

1 Broad anti-coronaviral activity of FDA approved drugs against
2 SARS-CoV-2 *in vitro* and SARS-CoV *in vivo*

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15
16 **Abstract**

17 SARS-CoV-2 emerged in China at the end of 2019 and has rapidly become a pandemic with
18 roughly 2.7 million recorded COVID-19 cases and greater than 189,000 recorded deaths by
19 April 23rd, 2020 (www.WHO.org). There are no FDA approved antivirals or vaccines for any
20 coronavirus, including SARS-CoV-2. Current treatments for COVID-19 are limited to supportive
21 therapies and off-label use of FDA approved drugs. Rapid development and human testing of
22 potential antivirals is greatly needed. A quick way to test compounds with potential antiviral
23 activity is through drug repurposing. Numerous drugs are already approved for human use and
24 subsequently there is a good understanding of their safety profiles and potential side effects,

25 making them easier to fast-track to clinical studies in COVID-19 patients. Here, we present data
26 on the antiviral activity of 20 FDA approved drugs against SARS-CoV-2 that also inhibit SARS-
27 CoV and MERS-CoV. We found that 17 of these also inhibit SARS-CoV-2 at a range of IC50
28 values at non-cytotoxic concentrations. We directly follow up with 7 of these to demonstrate all
29 are capable of inhibiting infectious SARS-CoV-2 production. Moreover, we have evaluated two
30 of these, chloroquine and chlorpromazine *in vivo* using a mouse-adapted SARS-CoV model and
31 found both drugs protect mice from clinical disease.

32

33 **Introduction**

34 At the end of December 2019, reports started to emerge from China of patients suffering from
35 pneumonia of unknown etiology. By early January, a new coronavirus had been identified and
36 determined as the cause (1). Since then, the virus originally known as novel coronavirus 2019
37 (nCoV-2019), now severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has spread
38 around the world. As of April 23th, 2020, there have been roughly 2.7 million confirmed cases of
39 COVID-19 (the disease caused by SARS-CoV-2 infection) with over to 189,000 recorded deaths
40 (www.WHO.org). Multiple countries have enacted social distancing and quarantine measures,
41 attempting to reduce person-to-person transmission of the virus. Healthcare providers lack
42 pharmaceutical countermeasures against SARS-CoV-2, beyond public health interventions, and
43 there remains a desperate need for rapid development of antiviral therapeutics. A potential route
44 to candidate antivirals is through repurposing of already approved drugs (for reviews; (2–4) and
45 for examples;(5–8). We have previously screened a library of FDA approved drugs for antiviral
46 activity against two other highly pathogenic human coronaviruses, SARS-CoV and Middle East
47 respiratory syndrome coronavirus (MERS-CoV)(6). We found 27 drugs that inhibited replication
48 of both of these coronaviruses, suggesting that they may have broad anti-coronaviral activity.
49 One of the hits from this work was imatinib with which we subsequently determined the

50 mechanism of action by demonstrating this drug inhibits fusion of coronaviruses with cellular
51 membranes, thus blocking entry (9, 10).

52

53 Here, we present our investigation of 20 priority compounds from our previous screening to test
54 if they can also inhibit SARS-CoV-2. Since these compounds are already approved for use in
55 humans, they make ideal candidates for drug repurposing and rapid development as antiviral
56 therapeutics. Our work found that 17 of the 20 of the drugs that inhibited SARS-CoV and
57 MERS-CoV could also inhibit SARS-CoV-2, with similar IC₅₀ values. We further assessed a
58 subset of these drugs for their effects on SARS-CoV-2 RNA and infectious virus production and
59 found all to have inhibitory activity. Our screening based on cytopathic effect therefore appears
60 a favorable approach to find drugs capable of inhibiting production of infectious virus. Currently
61 there are no established small animal model systems for SARS-CoV-2. However, there is a
62 well-established mouse-adapted system for SARS-CoV (MA15 strain(11)) and we present data
63 here assessing the *in vivo* efficacy of chloroquine (CQ) and chlorpromazine (CPZ) against
64 SARS-CoV. We found that drug treatment does not inhibit virus replication in mouse lungs, but
65 significantly improves clinical outcome. Based on both of these drugs inhibiting SARS-CoV-2
66 infection *in vitro* and providing protection *in vivo* against SARS-CoV clinical disease we believe
67 they may be beneficial for SARS-CoV-2 therapy.

68

69 **Materials and Methods**

70 **Cell lines and virus**

71 Vero E6 cells (ATCC# CRL 1586) were cultured in DMEM (Quality Biological), supplemented
72 with 10% (v/v) fetal bovine serum (Sigma), 1% (v/v) penicillin/streptomycin (Gemini Bio-
73 products) and 1% (v/v) L-glutamine (2 mM final concentration, Gibco). Cells were maintained at
74 37°C and 5% CO₂. Samples of SARS-CoV-2 were obtained from the CDC following isolation

75 from a patient in Washington State (WA-1 strain - BEI #NR-52281). Stocks were prepared by
76 infection of Vero E6 cells for two days when CPE was starting to be visible. Media were
77 collected and clarified by centrifugation prior to being aliquoted for storage at -80°C. Titer of
78 stock was determined by plaque assay using Vero E6 cells as described previously (12). All
79 work with infectious virus was performed in a Biosafety Level 3 laboratory and approved by our
80 Institutional Biosafety Committee. SARS-CoV stock was prepared as previously described (13).
81 SARS-CoV spike (S) psuedotype viruses were produced as previously described (9).

82

83 **Drug testing**

84 All drug screens were performed with Vero E6 cells. Cells were plated in opaque 96 well plates
85 one day prior to infection. Drug stocks were made in either DMSO, water or methanol. Drugs
86 were diluted from stock to 50 µM and an 8-point 1:2 dilution series made. Cells were pre-treated
87 with drug for 2 hour (h) at 37°C/5% CO₂ prior to infection at MOI 0.01 or 0.004. Vehicle controls
88 were used on every plate, and all treatments were performed in triplicate for each screen. In
89 addition to plates that were infected, parallel plates were left uninfected to monitor cytotoxicity of
90 drug alone. Three independent screens with this set-up were performed. Cells were incubated
91 at 37°C/5% CO₂ for 3 days before performing CellTiter-Glo (CTG) assays as per the
92 manufacturer's instruction (Promega). Luminescence was read using a Molecular Devices
93 Spectramax L plate reader. Fluphenazine dihydrochloride, benztropine mesylate, amodiaquine
94 hydrochloride, amodiaquine dihydrochloride dihydrate, thiethylperazine maleate, mefloquine
95 hydrochloride, triparanol, terconazole vetranal, anisomycin, fluspirilene, clomipramine
96 hydrochloride, hydroxychloroquine sulfate, promethazine hydrochloride, emetine dihydrochloride
97 hydrate and chloroquine phosphate were all purchased from Sigma. Chlorpromazine
98 hydrochloride, toremifene citrate, tamoxifen citrate, gemcitabine hydrochloride and imatinib
99 mesylate were all purchased from Fisher Scientific.

100

101 **Data analysis**

102 Cytotoxicity (%TOX) data was normalized according to cell-only uninfected (cell only) controls
103 and CTG-media-only (blank) controls:

$$\%TOX = \left(1 - \frac{(drug) - (blank)}{(cell\ only) - (blank)} \right) \times 100$$

104 Inhibition (%Inhibit) data was normalized according to cell only and the activity of the vehicle
105 controls:

$$\%Inhibit = \frac{(drug) - (vehicle)}{(cell\ only) - (vehicle)} \times 100$$

106 Nonlinear regression analysis was performed on the normalized %inhibit and %TOX data and
107 IC50s and CC50s were calculated from fitted curves (log [agonist] versus response - variable
108 slope [four parameters]) (GraphPad Software, LaJolla, CA), as described previously
109 (Dyall2018). Drug dilution points in a given run were excluded from IC50 analysis if the average
110 cytotoxicity was greater than 30% (arbitrary cutoff) across the 3 cytotoxicity replicates for that
111 screen. IC50 or CC50 values extrapolated outside the drug dilution range tested were reported
112 as greater than 50µM or less than 0.39µM. Selectivity indexes (SI) were also calculated by
113 dividing the CC50 by the IC50.

114

115 **Viral infection**

116 To further analyse candidate drugs, Vero E6 cells were grown in 24 well plate format for 24 h
117 prior to infection. As with the drug screens, cells were pre-treated with drug at a range of
118 concentrations, or vehicle control for 2 h. Cells were then infected with SARS-CoV-2 at MOI 0.1
119 for 24 hour (h). Supernatant was collected, centrifuged in a table-top centrifuge for 3 minutes
120 (min) at max speed and stored at -80°C. After a wash in PBS, infected cells were collected in

121 TRIZol (Ambion) for RNA analysis (described below). Supernatant was used to titer viral
122 production by TCID₅₀ assay (12).

123

124 **RNA extraction and qRT-PCR**

125 RNA was extracted from TRIZol samples using Direct-zol RNA miniprep kit (Zymo Research) as
126 per the manufacturer's instructions. RNA was converted to cDNA using RevertAid RT Kit
127 (Thermo Scientific), with 12 µl of extracted RNA per reaction. For qRT-PCR, 2 µl of cDNA
128 reaction product was mixed with PowerUp SYBR Green Master Mix (Applied Biosystems) and
129 WHO/Corman primers targeting N and RdRp: N FWD 5'-CACATTGGCACCCGCAATC-3', N
130 REV 5'-GAGGAACGAGAAGAGGCTTG-3', RdRp FWD 5'GTGARATGGTCATGTGTGGCGG-
131 3', RdRp REV 5'-CARATGTTAAASACACTATTAGCATA-3'. The qRT-PCR reactions were
132 performed with a QuantStudio 5 (Applied Biosystems). To normalize loading, 18S RNA was
133 used as a control, assessed with TaqMan Gene Expression Assays (Applied Biosystems) and
134 TaqMan Fast Advanced Master Mix. Fold change between drug treated and vehicle control was
135 determined by calculating $\Delta\Delta CT$ after normalization to the endogenous control of 18S.

136

137

138 **Pseudovirus fusion/entry assay**

139 The pseudovirion (PV) entry assay was performed as described (9, 14). Briefly, 2×10^4 BSC1
140 cells per well were in 96-well plates for 24 h, after which time cells were pre-treated with drug (1
141 h) and infected with PV (3 h). Media was removed and cells were washed with loading buffer
142 (47 ml clear DMEM, 5 mM Probenecid, 2 mM L-glutamine, 25 mM HEPES, 200 nM bafilomycin,
143 5 µM E64D) and incubated for 1 h in CCF2 solution (LB, CCF2-AM, Solution B [CCF2-AM kit
144 K1032] Thermo Fisher) in the dark. Cells were washed once with loading buffer and incubated
145 from 6 h to overnight with 10% FBS in loading buffer. Percentage CCF2 cleavage was assessed

146 by flow cytometry on the LSRII (Beckton Dickinson) in the flow cytometry core facility at the
147 University of Maryland, Baltimore. Data were analyzed using FlowJo.

148

149 **Mouse infections.**

150 All infections were performed in an animal biosafety level 3 facility at the University of Maryland,
151 Baltimore, using appropriate practices, including a HEPA-filtered bCON caging system, HEPA-
152 filtered powered air-purifying respirators (PAPRs), and Tyvek suiting. All animals were grown to
153 10 weeks of age prior to use in experiments. The animals were anesthetized using a mixture of
154 xylazine (0.38 mg/mouse) and ketamine (1.3 mg/mouse) in a 50 μ l total volume by
155 intraperitoneal injection. The mice were inoculated intranasally with 50 μ l of either PBS or 2.5 x
156 10^3 PFU of rMA15 SARS-CoV (11) after which all animals were monitored daily for weight loss.
157 Mice were euthanized at day 4 post-infection, and lung tissue was harvested for further
158 analysis. All animals were housed and used in accordance with the University of Maryland,
159 Baltimore, Institutional Animal Care and Use Committee guidelines.

160

161 **Plaque assay.**

162 Vero cells were seeded in 35 mm dishes with 5×10^5 cells per dish 24 h prior to
163 infection. Supernatants from homogenized were serially diluted 10^{-1} through 10^{-6} in serum-free
164 (SF) media. Cells were washed with SF media, 200 μ l of diluted virus was added to each well
165 and adsorption was allowed to proceed for 1 h at 37°C with gentle rocking every 10 min. 2X
166 DMEM and 1.6% agarose were mixed 1:1. Cells were washed with SF media, 2 ml DMEM-
167 agarose was added to each well, and cells were incubated for 72 h at 37°C, after which time
168 plaques were read.

169

170

171 **Results**

172 **Screening FDA approved compounds for anti-SARS-CoV-2 activity**

173 Previously, we performed a large-scale drug screen on 290 FDA approved compounds to
174 investigate which may have antiviral activity against SARS-CoV and MERS-CoV (6). With the
175 emergence of SARS-CoV-2, we prioritized testing 20 of the 27 hits that were determined to
176 inhibit both of the previously tested coronaviruses for antiviral activity against the novel virus.
177 The list of tested compounds is shown in Table 1. Our screening started at 50 μ M and used an
178 8-point, 1:2 dilution series with infections being performed at either MOI 0.01 or 0.004. CellTiter-
179 Glo (CTG) assays were performed 3 days post-infection to determine relative cell viability
180 between drug and vehicle control treated cells. Uninfected samples were used to measure the
181 cytotoxicity of drug alone. From the relative luminescence data of the CTG assay, percent
182 inhibition (of cell death caused by viral infection) could be measured and plotted along with the
183 percent cytotoxicity of drug alone. Fig. 1 shows these plotted graphs from one representative of
184 three independent screens at MOI 0.01. For those drugs demonstrating a cell toxicity rate lower
185 than 30%, we were able to calculate IC₅₀ values at both MOI from these graphs for 17 of the 20
186 drugs which is summarized in Table 1.

187

188 **Drug screen validation**

189 In order to validate our screening process as a means to determine compounds with antiviral
190 effect we decided to follow up with a subset of drugs. Chloroquine (CQ) has become the source
191 of much interest as a potential treatment for COVID19 (15), as such, we further investigated
192 hydroxychloroquine (HCQ) and CQ as both were present in our screen (Table 1). Vero E6 cells
193 were plated and pre-treated with drug for 2 h prior to infection with SARS-CoV-2 at MOI 0.1.
194 Supernatant was collected 24 h post-infection to determine titer of virus by TCID₅₀ assay and
195 cells were collected in TRIzol to assess production of viral mRNA. Treatment with both drugs
196 caused a significant reduction in viral mRNA levels, especially at higher concentrations, without
197 drug induced cytotoxicity (Fig. 1 and Table 1). There was a significant decrease in relative

198 expression levels of both RdRp and N mRNA across the range of concentrations used (Fig. 2A-
199 D). Along with causing a reduction in viral mRNA, treatment with both drugs caused a significant
200 reduction in viral replication (Fig. 2E and 2F). SARS-CoV-2 production was more sensitive to
201 HCQ than CQ with larger inhibition seen at the same concentration of treatment, which is in
202 agreement with HCQ having a lower IC50 in our cell viability assay (Table 1). We also
203 performed a time of addition assay with the highest concentration of HCQ to investigate whether
204 SARS-CoV-2 entry was the point of inhibition of this compound (Fig. 2G). Interestingly, while the
205 addition of HCQ at 2h post-infection did have some reduction in inhibitory activity but there was
206 not a complete loss, suggesting that HCQ treatment may impact other stages of the viral life
207 cycle than just entry.

208
209 We have previously used a β -lactamase-Vpr chimeric protein (Vpr-BlaM) pseudotype system to
210 demonstrate that imatinib (a drug also seen to inhibit SARS-CoV-2 [Table 1 and Fig. 1]) inhibits
211 SARS-CoV and MERS-CoV spike-mediated entry (9). We used this system to more directly
212 investigate whether CQ could inhibit viral entry mediated by coronavirus spike, and additionally
213 included chlorpromazine (CPZ) as this is known to inhibit clathrin-mediated endocytosis (16)
214 and was also part of our drug screening (Table 1 and Fig. 2). In this assay, when the
215 pseudovirus fuses with a cellular membrane, BlaM is released into the cytoplasm of the infected
216 cell. BlaM can cleave cytoplasmic loaded CCF2 to change its emission spectrum of from 520
217 nm (green) to ~450 nm (blue), which can be quantified using flow cytometry.

218
219 Cells were treated with CQ or CPZ for 1 h before infection with BlaM-containing SARS-S
220 pseudovirions (PV). Cells were then analyzed by flow cytometry to quantify the cleavage of
221 CCF2. In mock treated cells infected with SARS-S PV there was a shift in the CCF2 emission
222 spectrum indicating release of BlaM to the cytosol, and that spike-mediated fusion with cellular
223 had occurred. Upon treatment with CQ or CPZ there was a greater than 90% reduction in CCF2

224 cleavage caused by SARS-S PV (Fig. 2H). These data demonstrate that both drugs inhibit
225 SARS-CoV spike-mediated fusion with cellular membranes. These psuedotype assays suggest
226 that the inhibition of coronavirus replication caused by CQ and CPZ is at the stage of entry to
227 cells but combined with the time of addition assays (Fig. 2G), there is a suggestion that later
228 stages may also be impacted.

229
230 HCQ and CQ are used as anti-malarial drugs and are in the class of aminoquinolines which are
231 hemozoin inhibitors, similarly to 4-methanolquinolines. Interestingly, from our drug screening,
232 three other hemozoin inhibitors were identified: amodiaquine dihydrochloride dihydrate,
233 amodiaquine hydrochloride and mefloquine. We therefore decided to directly test these drugs
234 for antiviral activity against SARS-CoV-2. We directly tested CPZ against SARS-CoV-2 having
235 seen that it could inhibit SARS-CoV S-mediated entry to cells (Fig. 2H). And finally included
236 imatinib since we have previously shown that this can inhibit entry of both SARS-CoV and
237 MERS-CoV (9) and was a hit against SARS-CoV-2 (Fig. 1 and Table 1). Again, cells were pre-
238 treated with drugs at the indicated concentrations and infected with SARS-CoV-2 at MOI 0.1 for
239 24h, after which supernatant samples were collected. As can be seen in Fig. 3, at the highest
240 concentrations of all drugs there is significant inhibition of SARS-CoV-2 infection. All 5 drugs
241 showed very strong inhibition at 20 μ M.

242
243 Overall, the data from Fig. 2 and Fig. 3 indicate that there are various FDA approved drugs that
244 have broad-spectrum anti-coronavirus activity *in vitro*.

245
246 **Chloroquine and chlorpromazine do not inhibit SARS-CoV (MA15) replication in mouse**
247 **lungs, but significantly reduces weight loss and clinical signs**

248 CQ and CPZ treatment displayed significant inhibition of coronavirus replication *in vitro*, with our
249 data suggesting entry is inhibited. We therefore decided to investigate whether these drugs

250 were efficacious *in vivo* using SARS-CoV strain, MA15 in BALB/c mice. This model displays
251 ~15-20% weight loss by 4 days post infection (dpi), occasionally resulting in death. We tested
252 whether prophylactically administered CQ or CPZ could protect mice from severe MA15
253 infection. Mice were injected intraperitoneally with either water, 0.8 mg CQ, 1.6 mg CQ, 20 µg
254 CPZ, 100 µg CPZ or 200 µg CPZ at day -1 of infection and then were dosed every day through
255 the 4 days of infection. On day 0, mice were intranasally infected with 2.5×10^3 pfu of SARS-
256 CoV (MA15) or PBS as control. Weight loss was measured as a correlate of disease and mice
257 were euthanized at 4 dpi for analysis.

258
259 PBS inoculated mice showed no weight loss or clinical signs of disease when treated with either
260 water, CQ or CPZ over the experiment time course indicating drug treatment did not adversely
261 affect morbidity (Fig. 4A and 4D). Mice that were infected with MA15 and treated with water lost
262 ~15% of their starting body weight over 4 days and had significant clinical signs of disease
263 including ruffled fur, labored breathing and lethargy (Fig. 4A and 4D). Mice that were treated
264 with 0.8 mg of CQ each day, displayed similar weight loss as the water control through the first
265 3 days of infection, however by 4 dpi the weight loss was halted in the drug treated mice (Fig.
266 4A). Mice that were treated with 1.6 mg CQ per day showed markedly reduced weight loss
267 compared to the water control (Fig. 4A). Pathological analysis was also performed on H&E
268 stained sections. Mice infected with MA15 and treated with water displayed significant
269 inflammation and denuding bronchiolitis suggesting severe disease (Fig. 4B). By contrast, 0.8
270 mg CQ dose group had moderate inflammation that was reduced compared to control and the
271 1.6 mg dose group had minimal lung pathology (Fig. 4B). Interestingly, even though CQ
272 treatment appeared to protect against weight loss and inflammation in the lungs, the viral titer
273 was equivalent between drug treated and vehicle control mice (Fig. 4C).

274

275 Similar to the CQ results, CPZ treatment reduced weight loss in mice infected with MA15 at 100
276 μg and 200 μg , but the 20 μg treatment group were equivalent to vehicle control (Fig. 4D) and
277 the H&E sections showed protection against inflammation and denuding bronchiolitis at the
278 higher doses (Fig. 4E). Again, as with CQ treatment, even though there were reduced signs of
279 infection with CPZ treatment, there was no difference in MA15 titer in the mouse lungs (Fig. 4F).
280 Overall these data indicate that even though CQ and CPZ treatment do not inhibit viral
281 replication in the lungs, both can protect mice from signs of disease following SARS-CoV
282 (MA15) infection.

283

284 **Discussion**

285 The SARS-CoV-2 pandemic has demonstrated the desperate need for antiviral drugs. Since the
286 emergence of SARS-CoV in 2002, research has uncovered many details of coronavirus biology
287 and pathogenesis, however there are currently no approved therapeutics against this emerging
288 virus family. Whether being used for treating SARS-CoV-2 in this current pandemic or the next
289 unknown viral pathogen in the future, we must attempt to develop and validated antiviral drugs
290 that are ready to be used at the first signs of an outbreak. Many FDA approved drugs have been
291 found to have antiviral activity, in addition to their approved use (e.g; (5–8)) and since these are
292 extensively used in humans for other conditions, they could be streamlined for rapid approval
293 and repositioning as antivirals. In our previous work, 290 FDA approved drugs were screened
294 for antiviral activity and 27 were found to inhibit both SARS-CoV and MERS-CoV (6). We
295 prioritized testing these for antiviral activity against SARS-CoV-2 since they displayed broad-
296 spectrum antiviral activity. From multiple independent screens performed with two MOI, we
297 found that 17 of our 20 tested priority compounds display significant antiviral activity at non-
298 cytotoxic concentrations. Many of the compounds have IC₅₀ values under 10 μM and these will
299 be the source of follow up testing on additional cell lines and in mouse models of SARS-CoV-2.

300

301 We further investigated 7 of the hits to directly test if they inhibited SARS-CoV-2 replication. We
302 performed follow-up experiments with HCQ, CQ, amodiaquine and mefloquine because
303 chloroquine has garnered much interest as a potential treatment for COVID19 (15) and the
304 others are similarly used as anti-malarial compounds (17). In addition, we have previously
305 demonstrated that imatinib is an inhibitor of SARS-CoV, MERS-CoV and infectious bronchitis
306 virus entry to cells (9, 10) so included that here as the mechanism of coronavirus inhibition is
307 understood. Finally, CPZ inhibits clathrin function in cells (16) so can disrupt infection by many
308 viruses that require clathrin-mediated endocytosis and was therefore also chosen for further
309 analysis. Treatment of cells with all these drugs showed inhibition of infectious viral particle
310 production (measured by TCID₅₀ assay) at non-cytotoxic levels.

311
312 Having demonstrated that HCQ, CQ and CPZ can inhibit cytopathic effect, mRNA synthesis and
313 infectious viral particle production of SARS-CoV-2, we used a previously published system of
314 SARS-CoV psuedotype viruses carrying Vpr-BlaM to investigate whether CQ and CPZ inhibit
315 coronavirus spike-mediated entry to better define mechanism of action. We have previously
316 used this system to define imatinib as an entry inhibitor of these viruses (9) and found similar
317 results for CQ and CPZ, thus better defining their mechanism of a antiviral activity.

318
319 Finally, we investigated the efficacy of CQ and CPZ with an *in vivo* model using SARS-CoV
320 MA15. There is currently a lack of an established mouse model for SARS-CoV-2 so we used the
321 mouse adapted SARS-CoV (MA15) strain as a surrogate to assess the *in vivo* efficacy of these
322 drugs. We are of the opinion that this is a good model since both viruses use ACE2 as a
323 receptor (18–21) and therefore have a similar cellular tropism which is important since both of
324 these compounds appear to inhibit viral entry. Prophylactic dosing in MA15 infection
325 experiments demonstrated that, in contrast to the *in vitro* antiviral activity, CQ and CPZ did no
326 inhibit viral replication in mouse lungs based on viral titer recovered from lungs at 4 dpi.

327 However, both drugs resulted in reduced weight loss and improved clinical outcome, with the
328 higher dose giving greater protection. Along with being an anti-malarial, CQ is used in humans
329 for the treatment of systemic lupus erythematosus and rheumatoid arthritis because of anti-
330 inflammatory properties and has effects on antigen presentation (22–24). We speculate that
331 these properties may have a role in the protection we observe *in vivo* since much of the
332 pathology from SARS-CoV is a consequence of immunopathology during infection (in mice;
333 (25), in non-human primates (26) and for a detail review (27)). These results suggest that CQ
334 alone may not be a viable therapeutic but may be beneficial for treatment of SARS-CoV-2 in
335 combination with more directly acting antivirals such as remdesivir (28–30).

336
337 The development of antiviral drugs for emerging coronaviruses is a global priority. In the middle
338 of the COVID19 pandemic, we must identify rapidly accessible therapeutics that are validated in
339 both *in vitro* and *in vivo* models. FDA approved drugs being assessed for repurposing and other
340 experimental drugs in development must be properly validated in animal studies to best assess
341 their potential utility in people. We have presented here a list of FDA approved drugs that are
342 effective *in vitro* against SARS-CoV-2 as well as being effective against SARS-CoV and MERS-
343 CoV (6). Moreover, we have demonstrated that two of these, CQ and CPZ, can protect mice
344 from severe clinical disease from SARS-CoV. Future research will be aimed at testing these
345 compounds in SARS-CoV-2 animal models to further assess their potential utility for human
346 treatment.

347

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351

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457

458

459 **Figure legend**

460 **Figure 1 – Percentage inhibition and percentage cytotoxicity graphs from drug screens**

461 **starting at 50 μ M using an 8-point, 1:2 dilution series.** Results from one representative drug

462 screen of three showing percentage inhibition and cytotoxicity for each of the tested drugs.

463 Triplicate wells of cells were pre-treated with the indicated drug for 2 hours prior to infection with

464 SARS-CoV-2 at MOI 0.01. Cells were incubated for 72 hours prior to performing CellTiter-Glo

465 assays to assess cytopathic effect. Data are scored as percentage inhibition of relative cell

466 viability for drug treated versus vehicle control. Data are the mean percentages with error bars

467 displaying standard deviation between the triplicate wells.

468

469 **Figure 2 – Hydroxychloroquine and chloroquine phosphate inhibit production of SARS-**

470 **CoV-2 N and RdRp mRNA.**

471 Vero cells were pre-treated with hydroxychloroquine (A, C and E) or chloroquine sulfate (B, D

472 and F) at the indicated concentration (or 0.1% water as vehicle control) for 2 h prior to infection

473 with SARS-CoV-2 (WA-1 strain) at MOI 0.1. 24 h post-infection cells were collected in TRIzol.

474 RNA was extracted from TRIzol sample and qRT-PCR was performed for viral RdRp (A and B)

475 or N (C and D) mRNA using WHO primers. RNA levels were normalized with 18S RNA and fold

476 change for drug treated to vehicle control was calculated (dotted line to denote a fold change of

477 1 which is no change over control). Data are from 3 independent infections performed on

478 triplicate wells, the fold change was calculated in each independent experiment and the mean

479 fold change is plotted with error bars displaying standard deviation. Along with TRIzol samples

480 for RNA supernatant was collected from cells and used for TCID₅₀ assays to determine

481 infectious virus production following treatment with HCQ (E) or CQ (F) Data are from 3

482 independent infections performed on triplicate wells with the TCID₅₀/ml being averaged across

483 all wells. Error bars are the standard deviation. G) Cells were treated with 50 μ M HCQ or 0.1%
484 water as control. Drug was either added 2 h prior to infection, at the time of infection or 2 h after
485 infection with MOI 0.1 SARS-CoV-2. After 24 h infection, supernatant was collected and used
486 for TCID₅₀ assays to determine infectious virus production. Data are from 3 independent
487 infections performed on triplicate wells with the TCID₅₀/ml being averaged across all wells. Error
488 bars are the standard deviation. H) SARS-CoV spike psuedoviruses (PV) were used for
489 infection of BSC1 cells. The cells were treated with 10 μ M of CQ or CPZ for 1 h prior to infection
490 with PV for 3 h. The PV carry BlaM and cells were loaded with CCF2 to monitor cleavage and
491 shift in fluorescence output for evidence of S-mediated entry into cells. Data are normalised to
492 PV alone and are from 3 independent experiments with error bars representing standard
493 deviation.

494

495 **Figure 3 – Antiviral activity of additional FDA approved compounds against SARS-CoV-2.**

496 Other drugs that showed antiviral activity in our initial CellTiter-Glo screening were tested for
497 inhibition of productive virus infection. Cells were treated with the indicated concentrations of A)
498 amodiaquine dihydrochloride dihydrate, B) amodiaquine hydrochloride, C) chlorpromazine, D)
499 mefloquine and E) imatinib for 2 h prior to infection with SARS-CoV-2 at MOI 0.1 for 24 h.
500 Supernatant was collected and used for TCID₅₀ assay to quantify infectious virus production.
501 Data are from a representative experiment of four performed on triplicate wells. Data are the
502 mean TCID₅₀/ml with error bars being standard deviation.

503

504 **Figure 4 – CQ and CPZ are protective against SARS-CoV (MA15) infection *in vivo***

505 Mice were treated with CQ or CPZ 1 day prior to infection with SARS-CoV (MA15) and dosed
506 with each drug across the 4 day infection time course. Water was used as the vehicle control for
507 both drugs and PBS was used as a control for uninfected mice. A) Weight loss of mice treated
508 with CQ at two different dose levels (0.8 mg and 1.6 mg) over the 4 day infection. Data are

509 presented as relative weight loss compared to the mouse weight on day 0. In each treatment
510 group there were 5 mice and the data are mean average and standard deviation. B) At day 4,
511 mice were euthanized and lung sections were used for H&E staining. C) In addition to collecting
512 lungs for section staining, there was also collection to determine titer of virus by plaque assay.
513 D) Weight loss of mice treated with CPZ at three different doses (20 µg, 100 µg, and 200 µg)
514 with the same experimental set up as in A. E and F) As B and C but for CPZ treated mice.

515

516 **Table 1** - IC50 and CC50 values for 20 FDA approved drugs against SARS-CoV-2.

517 Abbreviations: MOI (multiplicity of infection), IC50 (half maximal inhibitory concentration), CC50
518 (half maximal cytotoxic concentration), avg. (average), ND (not determined).

519 A – Run totals listed as IC50,CC50

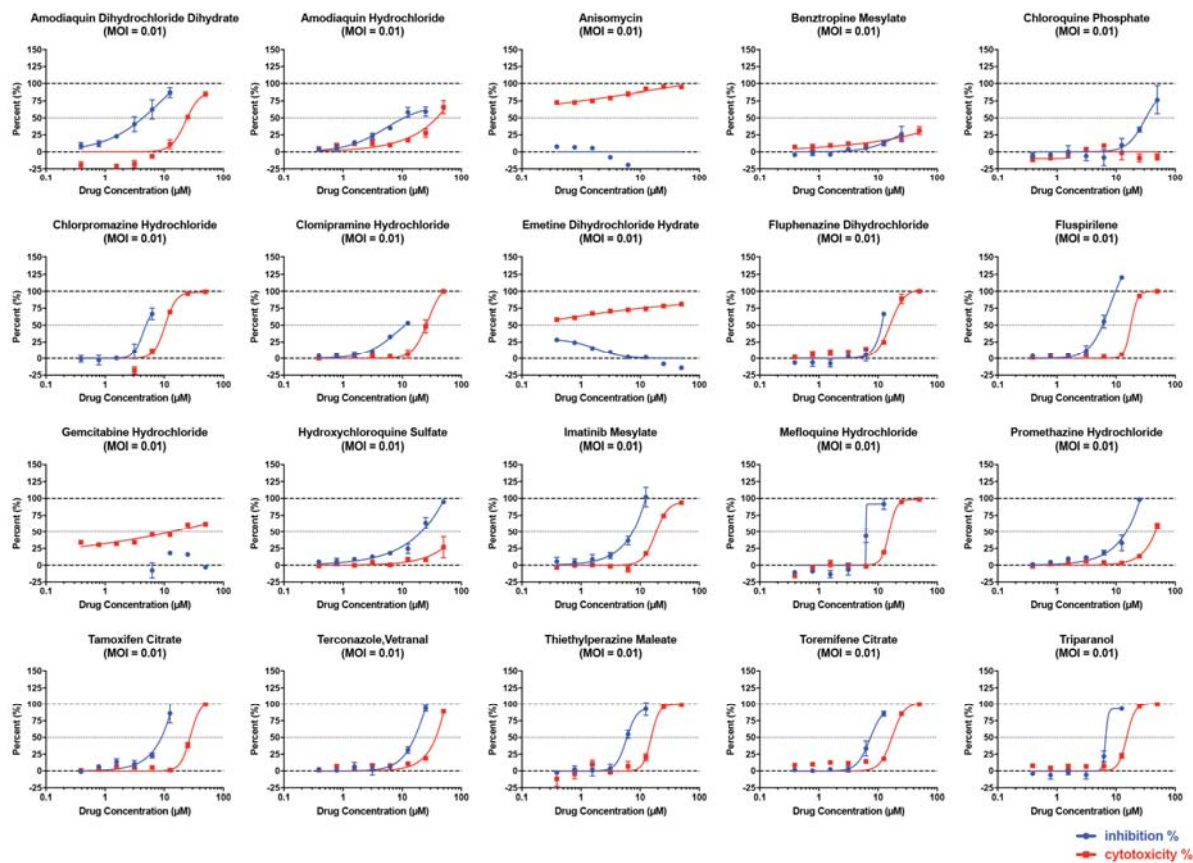
520 B – At least one CC50 could be extrapolated from the curve fit suggesting toxicity and SI are
521 slightly higher than listed

522 C – No CC50 could be extrapolated from the curve fit suggesting toxicity and SI are much
523 higher than listed.

524

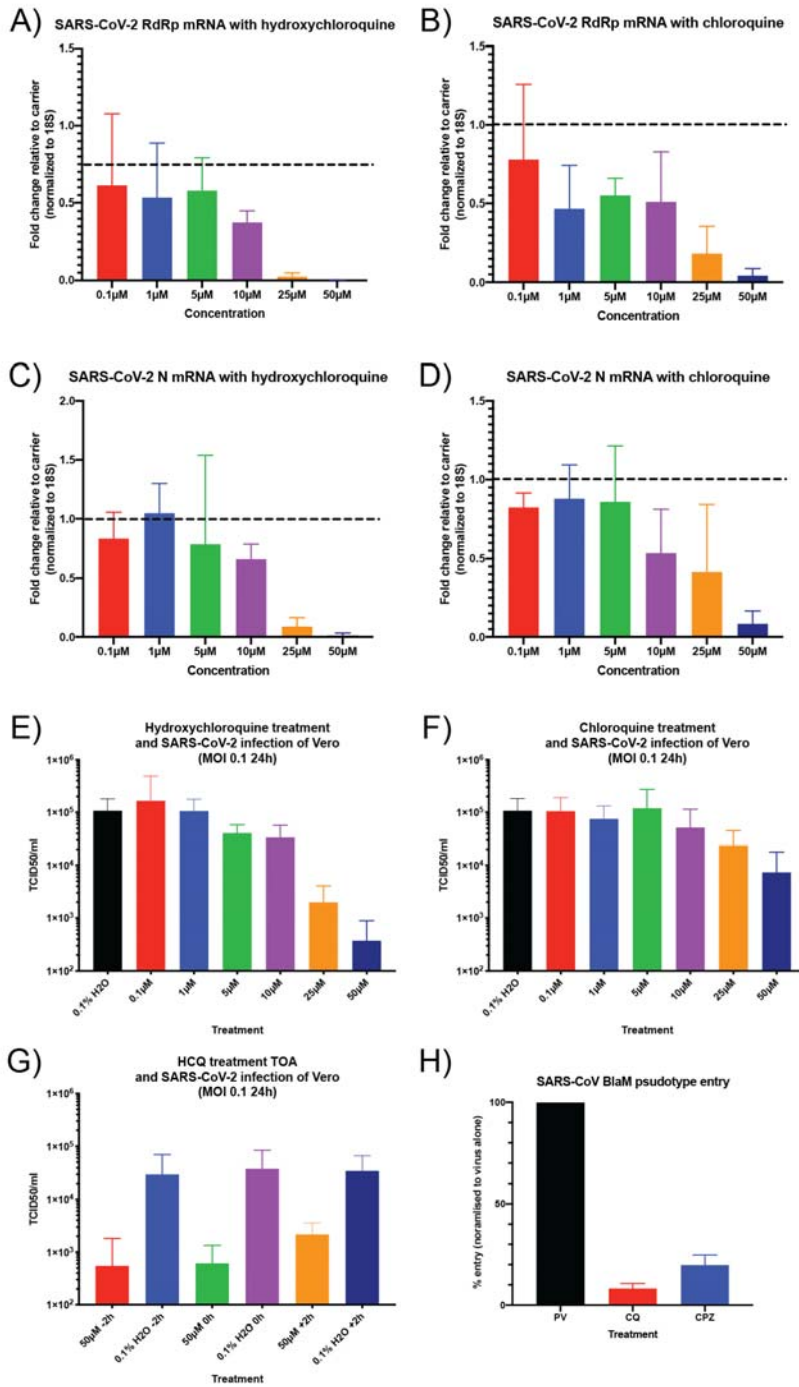
525 **Figure 1**

526



527

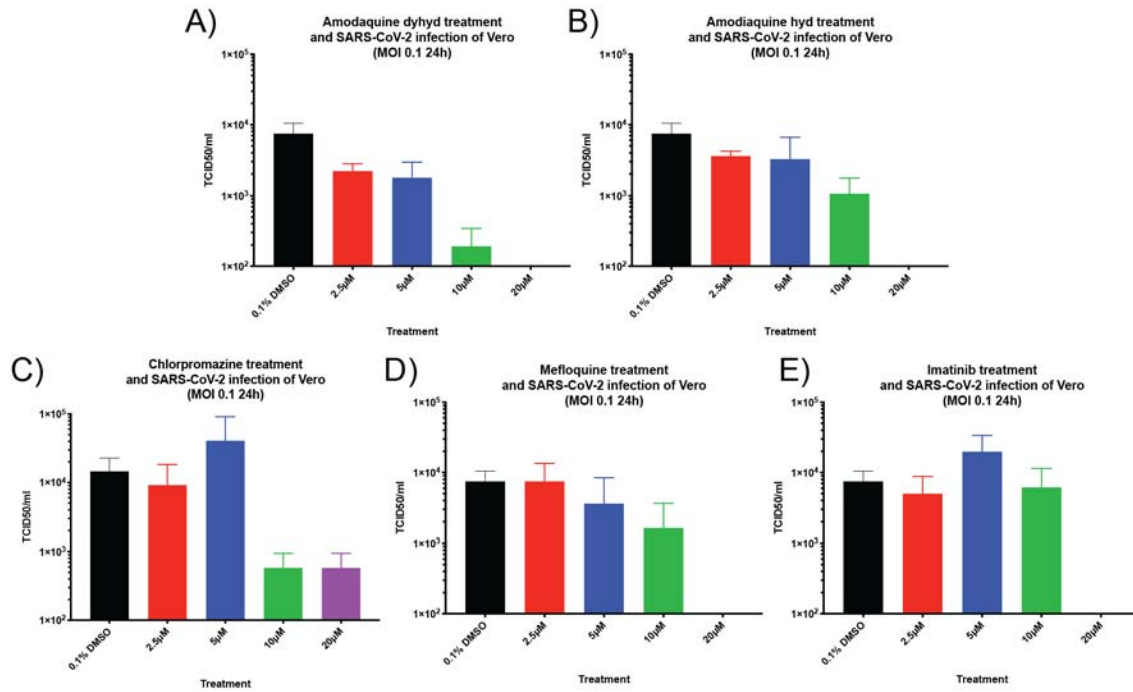
528 **Figure 2**



529

530

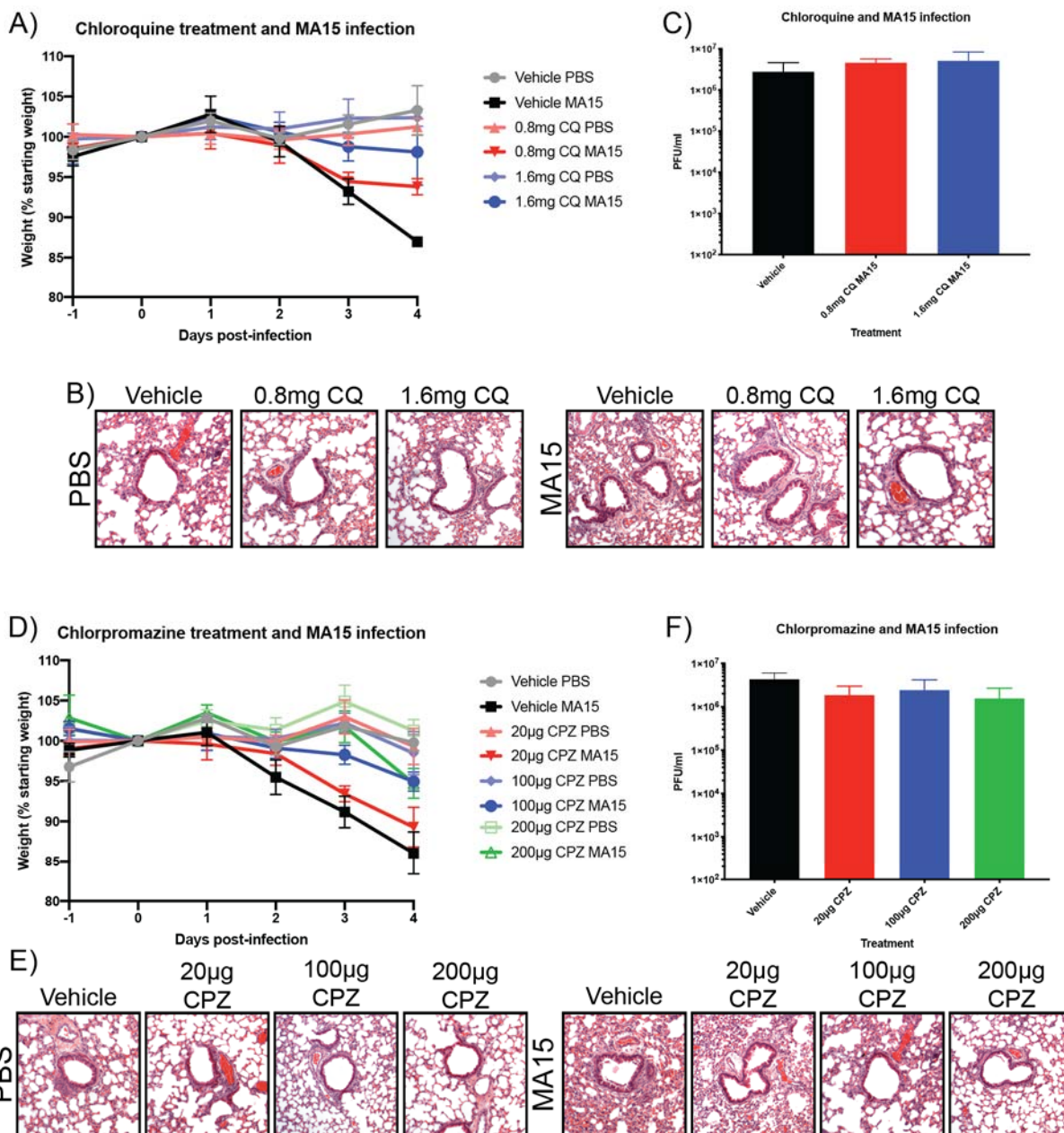
531 **Figure 3**



532

533

534 **Figure 4**



535