose the present antiviral
209 therapy for developing effective interventions against this novel coronavirus. One of the
210 promising antiviral drugs against RNA viruses *in vitro* and *in vivo* is Remdesivir which is a
211 nucleoside analogue that blocks viral RNA synthesis [46]. Furthermore, potential clinical trials
212 have been reported to combat COVID-19 using the HIV drug combination of Lopinavir–
213 Ritonavir [47]. Additionally, other nucleoside analogues *i.e.* Favipiravir, Ribavirin and
214 Galidesivir are in the pipeline and may have potential clinical application against COVID-19
215 [48]. The National Medical Products Administration of China has approved an antiviral drug
216 ‘Favilavir’ for the treatment for coronavirus [49].

217 Favilavir has reportedly shown efficacy in 70 patients in a clinical study. Allowing other
218 protease inhibitors such as but not limited to Lopinavir-Ritonavir that are previously approved
219 for HIV treatment to be studied. The two protease inhibitors mentioned above works by blocking
220 viral entry inside the cell and inhibiting viral particle maturation. This can be one of the effective
221 therapy for management and treatment of COVID-19. Other FDA approved drugs including
222 Chloroquine and Hydroxychloroquine are being tested in multiple studies as well. So far they
223 have demonstrated an efficient inhibition of COVID-19 infection *in vitro* and in clinical studies
224 [50]. Chloroquine phosphate has also been coined as a potential drug based on current clinical
225 trial data. The mechanism of hydroxychloroquine consists of inhibition of the virion assembly in
226 the endoplasmic reticulum and Golgi intermediate compartments. Followed by attenuating the
227 expression of pro-inflammatory factors, receptors and phosphatidylinositol binding clathrin
228 assembly protein (PICALM), which prevents endocytosis-mediated uptake of COVID-19 virus
229 [51]. Azithromycin along with hydroxychloroquine also has been found to be efficient on SARS-
230 CoV-2 infection treatment in Chinese COV-19 patients and found it reinforced with other drugs
231 because it inhibits the super infection of bacteria [52].

232 The subsequent drugs can be designed in a way that inhibits or blocks S protein binding
233 to ACE2 receptors. This can be an efficient way of attenuating or even completely ceasing the
234 infection of the virus in COVID-19 patients [53]. As with the current understanding of COVID-
235 19, the virus invades the host cells through an ACE2 receptor that is predominantly present in
236 lungs epithelial cells. The other major target for drug development via high affinity can also be
237 AP2-associated protein kinase 1 (AAK1) as some recent studies suggest [54]. Moreover, the
238 kinase inhibitor including ‘baricitinib’ can also be a potential candidate that can be evaluated in
239 the clinical trial settings for COVID-19 management [54]. Currently, the plasma/serum

240 immunoglobulin of patients recovered from COVID-19 can produce favorable results in
241 COVID-19 patients especially the one who are in later stages of disease progression [55]. There
242 are other potential therapies and as the science has grown so much we can also look at the
243 current generation of monoclonal antibodies (mAb), this can play a mechanism of neutralizing
244 the virus and hence reducing the symptoms of COVID-19. Also, by binding with the receptor-
245 binding domain of virus can also be developed as a potential candidate therapeutics of COVID-
246 19 infections [56]. We strongly believe and understand the importance of vaccination in the
247 current scenario and if it spreads with current prediction vaccination would be utmost important
248 and the best way to control the spread of disease. There have been various studies at different
249 levels of development currently underway in almost every part of the globe a lot of them have
250 been shown tested *in vivo* and shown promising results. The vaccination can be of different types
251 and these include live-attenuated, inactivated, subunit, recombinant DNA and proteins vaccines
252 [57]. The major disadvantage of any strategy or vaccine development is that it requires
253 approximately 2 years to be available for use with many more years and a well crafted strategy
254 for it to reach parts of the world. Being the need of the hour a lot of new pharmaceutical
255 molecules including HIV drugs and stem cell therapies are in testing phases of clinical trials.
256 Needless to mention the advent of modern technology has paved the path of other treatment
257 opportunities like siRNA, tumor necrosis factor-alpha inhibitors, interferons, neutralizing
258 antibodies, neuraminidase inhibitors, corticosteroids, pentoxifylline *etc.* Further evaluation of
259 these mechanisms can elaborate their usage and how these can help in combating the current
260 epidemic crisis [58].

261 We are still in a phase where we are unfolding the layers, evaluating and understanding
262 the virus. The new development in the current knowledge may open an opportunity for many

263 more therapeutic agents and targets to control the spread, manage and treat COVID-19. Needless
264 to mention this requires extensive research both in discovery science and clinical study settings.
265 The research community worldwide has come together and never before have we seen such a
266 sharing of research findings and joining of hands to work against a common goal to eradicate this
267 unwanted disease from our society. In the meantime when the research community is utilizing
268 every second available on the clock the people awaiting have to be strong and follow self care
269 and as a lot of scientists have suggested that boosting the immune system can play a pivotal role
270 to prevent the spread of COVID-19 infection. These trying times have provided us with an
271 opportunity to analyze and manage our lifestyle. This won't only help us with the current
272 epidemic but in general with other epidemiological diseases as well. Some of us have the
273 privilege to do workout routines, yoga, meditation, tai-chi, breathing exercise, aerobics and many
274 more ways to stay healthy, calm, eliminate anxiety and also take care of mental health needless
275 to mention as little of a task as getting proper sleep can be very essential and helpful to boost the
276 immune system [59]. Also, we can always access the ancient treasure and utilise the wisdom
277 from traditional systems of medicine like Ayurveda, herbal medicine, Chinese medicine etc.
278 These could also help or maybe possess a preventive potential measure while we still await a
279 tried and tested cure from modern medicine [60].

280

281 **Conclusion:**

282 The current available diagnostic technologies have enabled researchers to develop new realtime
283 efficient COVID-19 diagnostics tools. Based on the current knowledge of Coronavirus
284 pathogenesis and SARS-CoV-2 disease mechanisms known so far, newer tools could attainably

285 be developed to prevent, diagnose and treat people affected with COVID-19 and also to manage
286 the disease in general. In conclusion, we would like to stress out again on the importance of
287 diagnostics as a part of the important components to dealing with outbreaks in future. The proper
288 management and availability of diagnostic tools can aid in curbing the spread of pathogens like
289 coronavirus thus leading to a better survival rate. Some of the drugs may also have prophylactic
290 and therapeutic effects against COVID-19 infection, exploring the mechanisms of these drugs
291 and the keep on evaluating the effect of virus is absolutely necessary to explore and develop new
292 preventative and therapeutic strategies.

293

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299

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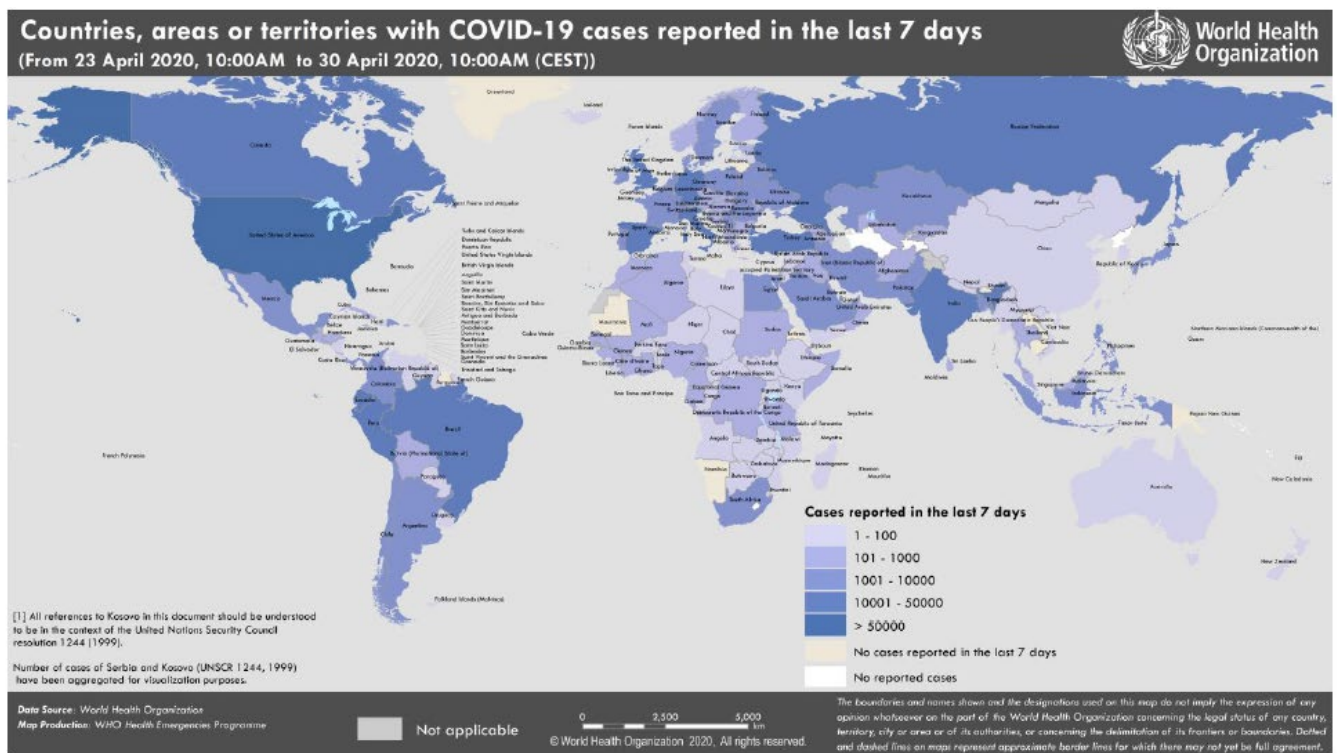
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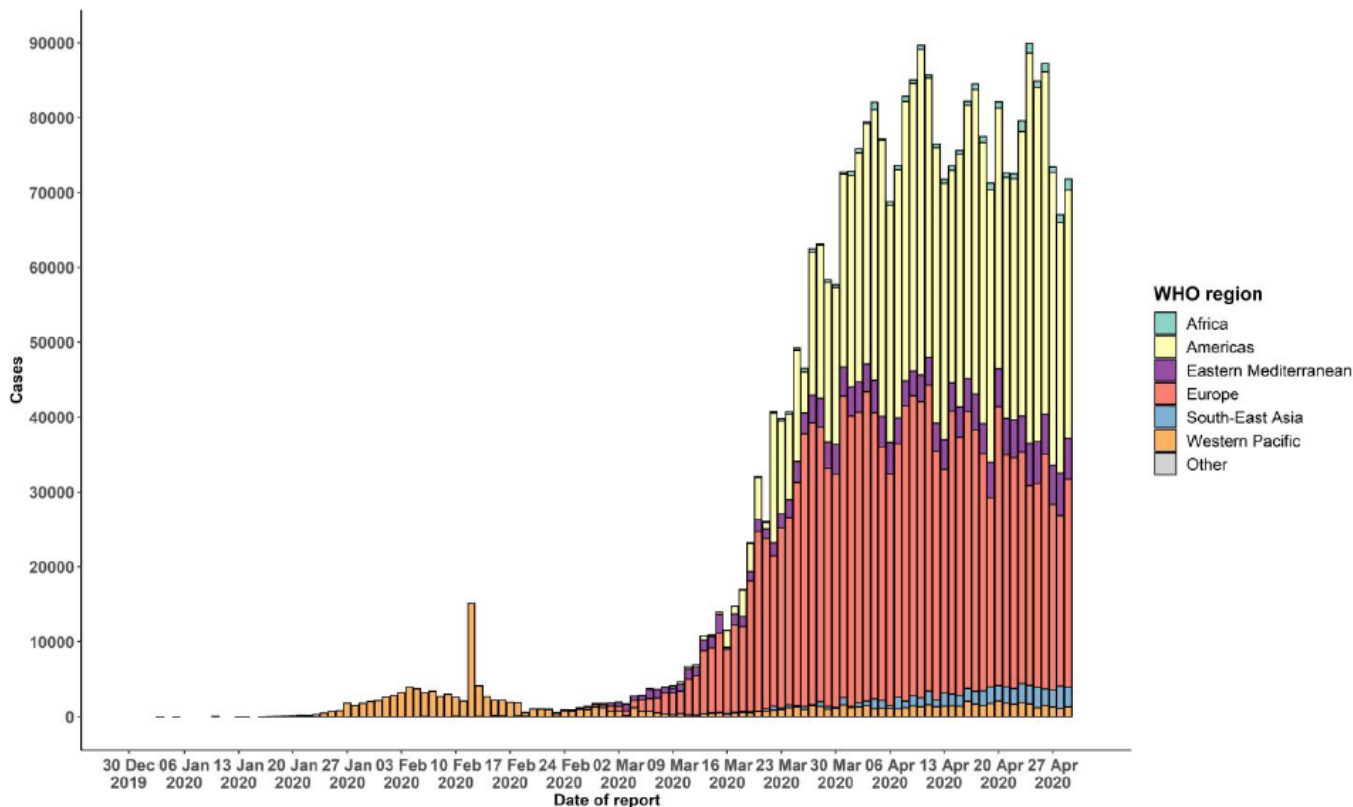
496 Figure Legends



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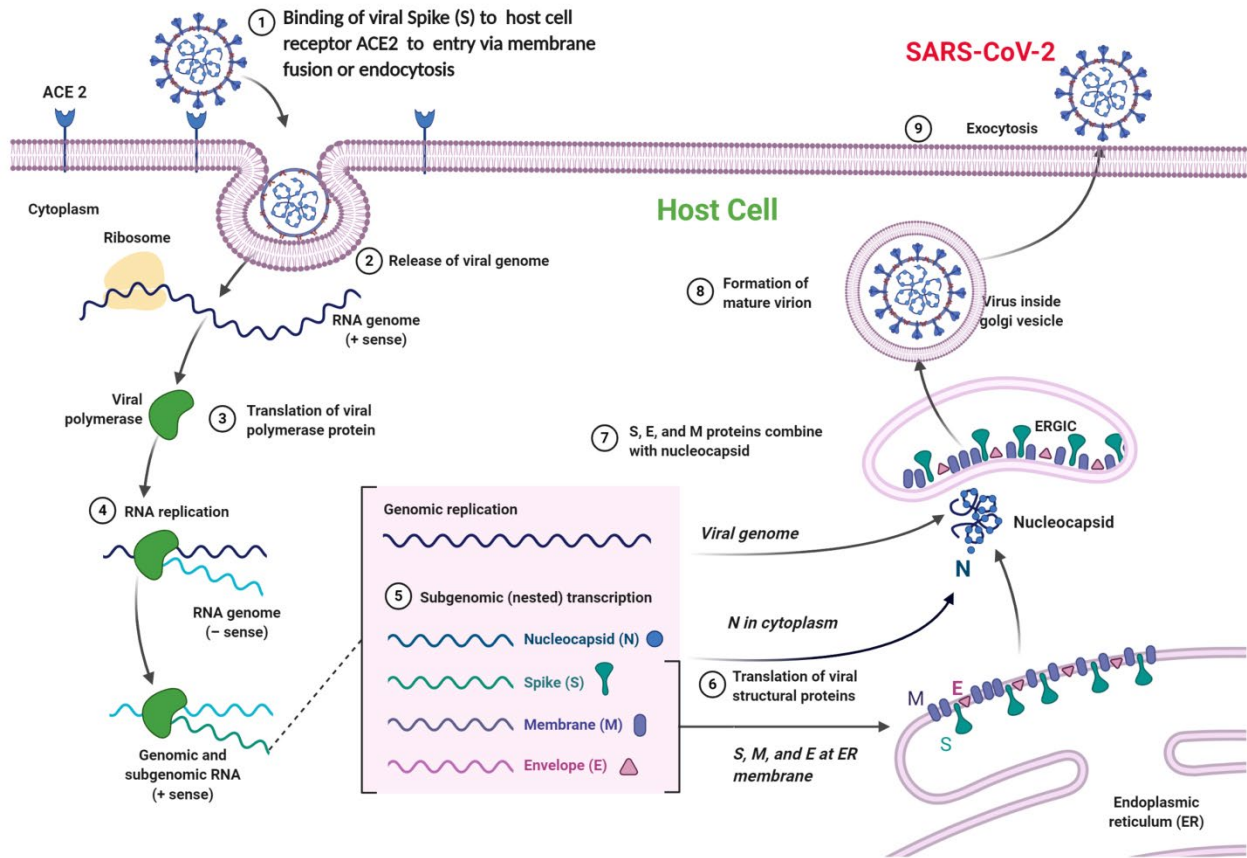
498 **Figure 1:** Represent the number of confirmed cases across countries, territories or areas as of 30
 499 April 2020 (with permission this figure was adopted from WHO situation report 101 on the

500 COVID-19;<https://www.who.int/emergencies/diseases/novel-coronavirus-2019/situation->
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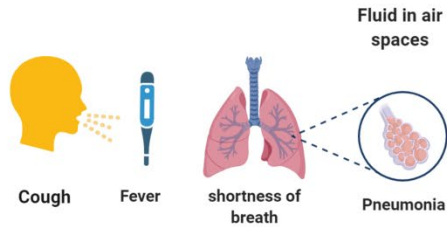


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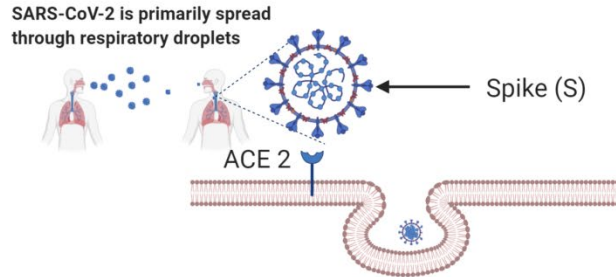
503 **Figure 2.** Represent the epidemic curve of confirmed COVID-19, by date of report and WHO
504 region through 30 April 2020 (with permission this figure was adopted from WHO situation
505 report 101 on the COVID-19; <https://www.who.int/emergencies/diseases/novel-coronavirus->
506 2019/situation-reports



Symptoms



Transmission



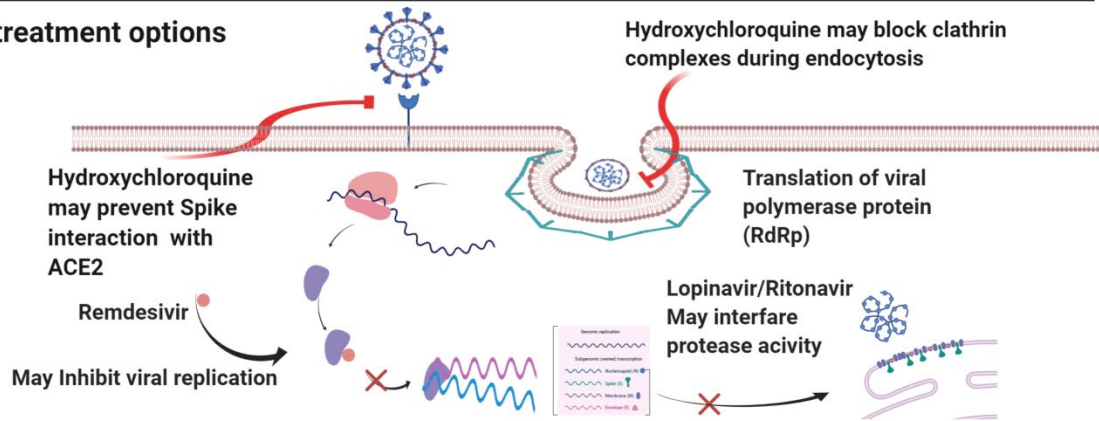
Diagnosis



Prevention



Possible treatment options



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515 **Graphical abstract:**

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