The adenosine analogue prodrug ATV006 is orally bioavailable and has potent

1

2 3	preclinical efficacy against SARS-CoV-2 and its variants
4	Liu Cao <sup>1, #</sup> , Yingjun Li <sup>2, #</sup> , *, Sidi Yang <sup>1, #</sup> , Guanguan Li <sup>2, 3, #</sup> , Qifan Zhou <sup>2, #</sup> , Jing Sun <sup>4</sup> ,
5	Tiefeng Xu <sup>1</sup> , Yujian Yang <sup>2</sup> , Tiaozhen Zhu <sup>2</sup> , Siyao Huang <sup>1</sup> , Yanxi Ji <sup>1</sup> , Feng Cong <sup>5</sup> ,
6	Yinzhu Luo <sup>5</sup> , Yujun Zhu <sup>5</sup> , Hemi Luan <sup>6</sup> , Huan Zhang <sup>7</sup> , Jingdiao Chen <sup>7</sup> , Xue Liu <sup>1</sup> , Ping
7	Wang <sup>2</sup> , Yang Yu <sup>2</sup> , Fan Xing <sup>1</sup> , Bixia Ke <sup>7</sup> , Huanying Zheng <sup>7</sup> , Xiaoling Deng <sup>7</sup> , Wenyong
8	Zhang <sup>6</sup> , Chun-Mei Li <sup>1</sup> , Yu Zhang <sup>5</sup> , Jincun Zhao <sup>4</sup> , Xumu Zhang <sup>2, 3, *</sup> , Deyin Guo <sup>1, *</sup>
9	
10	Affiliations:
11	<sup>1</sup> Centre for Infection and Immunity Studies (CIIS), School of Medicine, Shenzhen
12	Campus of Sun Yat-sen University, Guangdong 518107, China.
13	<sup>2</sup> Shenzhen Key Laboratory of Small Molecule Drug Discovery and Synthesis,
14	Department of Chemistry, College of Science, Academy for Advanced
15	Interdisciplinary Studies, Southern University of Science and Technology, Shenzhen,
16	Guangdong 518055, China
17	<sup>3</sup> Medi-X Pingshan, Southern University of Science and Technology, Shenzhen,
18	Guangdong 518118, China
19	<sup>4</sup> State Key Laboratory of Respiratory Disease, National Clinical Research Center for
20	Respiratory Disease, Guangzhou Institute of Respiratory Health, the First Affiliated
21	Hospital of Guangzhou Medical University, Guangzhou, Guangdong 510182, China
22	<sup>5</sup> Guangdong Province Key Laboratory of Laboratory Animals, Guangdong Laboratory
23	Animals Monitoring Institute, Guangzhou, Guangdong 510663, China.
24	<sup>6</sup> School of Medicine, Southern University of Science and Technology, Shenzhen,
25	Guangdong 518055, China
26	<sup>7</sup> Center for Disease Control and Prevention of Guangdong Province, Guangzhou,
27	Guangdong 511430, China
28	
29	

30 \* Correspondence to: Deyin Guo (guodeyin@mail.sysu.edu.cn), Xumu Zhang
31 (zhangxm@sustech.edu.cn) and Yingjun Li (liyj@sustech.edu.cn).

32 # These authors contributed equally to this work.

33 Abstract

34 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which causes the 35 COVID-19 pandemic, is rapidly evolving. Due to the limited efficacy of vaccination in 36 prevention of SARS-CoV-2 transmission and continuous emergence of variants of 37 concern (VOC), including the currently most prevalent Delta variant, orally 38 bioavailable and broadly efficacious antiviral drugs are urgently needed. Previously we 39 showed that adenosine analogue 69-0 (also known as GS-441524), possesses potent 40 anti-SARS-CoV-2 activity. Herein, we report that esterification of the 5'-hydroxyl 41 moieties of 69-0 markedly improved the antiviral potency. The 5'-hydroxyl -isobutyryl 42 prodrug, ATV006, showed excellent oral bioavailability in rats and cynomolgus 43 monkeys and potent antiviral efficacy against different VOCs of SARS-CoV-2 in cell 44 culture and three mouse models. Oral administration of ATV006 significantly reduced 45 viral loads, alleviated lung damage and rescued mice from death in the K18-hACE2 46 mouse model challenged with the Delta variant. Moreover, ATV006 showed broad 47 antiviral efficacy against different mammal-infecting coronaviruses. These indicate that 48 ATV006 represents a promising oral drug candidate against SARS-CoV-2 VOCs and 49 other coronaviruses.

50 Keywords:

51 SARS-CoV-2, Delta variant, antiviral drug, nucleoside analogue, ATV006,
52 Coronavirus.

53 Introduction

The outbreak of COVID-19 pandemic, caused by SARS-CoV-2, has been continuing for over one year and has resulted in over 228 million confirmed infections and over 4 million reported deaths worldwide as of 20 September 2021 (WHO 2021b). SARS-CoV-2 is a positive-sense, single-stranded RNA virus belonging to the genus 58 Betacoronavirus of the family Coronaviridae (Chen et al. 2020). Two other members 59 of the same genus, namely severe acute respiratory syndrome coronavirus (SARS-CoV) 60 and Middle East respiratory syndrome coronavirus (MERS-CoV), have caused 61 outbreaks with substantial fatality rates in 2002 and 2012, respectively (Al-Tawfig et 62 al. 2014; Perlman et al. 2009). Given the repeated and accelerating emergence of highly 63 pathogenic coronaviruses, it is increasingly important to develop broadly effective anti-64 coronaviral agents to combat the pandemics of COVID-19 and the future emerging 65 CoVs.

66 Although the coronavirus has a certain proofreading ability (Robson et al. 2020), it 67 still has a high mutation rate. Progressive mutational change in the virus is therefore 68 inevitable after a number of replicative cycles, which leads to the emergence of new 69 variants. At present, the WHO has classified many variants of the SARS-CoV-2 (WHO 70 2021a), including the variants of concern (VOC), such as the Alpha variant (B.1.1.7), 71 Beta variant (B.1.351) and Delta variant (B.1.671.2). Delta variant is highly contagious, 72 and has rapidly become the prevalent variant, contributing to the current wave of 73 pandemic in India and worldwide and leading to vaccine breakthrough infections 74 associated with higher viral load and long duration of shedding (Reardon 2021). These 75 SARS-CoV-2 variants reduced the vaccine effectiveness and antibody protection 76 (Harvey et al. 2021; Liu et al. 2021; Tegally et al. 2021). Currently available directly-77 acting antiviral drugs repurposed for treatment of COVID-19 patients showed limited 78 efficacy in large-scale clinical trials (Consortium et al. 2021). Therefore, it is in urgent 79 need to develop orally available and broadly efficacious anti-coronaviral agents against 80 SARS-CoV-2 and its variants.

81 The non-structural protein 12 (nsp12) of SARS-COV-2 acts as the RNA-dependent 82 RNA polymerase (RdRp), which catalyzes RNA template-dependent formation of 83 phosphodiester bonds between ribonucleotides using ribonucleoside triphosphates as 84 substrates, serving as the key component of the replication/transcription machinery

85 (Hillen et al. 2020; Wang et al. 2020b). RdRps are considered as the primary targets for 86 antiviral drug development in a wide variety of viruses (Kabinger et al. 2021; Picarazzi 87 et al. 2020). Several nucleoside or nucleotide analogues, including remdesivir, AT-527, 88 favipiravir and molnupiravir (EIDD-2801), originally developed by targeting the RdRp 89 of other RNA viruses, have been repurposed for SARS-CoV-2 since the COVID-19 outbreak (Ghasemnejad-Berenji et al. 2021; Goldman et al. 2020; Good et al. 2021; 90 91 Wahl et al. 2021; Wang et al. 2020a). Until now, remdesivir is the only FDA-approved 92 antiviral drug for the treatment of COVID-19 patients (Beigel et al. 2020). However, remdesivir is administered by intravenous (IV) injection and thus has a clinical 93 94 application limited to hospitalized patients with relatively advanced disease (Alanazi et 95 al. 2019; Beigel et al. 2020; Consortium et al. 2021). We and others reported that the 96 parent nucleoside of remdesivir, with CAS registry number 1191237-69-0 (named as 97 69-0, also known as GS-441524), potently inhibits SARS-CoV-2 infection in cell 98 culture and mouse models infected with SARS-CoV-2 (Li et al. 2021; Pruijssers et al. 99 2020). 69-0 is a 1'-cyano-substituted adenosine analogue with broad-spectrum antiviral 100 activities across multiple virus families (Cho et al. 2012; Huang et al. 2021; Murphy et 101 al. 2018; Pedersen et al. 2019). However, 69-0 suffers from its disadvantage of poor 102 solubility (water solubility of  $< 1 \mu g/mL$ ) and poor oral pharmacokinetic (PK) profile 103 (Li et al. 2021). In non-human primates, 69-0 has an oral bioavailability of 8.3% and 104 poor plasma exposure, preventing it from further development into an oral drug 105 (NCATS 2021).

In this study, we reported a series ester prodrugs of adenosine analogues 69-0 with improved antiviral potency. The isobutyryl nucleoside derivative of 69-0, named as ATV006, with improved oral PK profiles, potently inhibited the replication of SARS-CoV-2 and other CoVs. In three different animal models, ATV006 could efficaciously suppress the infection and pathogenesis of SARS-CoV-2 and its variants including the most prevalent Delta variant. These results indicate that ATV006 represents a 112 promising drug candidate for the further clinical development against COVID-19 and

113 other CoV diseases.

114 **Results** 

#### 115 Design and synthesis of prodrugs of adenosine analogs that target SARS-CoV-2

116 **RNA polymerase** 

117 Although the adenosine analogue 69-0 was effective against SARS-CoV-2 in vitro 118 and in mouse models, it suffers from poor oral bioavailability, which hampers its further 119 development as oral drug as shown in our previous study(Li et al. 2021). To overcome 120 this limitation, we devoted our efforts to synthesize 69-0 prodrugs by employing short 121 chain fatty acid (SCFA) or amino acid to mask the polar hydroxyl- or animo-groups. 122 For this, 21 compounds with different substitutions at the positions of  $\mathbf{R}^1$ ,  $\mathbf{R}^2$ ,  $\mathbf{R}^3$  and 123  $\mathbf{R}^4$  of 69-0 were designed and chemically synthesized (Fig 1; supplementary materials). 124 In brief, the compounds ATV001-004 were synthesized from 69-0 via one-step 125acylation reaction with related acid anhydride in the presence of 126 dimethylaminopyridine (DMAP) and ethylene glycol dimethacrylate (EDMA). To 127 synthesize 5'-hydroxyl-acetylated compounds, the 2',3'-hydroxyl moieties of 69-0 were 128 firstly protected with acetonide in the presence of sulfuric acid. Then, different aliphatic 129 acids, amino acids or benzoic acid were selectively coupled with the free hydroxyl 130 group on C5' position to produce the corresponding esters. Final deprotection of the 131 acetonide with 6N hydrochloride acid (HCl) afforded the target compounds ATV005-132 024. All the compounds were purified by high performance liquid chromatography 133 (HPLC), reaching a purity of above 95%.

The antiviral effect of the compounds was initially evaluated by using a biosafe SARS-CoV-2 replicon system (pBAC- SARS-CoV-2-Replicon-Luc), established in our previous work (Jin et al. 2021), which carries all the genes essential to genome replication including that for RdRp and the luciferase reporter gene but does not produce infectious virus particles. We first tested the percentage inhibition of the 139 replicon replication at 10 µM of each compound (Fig S1A) and then selected 17 140 compounds to measure their concentration for 50% of maximal effect (EC<sub>50</sub>) value, 141 which ranged from 0.217 to 2.351 µM (Fig S1B). As shown in Fig S1A and S1B, the 142 compounds ATV001 and ATV002 with isobutyryl amide or acetyl amide at the base 143 moiety showed decreased antiviral activities probably due to the biostable amide group 144 that obstructs the hydrogen bond formation between inhibitor and RNA template 145 (Kokic et al. 2021; Wang et al. 2020b). Tri-esterification of the hydroxyl groups on C5' 146  $(\mathbf{R}^1)$ , C2'  $(\mathbf{R}^2)$  and C3'  $(\mathbf{R}^3)$  positions (ATV003-004) did not significantly change the activity while the mono-isobutyryl- modification of 5'-hydroxyl group (ATV006) 147 148 markedly improved the inhibitory activities in the replicon system ( $EC_{50}$  value of 0.52 149  $\mu$ M, about one-fold more potent compared to parent 69-0). We then kept the 2' and 3' hydroxyl group unchanged, while the R<sup>1</sup> group was replaced with straight cyclic, 150151 branched SCFA, benzyl acyl- group or amino acid- group (ATV007-024). We found 152 that some of the SCFA ester prodrug compounds displayed an improvement in potency 153relative to 69-0. Six compounds (ATV019-024) bearing L- or D- amino acid ester were 154 designed to improve drug absorption by targeting the peptide transporter family 1 155 (PepT1) (Zhang et al. 2013). However, these compounds did not show improved 156 activities against the replicon or improved permeabilities in Caco-2 cells (Table S1).

Together, SCFA esterification on C5' position could generally improve the potency relative to 69-0. Then we selected six compounds from the SCFA group for further analysis of anti-SARS-CoV-2 activity with the live viruses of different variants in cell culture and animal models (Table 1).

### The adenosine analog prodrug ATV006 potently inhibits the replication of SARS CoV-2 and its variants of concern

163 The antiviral efficacy of the six SCFA prodrugs (ATV006, ATV009-011, ATV013 164 and ATV017) was first evaluated in a cell culture model infected with different strains 165 of SARS-CoV-2, including the early strain B.1, and two prevalent SARS-CoV variants 166 of concern (VOC), the Beta (B.1.351) and Delta (B.1.671.2) variants (Fig 2, Table 1). 167 The Vero E6 cells were treated with the compounds and then infected with SARS-CoV-168 2 at a multiplicity of infection (MOI) of 0.05, and the copy number of viral genome 169 RNA in the cell culture supernatant was measured by the quantitative real-time 170 polymerase chain reaction (qRT-PCR) 48 h post infection (hpi). As shown in Figure 2 171 and Table 1, the compounds showed improved potency against SARS-CoV-2 relative to remdesivir and 69-0, which was in consistency with the results of the SARS-CoV-2 172173replicon system. Among them, ATV010 exhibited low micromolar EC<sub>50</sub> value with 174 early strain B.1 and Beta variant, while ATV006 had an overall >4-fold potency 175improvement in inhibiting the replication of Delta variant, with EC<sub>50</sub> reaching 0.3485 176 $\mu$ M. We repeated the antiviral experiment of the compounds in Huh7 cells with B.1 177strain, and the results showed that these compounds exhibited similar antiviral activity 178 to that in Vero E6 cells (Fig S2). Together, these results indicate that SCFA-esterified 179 compounds could potently inhibit the replication of SARS-CoV-2 and its variants.

The cytotoxicity of the compounds was evaluated on Vero E6 cells with the CCK8 assays (Fig S3). The results showed that most of the compounds had low toxicity with  $CC_{50} > 50 \mu$ M, except for ATV010 with  $CC_{50}$  of 44.62  $\mu$ M, suggesting that most compounds had excellent safety. The therapeutic index ( $CC_{50}$ /  $EC_{50}$ ) of ATV006 was as high as 367 in Vero E6 cells. Considering the potent inhibition against Delta variant and high selectivity against cell proliferation, ATV006 was selected for further studies.

#### 186 **Pharmacokinetic properties of ATV006 in rats and cynomolgus monkeys.**

To assess the oral absorption of ATV006, the pharmacokinetic (PK) studies were conducted in rats and monkeys. Following oral dosing of 20 mg/kg in rats, ATV006 displayed high oral bioavailability (F%) of 98%, using the parent nucleoside 69-0 as an analyte (Table 2). The  $C_{max}$  of 11445 µg/L was achieved 0.83 h after the oral administration, indicating its effective blood exposure. Plasma concentration decayed with a half-life of 3.62 h (Fig 3A). In cynomolgus monkeys,  $C_{max}$  of averaged 2715 193 µg/L was reached in 1.5 h after the IG administration of 10 mg/kg ATV006. Plasma 194 concentration decayed with a  $T_{1/2}$  of 4 h and the oral bioavailability was about 30% (Fig. 195 3B). As a compound with an oral bioavailability of >10% has the potential for 196 development as an oral drug (Martin 2005), ATV006 well met such standard as an oral 197 drug candidate for further testing in animal models. Next, we explored the tissue 198 distribution of ATV006 in the mouse model by measuring its parent nucleoside 69-0. The results revealed that oral administration of ATV006 achieved a broad distribution 199 200 in liver and kidney as well as in lung, the major target organ to SARS-CoV-2 infection, 201 indicating its potential of an oral drug for the treatment of COVID-19 (Fig 3C).

## Orally administered ATV006 could effectively suppress SARS-CoV-2 replication in mouse models

204 We next investigated the in vivo antiviral activity of orally administered ATV006 in 205two different mouse models of SARS-CoV-2 (strain B.1) infection, one with 206 humanized angiotensin-converting enzyme 2 (hACE2) (Sun et al. 2020b)and the other 207 with adenovirus-delivered human ACE2 (Ad5-hACE2) (Sun et al. 2020a). The hACE2 transgenic mice were intranasally inoculated with SARS-CoV-2 (2x10<sup>5</sup> plaque forming 208 209 units (PFU) per mouse) and were treated with vehicle (control, n=6), ATV006 (500 210 mg/kg, IG, once daily, n=6) or ATV006 (250 mg/kg, IG, once daily, n=6), starting at 2 211 h prior to virus inoculation (Fig 4A) and continuing until 4 days post-infection. To 212 better determine the replication levels of SARS-CoV-2, we detected both the genomic 213 RNA (gRNA) and subgenomic RNA (sgRNA), the latter being produced by 214 discontinuous synthesis and representing a biomarker of coronavirus replication 215 (Hussain et al. 2005; Kim et al. 2020) (Fig S4). In the control group, both gRNA and 216 sgRNA of SARS-CoV-2 reached high levels in the lung, indicating that the mouse 217 infection model was well established. In contrast, SARS-CoV-2 RNAs were hardly 218 detectable at day 4 in the ATV006 treatment groups (Fig 4B and 4C), demonstrating 219 robust inhibition of SARS-CoV-2 replication by ATV006.

220 We further tested the antiviral potency of ATV006 in the Ad5-hACE2 mouse model, 221 which supports SARS-CoV-2 infection and pathogenesis in mouse lung (Sun et al. 222 2020a). The mice were inoculated intranasally with  $1 \times 10^5$  PFU virus per mouse and 223 were then treated with vehicle (control, n=8) or ATV006 (250 mg/kg, IG, once daily, 224 n=8) and EIDD-2801 (500 mg/kg, IG, once daily, n=8) starting at 1 day prior to virus 225 inoculation (Fig 4D). EIDD-2801 was previously shown to effectively inhibit SARS-226 CoV-2 replication at 500 mg/kg dosage (Wahl et al. 2021) and used as a positive control. 227 The virus titers were measured, and results showed that ATV006 (250 mg/kg) and 228 EIDD-2801 (500 mg/kg) could significantly reduce the viral load and pathological 229 damage of the lung (Fig 4E and 4F). Together, ATV006 showed potent anti-SARS-230 CoV-2 efficacy in different mouse models.

### ATV006 reduces lung damage and protects mice from death by infection of the Delta variant in the K18-hACE2 transgenic mice

233 We next tested the antiviral potency in the K18-hACE2 transgenic mice, which are 234 susceptible to SARS-CoV-2 and can lead to death of infected mice (Oladunni et al. 2352020). As the Delta variant of SARS-CoV-2 is the most prevalent variant, we 236 specifically tested the efficacy of ATV006 against the Delta variant. The K18 hACE2 237 mice were intranasally inoculated with SARS-CoV-2 Delta variant (1x10<sup>4</sup> PFU virus per mouse) and then treated with vehicle (control, n=11), ATV006 (250 mg/kg, IG, 238 239 once daily, n=11), ATV006 (100 mg/kg, IG, once daily, n=8) or EIDD-2801 (500 240 mg/kg, IG, once daily, n=8) starting at 2 h prior to virus inoculation (Fig 5A) and 241 continuing until 5 days post-infection (dpi). During the 9-day observation period, the 242 mice in the control group gradually lost weight from the fourth day and died from the 243 sixth day, and all died on the seventh day, but all mice of treatment groups survived 244 (Fig 5B and 5C). At 3 dpi, we evaluated the abundance of both the viral gRNA of 245 SARS-CoV-2 in the mouse lung and brain tissue by qPCR. The amount of viral RNAs 246 of the treatment groups was significantly lower than that of the control group, with

ATV006 (250 mg/kg) group having the strongest potency with reduction of viral RNA
for more than 10,000 times in the lung (Fig 5D).

249 Histopathological analysis and observation were performed with the lungs of the 250 mice infected with SARS-CoV-2 at 3 dpi. The vehicle-treated mice showed multiple 251injuries, including inflammatory cell infiltration ranging from the trachea, peri-alveolar 252space, to the interstitium whereas ATV006-treated animals had markedly alleviated 253 symptoms in the lungs (Fig 5E and 5F). Compared with the ATV006 treatment group, 254the spleen of the mice in the control group was significantly enlarged, and the white 255pulp was atrophied to varying degrees (Fig 5G and 5H). Furthermore, ATV006 256 markedly reduced the production of inflammatory cytokines and chemokines in the 257 lung tissues (Fig 5I).

Together, our results showed that intragastric administration of ATV006 could efficiently inhibit SARS-CoV-2 replication, ameliorate SARS-CoV-2-induced lung lesions in vivo and prevent death of the mice infected by the Delta variant of SARS-CoV-2. These demonstrated the potential of ATV006 as an orally bioavailable anti-SARS-CoV-2 drug.

#### 263 **ATV006** possesses broad antiviral activities against diversified coronaviruses.

264 To explore the broad-spectrum antiviral activity of ATV006, we further tested it 265 against other coronaviruses of the genera Alphacoronavirus and Betacoronavirus that 266 are known to infect humans (Chen et al. 2020; Fung et al. 2019). Six coronaviruses 267 were selected, including mouse hepatitis virus (MHV), feline infectious peritonitis virus 268 (FIPV), porcine epidemic diarrhea virus (PEDV), canine coronavirus (CCoV), 269 transmissible gastroenteritis virus (TGEV) and swine acute diarrhea syndrome 270 coronavirus (SADS-CoV). Among these coronaviruses, MHV is a beta-coronavirus 271that is distantly related to human coronaviruses SARS-CoV-2, SARS-CoV and MERS-272CoV.

We first compared the antiviral activities of ATV006 with that of remdesivir and 69-273 274 0 in MHV cell culture. The mouse L2 cells were infected with MHV-A59, a strain that 275 infects the liver and brain of mice and causes acute hepatitis, encephalitis, and chronic 276 demyelinating disease (Weiss et al. 2011), at a MOI of 0.1 and treated with different 277 dilutions of the compounds. Antiviral activities were evaluated by qRT-PCR 278 quantification of the viral copy number in the culture supernatant and intracellular 279 fraction after 16 hpi. ATV006 showed robust anti-MHV activity (EC<sub>50</sub> =  $0.265 \mu$ M) 280 compared to remdesivir (EC<sub>50</sub> = 1.338  $\mu$ M) and 69-0 (EC<sub>50</sub> = 0.874  $\mu$ M) in L2 cells 281 (Fig 6A). Then we tested the antiviral activity of ATV006 against FIPV, CCoV, PEDV, 282 TGEV and SADS-CoV, and the EC<sub>50</sub> was 1.040 µM, 0.186 µM, 1.040 µM, 3.045 µM 283 and 2.490 µM, respectively, in cell culture models (Fig 6B-6F). These results indicate 284 that ATV006 has a broad-spectrum anti-coronavirus efficacy.

285 We further evaluated the anti-coronavirus activity of ATV006 in MHV infection 286 mouse model. The Balb/c mice were inoculated intranasally with 1 x 10<sup>6</sup> PFU per 287 mouse of MHV-A59, and treated with vehicle (control, n=5) or ATV006 (500, 250, 288 100, 50 mg/kg, IG, twice daily, n=5), starting at 3 hours prior to virus inoculation and 289 continuing until 2 days post-infection (Fig 7A). We found that ATV006 treatments at 290 different dosages could prevent the weight loss of the mice (Fig 7B). At 2 dpi, we 291 measured the viral gRNA and sgRNA copy number and virus titer in mouse lung and 292 liver (Fig S4 and Fig 7). It was found that the treatments of high dosages of ATV006 293 (250, 500 mg/kg) could effectively inhibit virus replication both in the lung and liver, 294 and the 500 mg/kg group had 99% inhibition of virus replication in the lung and liver 295 (Fig 7C-7F) while the treatment of low dosages of ATV006 (50, 100 mg/kg) 296 significantly inhibited viral replication in the liver, but not significantly in the lung (Fig. 297 7C-7F). Histopathological analysis demonstrated that the vehicle-treated mice showed 298 inflammatory cell infiltration, whereas ATV006 (500 mg/kg)-treated mice had 299 alleviated symptoms in the lungs at 2 dpi (Fig 7G). In addition, ATV006 significantly 300 reduced the production of inflammatory cytokines and chemokines, such as IL-6, IL-

301 1β and IFN- $\gamma$  (Fig 7H). Due to the suppression of virus replication, IFN- $\beta$  and ISGs 302 (CXCL10) also significantly decreased (Fig 7H).

303 We next explored the lowest oral dosages of ATV006 that could protect MHV-A59-304 infected mice from death and weight loss. As shown in Figure S5, ATV006 could 305 prevent mouse death and weight loss of mice at dosages of 5-50 mg/kg while the mouse 306 treated with 2 mg/kg of ATV006 began to die 4 days post infection (dpi) and all mice 307 died till 10 dpi. In comparison, the mice of the control group all died at 8 dpi and even 308 the treatment with 2 mg/kg ATV006 prolonged the survival time and reduced the viral 309 load in the liver by about six times (Fig S5B-S5D). IP administration of remdesivir (20 310 mg/kg) and IG administration of 69-0 (50 mg/kg) could also prevent death of the MHV-311infected mice (Fig. S5) but the body weight of the mice was reduced in comparison 312 with the mice treated with ATV006.

313 MHV mainly infects the liver of mouse and causes hepatitis. Therefore, we also 314 performed intrahepatic (IH) inoculation to directly observe whether ATV006 had the 315 effect of inhibiting virus replication in the liver and alleviating the symptoms of 316 hepatitis (Fig S6A). Intriguingly, compared with the vehicle group, mice treated with 317 the lowest concentration (2 mg/kg) had a 100-fold reduction in virus replication by 318 measurement of the viral RNA load by qPCR, or the virus titer by plaque assay (Fig 319 S6B, S6C and S6D). The serum ALT and AST values of ATV006-treated mouse were 320 significantly lower than that of the control group (Fig S6E and S6F). The results of 321 histopathological analysis also showed that after ATV006 treatment, the inflammatory 322 cell infiltration in the liver was significantly reduced (Fig S6G). ATV006 could reduce 323 the production of various inflammatory cytokines (Fig S6H). The above results 324 demonstrate that ATV006 could also inhibit the replication of MHV-A59 in both the 325 lung and liver in the mouse model and prevent the death of MHV-infected mice, 326 indicating that ATV006 has a broad anti-coronavirus activity.

327 **Discussion** 

328 COVID-19, caused by SARS-CoV-2, is currently spreading globally, threatening 329 human health and economic development. The vaccine-induced or naturally acquired 330 protective herd immunity to interruption of transmission chains had been hampered by 331 the rapid evolution and recurrent emergence of SARS-CoV-2 variants, such as Delta 332 variant, one of the major variants of concern (VOC) (Liu et al. 2021; McCallum et al. 333 2021). Therefore, effective and broad-spectrum anti-SARS-CoV-2 drugs are 334 desperately needed. Until now, remdesivir is the only FDA-approved small molecule 335 antiviral drug for the treatment of COVID-19. However, the obligatory IV 336 administration of remdesivir limits its clinical application only for hospitalized patients 337 with advanced symptoms and its efficacy is limited in large scale clinical trials (Beigel 338 et al. 2020). As COVID-19 is an acute infectious disease, antiviral treatment can exert 339 its best effect at the early stage of the infection. In contrast, at the late stage of COVID-340 19 with hospitalized patients, anti-inflammatory therapy may play a major role in lessening the symptoms. Therefore, orally bioavailable anti-SARS-CoV-2 drugs 341 342 suitable for outpatients are superior to injectable drugs applied to hospitalized patients.

343 Previously, we reported 69-0, the major metabolite and the parent nucleotide of 344 remdesivir that targets the RdRp of SARS-CoV-2, has a better inhibitory activity on 345 SARS-CoV-2 and MHV-A59 in vitro and in vivo (Li et al. 2021). However, the 346 unfavorable oral PK prevents the further development of 69-0 (Li et al. 2021; NCATS 347 2021). To address this issue, herein, we synthesized a series of SCFA and amino acid 348 prodrug of 69-0 aiming at overcoming its limitations. Among the compounds 349 synthesized, the isobutyryl adenosine analogue ATV006 had improved oral absorption and potently inhibited the replication of SARS-CoV-2, especially the Delta variant (Fig 350 351 2). Compared to remdesivir, ATV006 is structurally simpler and easier to synthesize 352 via a three-step transformation with 69-0 as starting material, which would reduce the 353 cost and accelerate mass production. In addition, the orally active ATV006 is 354 potentially more useful for the management of SARS-CoV-2 infection at the early stage. After oral administration of ATV006, it is rapidly hydrolyzed by cellular esterases to produce the parent nucleoside 69-0 (Hsu et al. 2003; Lavis 2008), which then undergoes three steps of phosphorylation and is transformed to the active triphosphate form (Fig 3C), the same active component as that of remdesivir. Therefore, ATV006 shares the same mechanism of stalling SARS-CoV-2 polymerase as that of remdesivir (Kokic et al. 2021; Mackman et al. 2021; Wei et al. 2021; Yin et al. 2020a).

361 Several other small-molecule anti-SARS-CoV-2 antivirals are also under 362 development, including the ones that block viral entry, inhibit Mpro, and target host 363 immunity (Faheem et al. 2020; Good et al. 2021; Kabinger et al. 2021; Li et al. 2021; 364 Minghua et al. 2021; Pruijssers et al. 2020; Sabbah et al. 2021; Vuong et al. 2021; Wahl 365 et al. 2021; Zhang et al. 2021). Among them, EIDD-2801 was an orally available 366 prodrug of nucleoside analog at phase 3 clinical trials and it was granted provisional 367 approval in Australia in August, 2021. Clinical studies showed that a 5-day treatment 368 of EIDD-2801 has a 100% SARS-CoV-2 clearance rate in the non-hospitalized patient 369 (ClinicalTrials.gov NCT04405570) (Fischer et al. 2021). EIDD-2801 was reported to 370 induce a two-step mutagenesis of viral RNA that is different to the inhibitory 371 mechanism of ATV006 (Kabinger et al. 2021). We compared the antiviral efficiency of 372 ATV006 with that of EIDD-2801 in the same experimental settings, and the results 373 demonstrated that ATV006 possesses similar or higher anti-SARS-CoV-2 potency in 374 different mouse models (Fig. 4 and Fig. 5).

Delta variant recently causes the sharp rise in SARS-CoV-2 cases worldwide. Compared to the original strain, Alpha and Beta variant, the Delta variant is more infectious and pathogenic (Motozono et al. 2021; Reardon 2021; Teyssou et al. 2021). Intriguingly, ATV006 showed an improved antiviral potency against Delta variant and the EC<sub>50</sub> of ATV006 was about 3-4 times lower against Delta than the original strain and Beta variant (Fig 2). Oral treatments of ATV006 250 mg/kg, 100 mg/kg) as well as EIDD-2801 (500 mg/kg) effectively protected mice from severe weight loss and death 382 induced by the infection of Delta variant (Fig 5C), indicating the high efficacy of 383 ATV006 against the prevalent VOC. The signature mutations of the variants reside 384 mainly in the receptor binding domain of spike protein (McCallum et al. 2021; Salleh 385 et al. 2021) while the key residues of the nucleoside analog-binding and RdRp catalytic 386 sites are 100% conserved (Fig S7). Therefore, we speculate that the increased 387 sensitivity of Delta to ATV006 may be related to its high replication rate but not to the 388 mutations in RdRp. The detailed mechanism needs to be further investigated in the 389 future work. The analysis of 5,600 SARS-CoV-2 genomes also indicated that the 390 adenosine analog, the parent nucleoside of remdesivir, does not seem to exert high 391 selective pressure (Mari et al. 2021), suggesting that ATV006 and similar adenosine 392 analogs have low risk to induce escape mutation and drug resistance.

393 The RdRp-encoding nsp12 is the most conserved protein in the coronaviruses. The 394 key residues of nsp12 that are essential for RdRp enzymatic activity (Jin et al. 2021) 395 and the binding with the parent nucleoside of remdesivir (Yin et al. 2020b) are 100% 396 conserved throughout the coronaviruses of Alphacoronavirus and Betacoronavirus 397 (Fig). Indeed, our current study demonstrated that ATV006 has a broad antiviral 398 activity against different coronaviruses, including MHV, FIPV, CCoV, PEDV, TGEV 399 and SADS (Fig 2 and Fig 6) (Haake et al. 2020; Izes et al. 2020; Korner et al. 2020; 400 Laude et al. 1990; Lee 2015; Weiss et al. 2011; Zhou et al. 2018). The parent nucleoside 401 69-0 was previously reported to be broadly active against viruses that belong to the 402 families Paramyxoviridae, Coronaviridae and Filoviridae (Cho et al. 2012; Huang et 403 al. 2021; Lo et al. 2017; Murphy et al. 2018; Pedersen et al. 2019), indicating the 404 potential for more broad antiviral application of ATV006. Collectively, our results 405 demonstrated that ATV006 has potent and broad efficacy against SARS-CoV-2 and its 406 variants of concern as well as other coronaviruses, thus representing a promising orally 407 available drug candidate for the treatment for COVID-19 and emerging coronavirus 408 diseases in the future.

#### 409 Materials and Methods

#### 410 Compounds, cells, and viruses

- 411 The preparation of novel compounds, <sup>1</sup>H NMR, <sup>13</sup>C NMR, HRMS analysis and
- 412 HPLC purity are supplied in the Supplementary Information.
- 413 HEK 293T cells were obtained from American Tissue Culture Collection
- 414 (ATCC). Rat lung epithelial cells (L2) and wild-type MHV-A59 were kindly provided
- 415 by Rong Ye (Shanghai Medical School of Fudan University). African green monkey
- 416 kidney Vero E6 cell line (Vero E6) was kindly provided by Dr. Hui Zhang (Sun Yat-
- 417 sen University). Feline kidney cells (CRFK cells) and swine testicle cells (ST cells)
- 418 were provided from Guangdong Province Key Laboratory of Laboratory Animals.
- 419 HEK 293T, Vero E6, L2, CRFK and ST cells were cultured in DMEM supplemented
- 420 with 10% FBS, 100 U/mL penicillin and streptomycin at 37 °C in a humidified
- 421 atmosphere of 5% CO2.
- 422 SARS-CoV-2 (B.1, hCoV-19/CHN/SYSU-IHV/2020 strain, Accession ID on
- 423 GISAID: EPI\_ISL\_444969) was isolated from a sputum sample from a woman
- 424 admitted to the Eighth People's Hospital of Guangzhou.
- 425 SARS-CoV-2 (B.1.351, SARS CoV-2 human CHN 20SF18530 2020,
- 426 Accession ID on GWH: WHBDSE01000000) was isolated from a throat swab sample
  427 from a man admitted to the Eighth People's Hospital of Guangzhou.
- 428 SARS-CoV-2 (B.1.617.2, GDPCC 2.00096) was isolated from a patient infected
- 429 with SARS-CoV-2 Delta variant admitted in the Guangzhou Eighth People's Hospital
- 430 by Center for Disease Control and Prevention of Guangdong Province (Wang et al.
- 431 2021).
- 432 Canine coronavirus (CCoV), Porcine epidemic diarrhea virus (PEDV),
- 433 Transmissible gastroenteritis virus (TGEV), Swine acute diarrhea syndrome

- 435 Guangdong Province Key Laboratory of Laboratory Animals.
- 436 SARS-CoV-2 infection experiments were performed in the BSL-3 laboratory of
- 437 Sun Yat-sen University or Guangzhou Customs District Technology Center.
- 438 MHV-A59 infection experiments were performed in the Biosafety Level 2 (BSL
- 439 2) laboratory of Guangdong Laboratory Animals Monitoring Institute. All animal
- 440 studies protocols were approved by the Animal Welfare Committee and all
- 441 procedures used in animal studies complied with the guidelines and policies of the
- 442 Animal Care and Use Committee.
- 443 SARS-CoV-2 replicon assays
- 444 The assays were performed following the manufacturer's instructions (Promega
- 445 Corporation, Fitchburg, WI, USA). In brief, the cells in 24-well plate transfected with
- 446 500 ng pBAC- SARS-CoV-2-Replicon-Luciferase plasmid and 10 ng RL-TK
- 447 plasmid. After 6-8 h, the cells are transfected, the supernatant was discarded and
- 448 replaced with fresh DMEM medium, followed by adding each compound (described
- in Table 1) to the media with the final concentration of 50  $\mu$ M, 10  $\mu$ M, 5  $\mu$ M, 2  $\mu$ M, 1
- 450 μM, 0.1 μM or 0.01 μM. After 60 h, cells were lysed in 200 μL Passive Lysis Buffer
- 451 (PLB). Each lysate (20  $\mu$ L) was transferred into 96-well white plate and then mixed
- 452 with 20 µL Luciferase Assay Reagent II, followed by 20 µL of Stop & Glo solution.
- 453 The luminescence values of the two-step reaction were recorded using a luminescence
- 454 detector in Synergy H1 Hybrid Multi-Mode Reader.

#### 455 Anti-SARS-CoV-2 activity assays

- 456 Vero E6 and Huh7 cells were seeded at  $2 \times 10^4$  cells per well in 48-well plates.
- 457 Cells were allowed to adhere for 16-24 h and then infected at MOI of 0.05 with
- 458 SARS-CoV-2 for 1 h at 37°C. Then viral inoculum was removed, and cells were
- 459 washed 2 times with pre-warmed PBS. Medium containing dilutions of compounds,

460 or DMSO was added. At 48 hpi, supernatants or cells were harvested for qRT-PCR

461 analysis. The dose-response curves were plotted from viral RNA copies versus the

462 drug concentrations using GraphPad Prism 6 software.

463 Anti- MHV-A59 Activity Assays

464 L2 cells were seeded at  $1 \times 10^5$  cells per well in 6-well plates. Cells were allowed 465 to adhere for 16-24 h and then infected at MOI of 0.1 with MHV-A59 for 1h at 37°C. 466 Then viral inoculum was removed, and cells were washed 2 times with pre-warmed 467 PBS. Medium containing dilutions of compounds (ATV006, 69-0 and remdesivir), or 468 DMSO was added. At 20 hpi, supernatants were harvested for qRT-PCR analysis. The 469 EC<sub>50</sub> values were calculated from the dose response curve. Viruses and cells are listed 470 in Table S2. qPCR primers are listed in Table S3.

#### 471 Anti- FIPV, CCoV, PEDV TGEV and SADS Activity Assays

472 Cells were seeded in 48-cell plates for 24 h to reach 80% confluence and washed 473 thrice with serum-free medium. Cells were infected with virus (0.01 MOI) at 37°C for 474 1 h. Medium containing dilutions of ATV006 or DMSO was added. After incubating 475 at 37°C for 48h, cells and supernatants were harvested to determine viral loads using 476 qRT-PCR. The EC<sub>50</sub> values were calculated from the dose response curve. Viruses 477 and cells are listed in Table S2. qPCR primers are listed in Table S3.

#### 478 **qRT-PCR analysis**

479 For SARS-CoV-2 RNA quantification, RNA was isolated by Magbead Viral

480 DNA/RNA Kit (CWBIO). SARS-CoV-2 nucleic acid detection kit (Daan Company)
481 is used to detect the virus.

For the detection of cellular viruses and tissue viruses and cytokines, total RNA was isolated from cells or tissue samples with TRIzol reagent under the instruction of the manufacturer. The mRNAs were reverse transcribed into cDNA by PrimeScript 485RT reagent Kit (Takara). The cDNA was amplified by a fast two-step amplification486program using ChamQ Universal SYBR qPCR Master Mix (Vazyme Biotech Co.,487Ltd) or Taq Pro HS Universal Probe Master Mix (Vazyme Biotech Co., Ltd). GAPDH488was used to normalize the input samples via the  $\Delta$ Ct method. The relative mRNA489expression level of each gene was normalized to GAPDH housekeeping gene490expression in the untreated condition, and fold induction was calculated by the  $\Delta\Delta$ CT491method relative to those in untreated samples. The qRT-PCR primers are listed in

- 492 Table S3.
- 493 CCK-8 cell viability assay

To investigate the effect of drugs on cell viability, Vero E6 cells were seeded in
96-well plates at a density of 20,000 cells/well and were treated with drugs at
indicated concentrations (0, 0.01, 0.1, 1, 5, 10, 50 μM) for 48 h. Cell viability was
tested by using Cell Counting Kit-8 (CCK-8, Bimake, B34302). The figures were
plotted from viral RNA copies in supernatants versus the drug concentrations using
GraphPad Prism 6 software.

#### 500 **PK study in rat**

501 Male SD rats (180-220 g, N = 3) were fasted for 12 h before drug administration. ATV006 was administered intravenously at 4 mg/kg or intragastrically at 20 mg/kg. 502 503 Blood samples were collected from the jugular vein into anticoagulant EDTA-K2 504 tubes at 0.083, 0.25, 0.5, 1, 2, 3, 4, 6, 8 and 24 h for the IV group, and 0.25, 1, 0.5, 2, 505 3, 4, 6, 8 and 24 h for the IG group, respectively. All samples were centrifuged under 5064000 rpm/min for 10 min at 4 °C and the plasma (supernatants) were collected and 507 stored at -65 °C for future analysis. An aliquot of 50 µL each plasma sample was 508 treated with 250 µL of acetonitrile. The samples were centrifuged under 4000 509 rpm/min for 10 min and filtered through 0.2 µm membrane filters. The concentration 510 of analytes in each sample were analyzed by LC/MS/MS.

#### 511 **PK study in cynomolgus monkeys**

512 Three male cynomolgus monkeys (2 to 5 years of age, weighing 3 to 5 kg) were 513 used were orally received AVT006 of 10 mg/kg on day 1. After administration, the 514 blood samples for plasma were collected from a jugular vein into anticoagulant 515 EDTA-K<sub>2</sub> tubes at 0.083, 0.25, 0.5, 1, 2, 4, 8, 24 and 48 h. After a 3-day washout 516 period, each animal was administered intravenously with ATV006 at a dose of 5 517 mg/kg, followed by bold collection from the jugular vein at the specified time points. 518 All samples were centrifuged under 4000 rpm/min for 10 min at 4 °C and the plasma 519 (supernatants) were collected and stored at -65°C for future analysis. The 520 concentration of analytes in each sample were analyzed by LC/MS/MS. 521**Tissue distribution study in mice** 522 Five C57BL/6 mice were fasted for 12-16 h before administered orally with a 523 single dose of 100 mg/kg ATV006. After 1 h of dosing, 0.5 mL of the blood sample 524 was taken from heart. Liver, kidney, lung, brain tissues were harvested. The blood sample was processed to the same method as in the rat PK studies. Tissue samples 525 526were homogenized and extracted with 70% methanol, after centrifuge under 4000 rpm 527 for 10min, the supernatants were transfer to a clean tube and the concentration of

528 analytes in each sample were analyzed by LC/MS/MS.

#### 529 Ad5-hACE2 Mice Study

530 The experiments were performed as previously described (Sun et al. 2020a). Mice

531 were lightly anesthetized with isoflurane and transduced intranasally with  $2.5 \times 10^8$ 

532 FFU of Ad5-ACE2 in 75  $\mu$ L DMEM. Five days post transduction, mice were infected

533 intranasally with SARS-CoV-2 (1  $\times$  10<sup>5</sup> PFU) in a total volume of 50µL DMEM.

534 Focus forming assay (FFA)

535 Vero E6 cells were seeded in 96-well plates one day before infection. Tissue
536 homogenates were serially diluted and used to inoculate Vero E6 cells at 37°C for 1 h.
537 Inoculate were then removed before adding 125 µL 1.6% carboxymethylcellulose per

538 well and warmed to 37°C. After 24 h, cells were fixed with 4% paraformaldehyde and

- 539 permeabilized with 0.2% Triton X-100. Cells were then incubated with a rabbit anti-
- 540 SARS-CoV-2 nucleocapsid protein polyclonal antibody (Cat. No.: 40143-T62, Sino
- 541 Biological), followed by an HRP-labeled goat anti-rabbit secondary antibody (Cat.
- 542 No.: 109-035-088, Jackson Immuno Research Laboratories). The foci were visualized
- 543 by TrueBlue Peroxidase Substrate (KPL) and counted with an ELISPOT reader
- 544 (Cellular Technology). Viral titers were calculated as per gram tissue.

#### 545 MHV plaque assay

- 546 L2 cells were grown in 60-mm dishes to 70-80% confluence and infected with
- 547 1 mL of media containing viruses at dilutions ranging from 10-3 to 10-6. After 1 h at
- 548 37°C, the inoculate were removed. Seven ml of 0.95% agar (Amresco) in DMEM
- 549 with 5% FBS was overlaid onto cells at 1-2h. Plaques were picked between 24 and
- 550 36 h. For plaque staining, 3 mL of agar containing 0.02% neutral red (Sigma-Aldrich)
- 551 was overlaid onto cells. Six to eight hours later, the stained plaques were counted.

#### 552 AST/ALT assay

Blood was incubated at room temperature to allow coagulation and was then
centrifuged to obtain serum; the serum was used for measurements of alanine
transaminase (ALT) and aspartate aminotransferase (AST) levels using Assay Kit
(Nanjing Jiancheng Bioengineering Institute).

#### 557 H&E Staining.

558 Mice lung, liver and spleen dissections were fixed in zinc formalin and embedded 559 with paraffin. Tissue sections ( $\sim 4 \mu m$ ) were stained with hematoxylin and eosin.

#### 560 Analysis of RdRp mutations of SARS-CoV-2 and its variants

561There were four subtypes high-quality SARS-CoV-2 genomes sequences562available in Global Initiative on Sharing All Influenza Data (GISIAD) (Shu et al.

563 2017). These high-quality genomic sequences of SARS-CoV-2 were screened out

- under the following criteria: 1) complete genome (>29000bp); 2) high coverage (only
- 565 entries with <1% Ns and <0.05% unique amino acid mutations and no
- 566 insertion/deletion unless verified by submitter); 3) low coverage excl (exclude entries
- 567 with >5% NNNs). In total, we collected Alpha (B.1.1.7) (2021.5.1-6.10), Beta
- 568 (B.1.351) (2019-2021.6.10), Gamma (P.1) (2019-2021.6.10), Delta (B.1.617.2)
- 569 (2019-2021.6.10) were 108791, 13871, 18808, 29152 strains respectively.
- 570 Subsequently, we eliminated 50 Ns or ns sequences. After screening, the remaining
- 571 sequences were 92214, 6341, 13655, 21611 (Table S4).
- 572 SARS-CoV-2 genome of NC 045512.2 (Wu et al. 2020) was utilized as the
- 573 reference sequence. Multiple sequences alignments were performed using the
- 574 progressive method (FFT-NS-2) implemented in MAFFT (version 7.4) (Katoh et al.
- 575 2002). The whole genome mutation analysis was carried out used the pipeline
- 576 provided by CoVa (Ali. et al. 2021) (version 0.2) software. Finally, the ggplot2 was

577 used for drawing in R.

#### 578 Coronavirus RdRp Conservation Analysis

Virus data collection is derived from International Committee on Taxonomy of
Viruses (ICTV) and NCBI, multi-sequence comparisons using mega 6, and data
visualization using texshade (Beitz 2000) (Table S4). The sequence of 523aa-734aa of
RdRp is selected for display, where K545, R555, S682, N691 and D760 are the key
sites for remdesivir binding to RdRp, and 759-761 (SDD) is the key site enzyme
activity of RdRp.

585 Statistical analysis.

586 All values are mean  $\pm$  SD or SEM of individual samples. Data analysis was

- 587 performed with GraphPad Prism Software (GraphPad Software Inc., version 6.01).
- 588 The statistical tests utilized are two-tailed and respective details have been indicated

589	in figure legends.	p-value of $< 0.05$ we	re considered statisticall	y significant. (*, r	)-
					·

590 value of  $\leq 0.05$ . \*\*, p-value of  $\leq 0.005$ . \*\*\*, p-value of  $\leq 0.0005$ . \*\*\*\*, p-value of  $\leq$ 

591 **0.0001**).

592

#### 593 ACKNOWLEDGEMENTS

- 594 The project was supported by Shenzhen Science and Technology Program
- 595 (JSGG20200225150431472, ZDSYS20190902093215877 &
- 596 KQTD20180411143323605), Shenzhen Bay Laboratory (SZBL2019062801006),
- 597 Guangdong Basic and Applied Basic Research Foundation (Grant
- 598 #2020A1515110361) and National Natural Science Foundation of China (grant
- 599 #32041002 & #81620108020). D.G. is also supported by Guangdong Zhujiang
- 600 Talents Program and National Ten-thousand Talents Program. We thank Dr. Chuwen
- 601 Lin from School of Medicine, Sun Yat-Sen University for the help in lung pathology
- analysis. We thank the Center for Disease Control and Prevention of Guangdong
- 603 Province for providing the Delta variant of SARS-CoV-2.
- 604
- 605

#### 606 **REFERENCES**

- 607
- Al-Tawfiq, J. A., and Z. A. Memish. 2014. 'Middle East respiratory
  syndrome coronavirus: transmission and phylogenetic evolution', *Trends Microbiol*, 22: 573-9.
- Alanazi, A. S., E. James, and Y. Mehellou. 2019. 'The ProTide Prodrug
  Technology: Where Next?', ACS Med Chem Lett, 10: 2-5.
- Ali., Farhan, Mohak Sharda., and Aswin Sai Narain Seshasayee. 2021.
  'SARS-CoV-2 sequence typing, evolution and signatures of
  selection using CoVa, a Python-based command-line utility', *biovix*.
- Beigel, J. H., K. M. Tomashek, L. E. Dodd, A. K. Mehta, B. S.
  Zingman, A. C. Kalil, E. Hohmann, H. Y. Chu, A. Luetkemeyer, S.
  Kline, D. Lopez de Castilla, R. W. Finberg, K. Dierberg, V.
  Tapson, L. Hsieh, T. F. Patterson, R. Paredes, D. A. Sweeney,
  W. R. Short, G. Touloumi, D. C. Lye, N. Ohmagari, M. D. Oh, G.
  M. Ruiz-Palacios, T. Benfield, G. Fatkenheuer, M. G.
  Kortepeter, R. L. Atmar, C. B. Creech, J. Lundgren, A. G.

624 625 626 627	Babiker, S. Pett, J. D. Neaton, T. H. Burgess, T. Bonnett, M. Green, M. Makowski, A. Osinusi, S. Nayak, H. C. Lane, and Actt-Study Group Members. 2020. 'Remdesivir for the Treatment of Covid-19 - Final Report', <i>N Engl J Med</i> , 383: 1813-26.
628	Beitz, E. 2000. 'TEXshade: shading and labeling of multiple sequence
629	alignments using LATEX2 epsilon', <i>Bioinformatics</i> , 16: 135-9.
630	Chen, Y., Q. Liu, and D. Guo. 2020. 'Emerging coronaviruses: Genome
631	structure, replication, and pathogenesis', <i>J Med Virol</i> , 92:
632	2249.
633	Cho, A., O. L. Saunders, T. Butler, L. Zhang, J. Xu, J. E. Vela, J.
634	Y. Feng, A. S. Ray, and C. U. Kim. 2012. 'Synthesis and
635	antiviral activity of a series of 1'-substituted 4-aza-7,9-
636	dideazaadenosine C-nucleosides', <i>Bioorg Med Chem Lett</i> , 22:
637	2705-7.
638	Consortium, W. H. O. Solidarity Trial, H. Pan, R. Peto, A. M. Henao-
639	Restrepo, M. P. Preziosi, V. Sathiyamoorthy, Q. Abdool Karim,
640	M. M. Alejandria, C. Hernandez Garcia, M. P. Kieny, R.
641	Malekzadeh, S. Murthy, K. S. Reddy, M. Roses Periago, P. Abi
642	Hanna, F. Ader, A. M. Al-Bader, A. Alhasawi, E. Allum, A.
643	Alotaibi, C. A. Alvarez-Moreno, S. Appadoo, A. Asiri, P.
644	Aukrust, A. Barratt-Due, S. Bellani, M. Branca, H. B. C.
645	Cappel-Porter, N. Cerrato, T. S. Chow, N. Como, J. Eustace, P.
646	J. Garcia, S. Godbole, E. Gotuzzo, L. Griskevicius, R. Hamra,
647	M. Hassan, M. Hassany, D. Hutton, I. Irmansyah, L. Jancoriene,
648	J. Kirwan, S. Kumar, P. Lennon, G. Lopardo, P. Lydon, N.
649	Magrini, T. Maguire, S. Manevska, O. Manuel, S. McGinty, M. T.
650	Medina, M. L. Mesa Rubio, M. C. Miranda-Montoya, J. Nel, E. P.
651	Nunes, M. Perola, A. Portoles, M. R. Rasmin, A. Raza, H. Rees,
652	P. P. S. Reges, C. A. Rogers, K. Salami, M. I. Salvadori, N.
653	Sinani, J. A. C. Sterne, M. Stevanovikj, E. Tacconelli, K. A.
654	O. Tikkinen, S. Trelle, H. Zaid, J. A. Rottingen, and S.
655	Swaminathan. 2021. 'Repurposed Antiviral Drugs for Covid-19 -
656	Interim WHO Solidarity Trial Results', <i>N Engl J Med</i> , 384: 497-
657	511.
658	Faheem, B. K. Kumar, Kvgc Sekhar, S. Kunjiappan, J. Jamalis, R.
659	Balaña-Fouce, B. L. Tekwani, and M. Sankaranarayanan. 2020.
660	'Druggable targets of SARS-CoV-2 and treatment opportunities
661	for COVID-19', <i>Bioorg Chem</i> , 104: 104269.
662	Fischer, W., J. J. Eron, W. Holman, M. S. Cohen, L. Fang, L. J.
663	Szewczyk, T. P. Sheahan, R. Baric, K. R. Mollan, C. R. Wolfe,
664 665	E. R. Duke, M. M. Azizad, K. Borroto-Esoda, D. A. Wohl, A. J.
665	Loftis, P. Alabanza, F. Lipansky, and W. P. Painter. 2021.

666 'Molnupiravir, an Oral Antiviral Treatment for COVID-19', 667 medRxiv. 668 T. S., and D. X. Liu. 2019. 'Human Coronavirus: Host-Pathogen Fung, 669 Interaction', Annu Rev Microbiol, 73: 529-57. 670 Ghasemnejad-Berenji, M., and S. Pashapour. 2021. 'Favipiravir and COVID-19: A Simplified Summary', Drug Res (Stuttg), 71: 166-70. 671 Goldman, J. D., D. C. B. Lye, D. S. Hui, K. M. Marks, R. Bruno, R. 672 673 Montejano, C. D. Spinner, M. Galli, M. Y. Ahn, R. G. Nahass, Y. 674 S. Chen, D. SenGupta, R. H. Hyland, A. O. Osinusi, H. Cao, C. 675 Blair, X. Wei, A. Gaggar, D. M. Brainard, W. J. Towner, J. 676 Munoz, K. M. Mullane, F. M. Marty, K. T. Tashima, G. Diaz, A. Subramanian, and Gs-Us- Investigators. 2020. 'Remdesivir for 5 677 or 10 Days in Patients with Severe Covid-19', N Engl J Med, 678 679 383: 1827-37. Good, S. S., J. Westover, K. H. Jung, X. J. Zhou, A. Moussa, P. La 680 Colla, G. Collu, B. Canard, and J. P. Sommadossi. 2021. 'AT-681 527, a Double Prodrug of a Guanosine Nucleotide Analog, Is a 682 Potent Inhibitor of SARS-CoV-2 In Vitro and a Promising Oral 683 684 Antiviral for Treatment of COVID-19', Antimicrob Agents Chemother, 65. 685 686 Haake, C., S. Cook, N. Pusterla, and B. Murphy. 2020. 'Coronavirus 687 Infections in Companion Animals: Virology, Epidemiology, 688 Clinical and Pathologic Features', Viruses, 12. 689 Harvey, W. T., A. M. Carabelli, B. Jackson, R. K. Gupta, E. C. 690 Thomson, E. M. Harrison, C. Ludden, R. Reeve, A. Rambaut, 691 Covid- Genomics UK Consortium, S. J. Peacock, and D. L. 692 Robertson. 2021. 'SARS-CoV-2 variants, spike mutations and immune escape', Nat Rev Microbiol, 19: 409-24. 693 Hillen, H. S., G. Kokic, L. Farnung, C. Dienemann, D. Tegunov, and P. 694 695 Cramer. 2020. 'Structure of replicating SARS-CoV-2 polymerase', 696 Nature, 584: 154-56. Hsu, C. H., M. Jay, P. M. Bummer, and H. J. Lehmler. 2003. 'Chemical 697 stability of esters of nicotinic acid intended for pulmonary 698 699 administration by liquid ventilation', *Pharm Res*, 20: 918-25. 700 Huang, Z., L. Gong, Z. Zheng, Q. Gao, X. Chen, Y. Chen, X. Chen, R. 701 Xu, J. Zheng, Z. Xu, S. Zhang, H. Wang, and G. Zhang. 2021. 702 'GS-441524 inhibits African swine fever virus infection in vitro', Antiviral Res, 191: 105081. 703 Hussain, S., J. Pan, Y. Chen, Y. Yang, J. Xu, Y. Peng, Y. Wu, Z. Li, 704 705 Y. Zhu, P. Tien, and D. Guo. 2005. 'Identification of novel 706 subgenomic RNAs and noncanonical transcription initiation

707 signals of severe acute respiratory syndrome coronavirus', J708 Virol, 79: 5288-95. 709 Izes, A. M., J. Yu, J. M. Norris, and M. Govendir. 2020. 'Current 710 status on treatment options for feline infectious peritonitis 711 and SARS-CoV-2 positive cats', Vet Q, 40: 322-30. Jin, Y., H. Lin, L. Cao, W. C. Wu, Y. Ji, L. Du, Y. Jiang, Y. Xie, 712 K. Tong, F. Xing, F. Zheng, M. Shi, J. A. Pan, X. Peng, and D. 713 714 Guo. 2021. 'A Convenient and Biosafe Replicon with Accessory Genes of SARS-CoV-2 and Its Potential Application in Antiviral 715 716 Drug Discovery', Virol Sin. Kabinger, F., C. Stiller, J. Schmitzova, C. Dienemann, G. Kokic, H. 717 S. Hillen, C. Hobartner, and P. Cramer. 2021. 'Mechanism of 718 molnupiravir-induced SARS-CoV-2 mutagenesis', Nat Struct Mol 719 720 Biol. Katoh, K., K. Misawa, K. Kuma, and T. Miyata. 2002. 'MAFFT: a novel 721 722 method for rapid multiple sequence alignment based on fast Fourier transform', Nucleic Acids Res, 30: 3059-66. 723 Kim, D., J. Y. Lee, J. S. Yang, J. W. Kim, V. N. Kim, and H. Chang. 724 725 2020. 'The Architecture of SARS-CoV-2 Transcriptome', Cell, 726 181: 914-21 e10. 727 Kokic, G., H. S. Hillen, D. Tegunov, C. Dienemann, F. Seitz, J. 728 Schmitzova, L. Farnung, A. Siewert, C. Hobartner, and P. 729 Cramer. 2021. 'Mechanism of SARS-CoV-2 polymerase stalling by 730 remdesivir', Nat Commun, 12: 279. 731 Korner, R. W., M. Majjouti, M. A. A. Alcazar, and E. Mahabir. 2020. 732 'Of Mice and Men: The Coronavirus MHV and Mouse Models as a 733 Translational Approach to Understand SARS-CoV-2', Viruses, 12. 734 Laude, H., D. Rasschaert, B. Delmas, M. Godet, J. Gelfi, and B. Charley. 1990. 'Molecular biology of transmissible 735 gastroenteritis virus', Vet Microbiol, 23: 147-54. 736 737 Lavis, L. D. 2008. 'Ester bonds in prodrugs', ACS Chem Biol, 3: 203-738 6. Lee, C. 2015. 'Porcine epidemic diarrhea virus: An emerging and re-739 740 emerging epizootic swine virus', Virol J, 12: 193. 741 Li, Y., L. Cao, G. Li, F. Cong, Y. Li, J. Sun, Y. Luo, G. Chen, G. 742 Li, P. Wang, F. Xing, Y. Ji, J. Zhao, Y. Zhang, D. Guo, and X. 743 Zhang. 2021. 'Remdesivir Metabolite GS-441524 Effectively 744 Inhibits SARS-CoV-2 Infection in Mouse Models', J Med Chem. Liu, C., H. M. Ginn, W. Dejnirattisai, P. Supasa, B. Wang, A. 745 Tuekprakhon, R. Nutalai, D. Zhou, A. J. Mentzer, Y. Zhao, H. M. 746 747 E. Duyvesteyn, C. Lopez-Camacho, J. Slon-Campos, T. S. Walter, 748 D. Skelly, S. A. Johnson, T. G. Ritter, C. Mason, S. A. Costa

749	Clamona E Camaa Navaaa V Nagaimenta E Nagaimenta C
749 750	Clemens, F. Gomes Naveca, V. Nascimento, F. Nascimento, C.
	Fernandes da Costa, P. C. Resende, A. Pauvolid-Correa, M. M.
751 759	Siqueira, C. Dold, N. Temperton, T. Dong, A. J. Pollard, J. C.
752 752	Knight, D. Crook, T. Lambe, E. Clutterbuck, S. Bibi, A.
753	Flaxman, M. Bittaye, S. Belij-Rammerstorfer, S. C. Gilbert, T.
754	Malik, M. W. Carroll, P. Klenerman, E. Barnes, S. J. Dunachie,
755	V. Baillie, N. Serafin, Z. Ditse, K. Da Silva, N. G. Paterson,
756	M. A. Williams, D. R. Hall, S. Madhi, M. C. Nunes, P. Goulder,
757	E. E. Fry, J. Mongkolsapaya, J. Ren, D. I. Stuart, and G. R.
758	Screaton. 2021. 'Reduced neutralization of SARS-CoV-2 B.1.617
759	by vaccine and convalescent serum', <i>Cell</i> , 184: 4220-36 e13.
760	Lo, M. K., R. Jordan, A. Arvey, J. Sudhamsu, P. Shrivastava-Ranjan,
761	A. L. Hotard, M. Flint, L. K. McMullan, D. Siegel, M. O.
762	Clarke, R. L. Mackman, H. C. Hui, M. Perron, A. S. Ray, T.
763	Cihlar, S. T. Nichol, and C. F. Spiropoulou. 2017. 'GS-5734 and
764	its parent nucleoside analog inhibit Filo-, Pneumo-, and
765	Paramyxoviruses', <i>Sci Rep</i> , 7: 43395.
766	Mackman, R. L., H. C. Hui, M. Perron, E. Murakami, C. Palmiotti, G.
767	Lee, K. Stray, L. Zhang, B. Goyal, K. Chun, D. Byun, D. Siegel,
768	S. Simonovich, V. Du Pont, J. Pitts, D. Babusis, A.
769	Vijjapurapu, X. Lu, C. Kim, X. Zhao, J. Chan, B. Ma, D. Lye, A.
770	Vandersteen, S. Wortman, K. T. Barrett, M. Toteva, R. Jordan,
771	R. Subramanian, J. P. Bilello, and T. Cihlar. 2021. 'Prodrugs
772	of a 1'-CN-4-Aza-7,9-dideazaadenosine C-Nucleoside Leading to
773	the Discovery of Remdesivir (GS-5734) as a Potent Inhibitor of
774	Respiratory Syncytial Virus with Efficacy in the African Green
775	Monkey Model of RSV', <i>J Med Chem</i> , 64: 5001-17.
776	Mari, A., T. Roloff, M. Stange, K. K. Sogaard, E. Asllanaj, G.
777	Tauriello, L. T. Alexander, M. Schweitzer, K. Leuzinger, A.
778	Gensch, A. E. Martinez, J. Bielicki, H. Pargger, M. Siegemund,
779	C. H. Nickel, R. Bingisser, M. Osthoff, S. Bassetti, P. Sendi,
780	M. Battegay, C. Marzolini, H. M. B. Seth-Smith, T. Schwede, H.
781	H. Hirsch, and A. Egli. 2021. 'Global Genomic Analysis of SARS-
782	CoV-2 RNA Dependent RNA Polymerase Evolution and Antiviral Drug
783	Resistance', <i>Microorganisms</i> , 9.
784	Martin, Y. C. 2005. 'A bioavailability score', <i>J Med Chem</i> , 48: 3164-
785	70.
786	McCallum, M., J. Bassi, A. Marco, A. Chen, A. C. Walls, J. D. Iulio,
780 787	M. A. Tortorici, M. J. Navarro, C. Silacci-Fregni, C. Saliba,
788	M. Agostini, D. Pinto, K. Culap, S. Bianchi, S. Jaconi, E.
789 700	Cameroni, J. E. Bowen, S. W. Tilles, M. S. Pizzuto, S. B.
790	Guastalla, G. Bona, A. F. Pellanda, C. Garzoni, W. C. Van

791 792	Voorhis, L. E. Rosen, G. Snell, A. Telenti, H. W. Virgin, L. Piccoli, D. Corti, and D. Veesler. 2021. 'SARS-CoV-2 immune
793	evasion by variant B.1.427/B.1.429', <i>bioRxiv</i> .
794	Minghua, Li., Ferretti. Max, Ying. Baoling, Descamps. Hélène, Lee.
795	Emily, Dittmar. Mark, Lee. Jae Seung, Whig. Kanupriya, Brinda
796	Kamalia., Dohnalová. Lenka, Uhr. Giulia, Zarkoob. Hoda, Chen.
797	Yu-Chi, Ramage. Holly, Ferrer. Marc, Lynch. Kristen, Schultz.
798	David C., Christoph A. Thaiss., Diamond. Michael S., and
799	Cherry. Sara. 2021. 'Pharmacological activation of STING blocks
800	SARS-CoV-2 infection', <i>Sci Immunol</i> , 6.
801	Motozono, C., M. Toyoda, J. Zahradnik, A. Saito, H. Nasser, T. S.
802	Tan, I. Ngare, I. Kimura, K. Uriu, Y. Kosugi, Y. Yue, R.
803	Shimizu, J. Ito, S. Torii, A. Yonekawa, N. Shimono, Y.
804	Nagasaki, R. Minami, T. Toya, N. Sekiya, T. Fukuhara, Y.
805	Matsuura, G. Schreiber, Consortium Genotype to Phenotype Japan,
806	T. Ikeda, S. Nakagawa, T. Ueno, and K. Sato. 2021. 'SARS-CoV-2
807	spike L452R variant evades cellular immunity and increases
808	infectivity', Cell Host Microbe, 29: 1124-36 ell.
809	Murphy, B. G., M. Perron, E. Murakami, K. Bauer, Y. Park, C.
810	Eckstrand, M. Liepnieks, and N. C. Pedersen. 2018. 'The
811	nucleoside analog GS-441524 strongly inhibits feline infectious
812	peritonitis (FIP) virus in tissue culture and experimental cat
813	infection studies', Vet Microbiol, 219: 226-33.
814	NCATS. 2021. 'GS-441524 Studies', National Center for Advancing
815	Translational Sciences.
816	https://opendata.ncats.nih.gov/covid19/GS-441524.
817	Oladunni, F. S., J. G. Park, P. A. Pino, O. Gonzalez, A. Akhter, A.
818	Allue-Guardia, A. Olmo-Fontanez, S. Gautam, A. Garcia-Vilanova,
819	C. Ye, K. Chiem, C. Headley, V. Dwivedi, L. M. Parodi, K. J.
820	Alfson, H. M. Staples, A. Schami, J. I. Garcia, A. Whigham, R.
821	N. Platt, 2nd, M. Gazi, J. Martinez, C. Chuba, S. Earley, O. H.
822	Rodriguez, S. D. Mdaki, K. N. Kavelish, R. Escalona, C. R. A.
823	Hallam, C. Christie, J. L. Patterson, T. J. C. Anderson, R.
824	Carrion, Jr., E. J. Dick, Jr., S. Hall-Ursone, L. S.
825	Schlesinger, X. Alvarez, D. Kaushal, L. D. Giavedoni, J.
826	Turner, L. Martinez-Sobrido, and J. B. Torrelles. 2020.
827	'Lethality of SARS-CoV-2 infection in K18 human angiotensin-
828	converting enzyme 2 transgenic mice', <i>Nat Commun</i> , 11: 6122.
829	Pedersen, N. C., M. Perron, M. Bannasch, E. Montgomery, E. Murakami,
830	M. Liepnieks, and H. Liu. 2019. 'Efficacy and safety of the
831	nucleoside analog GS-441524 for treatment of cats with

naturally occurring feline infectious peritonitis', J Feline 832 Med Surg, 21: 271-81. 833 834 Perlman, S., and J. Netland. 2009. 'Coronaviruses post-SARS: update 835 on replication and pathogenesis', Nat Rev Microbiol, 7: 439-50. 836 Picarazzi, F., I. Vicenti, F. Saladini, M. Zazzi, and M. Mori. 2020. 'Targeting the RdRp of Emerging RNA Viruses: The Structure-837 Based Drug Design Challenge', Molecules, 25. 838 839 Pruijssers, A. J., A. S. George, A. Schafer, S. R. Leist, L. E. 840 Gralinksi, K. H. Dinnon, 3rd, B. L. Yount, M. L. Agostini, L. J. Stevens, J. D. Chappell, X. Lu, T. M. Hughes, K. Gully, D. 841 842 R. Martinez, A. J. Brown, R. L. Graham, J. K. Perry, V. Du 843 Pont, J. Pitts, B. Ma, D. Babusis, E. Murakami, J. Y. Feng, J. P. Bilello, D. P. Porter, T. Cihlar, R. S. Baric, M. R. 844 845 Denison, and T. P. Sheahan. 2020. 'Remdesivir Inhibits SARS-CoV-2 in Human Lung Cells and Chimeric SARS-CoV Expressing the 846 SARS-CoV-2 RNA Polymerase in Mice', Cell Rep, 32: 107940. 847 Reardon, S. 2021. 'How the Delta variant achieves its ultrafast 848 spread', Nature. 849 850 Robson, F., K. S. Khan, T. K. Le, C. Paris, S. Demirbag, P. Barfuss, P. Rocchi, and W. L. Ng. 2020. 'Coronavirus RNA Proofreading: 851 Molecular Basis and Therapeutic Targeting', Mol Cell, 79: 710-852 853 27. Sabbah, D. A., R. Hajjo, S. K. Bardaweel, and H. A. Zhong. 2021. 'An 854 Updated Review on SARS-CoV-2 Main Proteinase (M(Pro)): Protein 855 Structure and Small-Molecule Inhibitors', Curr Top Med Chem, 856 21: 442-60. 857 Salleh, M. Z., J. P. Derrick, and Z. Z. Deris. 2021. 'Structural 858 859 Evaluation of the Spike Glycoprotein Variants on SARS-CoV-2 Transmission and Immune Evasion', Int J Mol Sci, 22. 860 Shu, Y., and J. McCauley. 2017. 'GISAID: Global initiative on sharing 861 862 all influenza data - from vision to reality', Euro Surveill, 863 22. 864 Sun, J., Z. Zhuang, J. Zheng, K. Li, R. L. Wong, D. Liu, J. Huang, J. He, A. Zhu, J. Zhao, X. Li, Y. Xi, R. Chen, A. N. Alshukairi, 865 Z. Chen, Z. Zhang, C. Chen, X. Huang, F. Li, X. Lai, D. Chen, 866 L. Wen, J. Zhuo, Y. Zhang, Y. Wang, S. Huang, J. Dai, Y. Shi, 867 K. Zheng, M. R. Leidinger, J. Chen, Y. Li, N. Zhong, D. K. 868 Meyerholz, P. B. McCray, Jr., S. Perlman, and J. Zhao. 2020a. 869 'Generation of a Broadly Useful Model for COVID-19 870 Pathogenesis, Vaccination, and Treatment', Cell, 182: 734-43 871 872 e5.

873 Sun, S. H., Q. Chen, H. J. Gu, G. Yang, Y. X. Wang, X. Y. Huang, S. 874 S. Liu, N. N. Zhang, X. F. Li, R. Xiong, Y. Guo, Y. Q. Deng, W. 875 J. Huang, Q. Liu, Q. M. Liu, Y. L. Shen, Y. Zhou, X. Yang, T. Y. Zhao, C. F. Fan, Y. S. Zhou, C. F. Qin, and Y. C. Wang. 876 877 2020b. 'A Mouse Model of SARS-CoV-2 Infection and Pathogenesis', Cell Host Microbe, 28: 124-33 e4. 878 Tegally, H., E. Wilkinson, M. Giovanetti, A. Iranzadeh, V. Fonseca, 879 880 J. Giandhari, D. Doolabh, S. Pillay, E. J. San, N. Msomi, K. 881 Mlisana, A. von Gottberg, S. Walaza, M. Allam, A. Ismail, T. 882 Mohale, A. J. Glass, S. Engelbrecht, G. Van Zyl, W. Preiser, F. 883 Petruccione, A. Sigal, D. Hardie, G. Marais, N. Y. Hsiao, S. Korsman, M. A. Davies, L. Tyers, I. Mudau, D. York, C. Maslo, 884 885 D. Goedhals, S. Abrahams, O. Laguda-Akingba, A. Alisoltani-886 Dehkordi, A. Godzik, C. K. Wibmer, B. T. Sewell, J. Lourenco, L. C. J. Alcantara, S. L. Kosakovsky Pond, S. Weaver, D. 887 Martin, R. J. Lessells, J. N. Bhiman, C. Williamson, and T. de 888 Oliveira. 2021. 'Detection of a SARS-CoV-2 variant of concern 889 in South Africa', Nature, 592: 438-43. 890 891 Teyssou, E., H. Delagreverie, B. Visseaux, S. Lambert-Niclot, S. Brichler, V. Ferre, S. Marot, A. Jary, E. Todesco, A. 892 893 Schnuriger, E. Ghidaoui, B. Abdi, S. Akhavan, N. Houhou-Fidouh, 894 C. Charpentier, L. Morand-Joubert, D. Boutolleau, D. Descamps, V. Calvez, A. G. Marcelin, and C. Soulie. 2021. 'The Delta 895 SARS-CoV-2 variant has a higher viral load than the Beta and 896 897 the historical variants in nasopharyngeal samples from newly 898 diagnosed COVID-19 patients', J Infect. Vuong, W., C. Fischer, M. B. Khan, M. J. van Belkum, T. Lamer, K. D. 899 900 Willoughby, J. Lu, E. Arutyunova, M. A. Joyce, H. A. Saffran, J. A. Shields, H. S. Young, J. A. Nieman, D. L. Tyrrell, M. J. 901 Lemieux, and J. C. Vederas. 2021. 'Improved SARS-CoV-2 M(pro) 902 903 inhibitors based on feline antiviral drug GC376: Structural 904 enhancements, increased solubility, and micellar studies', Eur 905 J Med Chem, 222: 113584. 906 Wahl, A., L. E. Gralinski, C. E. Johnson, W. Yao, M. Kovarova, K. H. 907 Dinnon, 3rd, H. Liu, V. J. Madden, H. M. Krzystek, C. De, K. K. White, K. Gully, A. Schafer, T. Zaman, S. R. Leist, P. O. 908 Grant, G. R. Bluemling, A. A. Kolykhalov, M. G. Natchus, F. B. 909 Askin, G. Painter, E. P. Browne, C. D. Jones, R. J. Pickles, R. 910 S. Baric, and J. V. Garcia. 2021. 'SARS-CoV-2 infection is 911 effectively treated and prevented by EIDD-2801', Nature, 591: 912 451-57. 913

914	Wang, M., R. Cao, L. Zhang, X. Yang, J. Liu, M. Xu, Z. Shi, Z. Hu, W.
915	Zhong, and G. Xiao. 2020a. 'Remdesivir and chloroquine
916	effectively inhibit the recently emerged novel coronavirus
917	(2019-nCoV) in vitro', <i>Cell Res</i> , 30: 269-71.
918	Wang, Q., J. Wu, H. Wang, Y. Gao, Q. Liu, A. Mu, W. Ji, L. Yan, Y.
919	Zhu, C. Zhu, X. Fang, X. Yang, Y. Huang, H. Gao, F. Liu, J. Ge,
920	Q. Sun, X. Yang, W. Xu, Z. Liu, H. Yang, Z. Lou, B. Jiang, L.
921	W. Guddat, P. Gong, and Z. Rao. 2020b. 'Structural Basis for
922	RNA Replication by the SARS-CoV-2 Polymerase', <i>Cell</i> , 182: 417-
923	28  e13.
924	Wang, Y., R. Chen, F. Hu, Y. Lan, Z. Yang, C. Zhan, J. Shi, X. Deng,
925	M. Jiang, S. Zhong, B. Liao, K. Deng, J. Tang, L. Guo, M.
926	Jiang, Q. Fan, M. Li, J. Liu, Y. Shi, X. Deng, X. Xiao, M.
927	Kang, Y. Li, W. Guan, Y. Li, S. Li, F. Li, N. Zhong, and X.
928	Tang. 2021. 'Transmission, viral kinetics and clinical
929	characteristics of the emergent SARS-CoV-2 Delta VOC in
930	Guangzhou, China', <i>EClinicalMedicine</i> , 40: 101129.
931	Wei, D., T. Hu, Y. Zhang, W. Zheng, H. Xue, J. Shen, Y. Xie, and H.
932	A. Aisa. 2021. 'Potency and pharmacokinetics of GS-441524
933	derivatives against SARS-CoV-2', <i>Bioorg Med Chem</i> , 46: 116364.
934 934	Weiss, S. R., and J. L. Leibowitz. 2011. 'Coronavirus pathogenesis',
935	Adv Virus Res, 81: 85–164.
936 936	WHO. 2021a. 'SARS-CoV-2 Variants, Working Definitions and Actions
937	Taken'. https://www.who.int/en/activities/tracking-SARS-CoV-2-
938	variants.
939 939	———. 2021b. 'WHO Coronavirus (COVID-19) Dashboard'.
939 940	https://covid19.who.int.
940 941	Wu, F., S. Zhao, B. Yu, Y. M. Chen, W. Wang, Z. G. Song, Y. Hu, Z. W.
942	Tao, J. H. Tian, Y. Y. Pei, M. L. Yuan, Y. L. Zhang, F. H. Dai,
943	Y. Liu, Q. M. Wang, J. J. Zheng, L. Xu, E. C. Holmes, and Y. Z.
944 944	Zhang. 2020. 'A new coronavirus associated with human
945	respiratory disease in China', <i>Nature</i> , 579: 265-69.
946	Yin, W., X. Luan, Z. Li, Y. Xie, Z. Zhou, J. Liu, M. Gao, X. Wang, F.
947	Zhou, Q. Wang, Q. Wang, D. Shen, Y. Zhang, G. Tian, Haji A.
948	
	Aisa, T. Hu, D. Wei, Y. Jiang, G. Xiao, H. Jiang, L. Zhang, X. Vu. I. Shop, S. Zhang, and H. F. Vu. 2020a, 'Structural basis
949 950	Yu, J. Shen, S. Zhang, and H. E. Xu. 2020a. 'Structural basis for repurpose and design of nucleoside drugs for treating
	for repurpose and design of nucleoside drugs for treating COVID-19'.
951 952	
952 953	Yin, W., C. Mao, X. Luan, D. D. Shen, Q. Shen, H. Su, X. Wang, F.
	Zhou, W. Zhao, M. Gao, S. Chang, Y. C. Xie, G. Tian, H. W. Liong S. C. Too, J. Shop, V. Liong, H. Liong, V. Yu, S. Zhong
954 055	Jiang, S. C. Tao, J. Shen, Y. Jiang, H. Jiang, Y. Xu, S. Zhang, V. Zhang, and H. F. Xu. 2020b, 'Structural basis for inhibition
955	Y. Zhang, and H. E. Xu. 2020b. 'Structural basis for inhibition

956	of the RNA-dependent RNA polymerase from SARS-CoV-2 by
957	remdesivir', <i>Science</i> , 368: 1499-504.
958	Zhang, Q., R. Xiang, S. Huo, Y. Zhou, S. Jiang, Q. Wang, and F. Yu.
959	2021. 'Molecular mechanism of interaction between SARS-CoV-2
960	and host cells and interventional therapy', Signal Transduct
961	Target Ther, 6: 233.
962	Zhang, Y., J. Sun, Y. Sun, Y. Wang, and Z. He. 2013. 'Prodrug design
963	targeting intestinal PepT1 for improved oral absorption: design
964	and performance', Curr Drug Metab, 14: 675-87.
965	Zhou, P., H. Fan, T. Lan, X. L. Yang, W. F. Shi, W. Zhang, Y. Zhu, Y.
966	W. Zhang, Q. M. Xie, S. Mani, X. S. Zheng, B. Li, J. M. Li, H.
967	Guo, G. Q. Pei, X. P. An, J. W. Chen, L. Zhou, K. J. Mai, Z. X.
968	Wu, D. Li, D. E. Anderson, L. B. Zhang, S. Y. Li, Z. Q. Mi, T.
969	T. He, F. Cong, P. J. Guo, R. Huang, Y. Luo, X. L. Liu, J.
970	Chen, Y. Huang, Q. Sun, X. L. Zhang, Y. Y. Wang, S. Z. Xing, Y.
971	S. Chen, Y. Sun, J. Li, P. Daszak, L. F. Wang, Z. L. Shi, Y. G.
972	Tong, and J. Y. Ma. 2018. 'Fatal swine acute diarrhoea syndrome
973	caused by an HKU2-related coronavirus of bat origin', <i>Nature</i> ,
974	556: 255-58.
975	

976 Figure legends

#### 977 Figure 1. The chemical structure and synthesis of 69-0 prodrugs.

- 978 Reagent and condition: i) anhydrides, DMAP, EDMA, ACN, 40 °C, 0.5 h; ii) 2,2-
- 979 Dimethoxypropane, Conc. H<sub>2</sub>SO<sub>4</sub>, Acetone, rt~45 °C, 4 h; iii) carboxylic acid, DCC,
- 980 DMAP, DCM, rt, 12 h; iv) 6 N HCl, THF, 0 °C, 7 h.

#### 981 Figure 2. Antiviral activity of compounds against SARS-CoV-2 variants (B.1, Beta

- 982 and Delta) in Vero E6 cells.
- 983 Vero-E6 cells were infected with different strains of SARS-CoV-2 variant (B.1, Beta
- and Delta) at an MOI of 0.05 and treated with dilutions of compounds (0, 0.01, 0.1, 0.5,
- 985 1, 2, 5, 10 and 50  $\mu$ M) for 48 h. Viral yield in the cell supernatant was then quantified
- 986 by qRT-PCR. The values of  $EC_{50}$  of each compound were analyzed.

#### 987 Figure 3. Pharmacokinetic profile of ATV006 in SD rats and cynomolgus monkeys.

988 (A) Time-plasma concentration curve of the nucleoside following single IV (4mg/kg) 989 or IG (20 mg/kg) administration of ATV006 to SD rats (n = 3, Mean  $\pm$  SD). (B) Plasma 990 concentration of the nucleoside following single IV (5 mg/kg) or IG (10 mg/kg) 991 administration to cynomolgus monkeys (n = 3, Mean  $\pm$  SD). (C) Tissue distribution of 992 the parent nucleoside followed a single oral dose of 100 mg/kg ATV006 to C57BL/6 993 mice (n = 5, Mean  $\pm$  SD).

### Figure 4. Anti-SARS-CoV-2 efficacy of ATV006 in hACE2 humanized and Ad5hACE2 mouse model.

996 (A) Schematic of the experiment viral infection in hACE2 humanized mice. hACE2 997 humanized mice were intranasally inoculated with B.1 original strain of SARS-CoV-2 998 (2x10<sup>5</sup> PFU virus per mouse) and were administered with vehicle (control), ATV006 999 (250 mg/kg, IG, once daily), ATV006 (500 mg/kg, IG, once daily). Viral titers in the 1000 lungs at 4 dpi were measured by qRT-PCR analysis of gRNA N (B) and sgRNA N (C) 1001 of SARS-CoV-2. (D) Schematic of the experiment viral infection in Ad5-hACE2 mice. 1002 B.1 original strain of SARS-CoV-2 (1 x 10<sup>5</sup> PFU virus per mouse) infected Ad5-hACE2 1003 mice were administered with vehicle (control), ATV006 (250 mg/kg, IG, once daily), 1004 EIDD-2801 (500 mg/kg, IG, once daily) beginning at -2 hpi. (E) Viral titers in the lungs 1005 of treated or untreated Ad5-hACE2 mice at 2 dpi were measured by plaque assay. (F) 1006 Histopathology analysis of lung from vehicle group and ATV006 (250 mg/kg) group at 1007 4 dpi. \*p values  $\le 0.05$ ; \*\*p values  $\le 0.005$ ; \*\*\*p values  $\le 0.0005$ ; \*\*\*\*p values  $\le$ 1008 0.0001.

## Figure 5. Anti-SARS-CoV-2 Delta variant efficacy of ATV006 in K18 hACE2 mouse model.

1011 (A) Schematic of the experiment. K18 hACE2 mice were intranasally inoculated with

1012 SARS-CoV-2 Delta variant (1 x 104 plaque forming units (PFU) virus per mouse) and

- 1013 were treated with vehicle (control, n=11), ATV006 (250 mg/kg, IG, once daily, n=11),
- 1014 ATV006 (100 mg/kg, IG, once daily, n=8) or EIDD-2801 (500 mg/kg, IG, once daily,

1015 n=8). (B) Survival curve. (C) Body weight curve. (D) Viral titers from lungs and brains 1016 tissue were harvested at 3 dpi and analyzed by qRT-PCR. Histopathology (F, H) and 1017 gross pathology (E, G) of lungs and spleens from vehicle group and ATV006 (250 1018 mg/kg, IG, once daily). (I) Representative chemokines and cytokines assessment of the 1019 lung tissues harvested at 3 dpi of the vehicle group and 250 mg/kg ATV006 group. 1020 Total RNA were extracted from lung homogenates and IFN-β, IFN-γ, CXCL10 and 1021 CCL2 were analyzed by qRT-PCR. \*p values  $\leq 0.05$ ; \*\*p values  $\leq 0.005$ ; \*\*\*p values

1022  $\leq 0.0005$ ; \*\*\*\*p values  $\leq 0.0001$ .

## Figure 6. ATV006 has broad-spectrum antiviral activity among different coronaviruses.

(A) L2 cells were infected with MHV-A59 at a multiplicity of infection (MOI) of 0.1
and treated with dilutions of remdesivir, 69-0 and ATV006. Antiviral activities were
evaluated by qRT-PCR quantification of a viral copy numbers in the cultured
supernatant after 16 h post infection. (B-F) The values of EC<sub>50</sub> of ATV006 in (FIPV,
CCoV, PEDV, TGEV and SADS) were analyzed.

#### 1030 Figure 7. Anti-viral efficacy of ATV006 against MHV in vivo.

1031 (A)Schematic of the experiment. 5 weeks Balb/c mouse were intranasally inoculated 1032 with  $1 \ge 10^6$  PFU per mouse of MHV-A59, and treated with vehicle (control, n=5) and 1033 ATV006 (500, 250, 100, 50 mg/kg, IG, twice daily, n=5 per group). (B) Body weight. 1034 (C, D, E, F) Viral titers from lungs and livers tissue were harvested at 2 dpi and analyzed 1035 by plaque assay and qRT-PCR. (G) Histopathology analysis of lung from vehicle group 1036 and ATV006 (500 mg/kg, IG, twice daily). (H) Representative chemokine and cytokine 1037 assessment of the lung tissues of the indicated groups, as detected in lung tissue 1038 homogenate at 2 dpi. \*p values  $\leq 0.05$ ; \*\*p values  $\leq 0.005$ ; \*\*\*p values  $\leq 0.0005$ ; \*\*\*\*p 1039 values  $\leq 0.0001$ .

#### 1040 Figure S1. Antiviral activity of 21 compounds in SARS-CoV-2 Replicon system.

- 1041 (A) HEK 293T cells transfected with SARS-CoV-2-Rep-Luci were treated with DMSO
- and 24 compounds with 10  $\mu$ M. 60 h post-transfection, the cells were subjected to the
- 1043 Dual-Luciferase® Reporter (DLR<sup>TM</sup>) Assay.
- 1044 (B) The values of  $EC_{50}$  of each compound were analyzed.

### Figure S2. Antiviral activity of compounds against SARS-CoV-2 (B.1) in Huh7 cells.

- 1047 Huh7 cells were infected with B.1 original strain SARS-CoV-2 at an MOI of 0.05 and
- 1048 treated with dilutions of each compound  $(0, 0.01, 0.1, 0.5, 1, 2, 5 \text{ and } 10 \,\mu\text{M})$  for 48 h.
- 1049 Viral yield in the cultured supernatant was then quantified by qRT-PCR. The values of
- 1050  $EC_{50}$  of each compound was analyzed.
- 1051 **Figure S3. Cytotoxicity assay of compounds.**
- 1052 Vero-E6 cells were plated in 96-well plate and treated with increasing concentrations
- 1053 compound ranging from 0 to 200 µM for 48 h. Cell viability was tested using Cell
- 1054 Counting Kit-8 (CCK-8). (A) ATV006. (B) other compounds.

#### 1055 Figure S4. Genomic RNA (gRNA) and Subgenomic RNA (sgRNA) of Coronavirus.

1056 Detect the target position of the primer/probe sets of genomic RNA (gRNA) and

subgenomic RNA (sgRNA) of Coronavirus. In this article, only FP and RP were used
to detect sgRNA of SARS-CoV-2, and no Prb was used. The sequences of primer/probe

1059 sets are listed in Table S3.

### 1060 Figure S5. Dose-response in vivo anti-MHV efficacy of ATV006, remdesivir and

#### 1061 **69-0 via intranasal inoculation.**

1062 (A) Schematic of the experiment. Mouse were divided them into the following groups:

- 1063 vehicle (control), ATV006 (50, 20, 10, 5, 2 mg/kg, IG, once daily), remdesivir (20
- 1064 mg/kg, IV, once daily) or 69-0 (50 mg/kg, IG, once daily). After MHV-A59 infects the
- 1065 mice, the administration is continued for 4 days, and the body weight curve (C) and

survival curve (B) of the mice were recorded for 14 days. (D) Viral titers from liverswere harvested at 3 dpi and analyzed by qRT-PCR.

### Figure S6. Dose-response anti-HMV efficacy of ATV006 via intrahepatic inoculation.

1070 (A) Schematic of the experiment. 5 weeks Balb/c mouse were intranasally

- 1071 intrahepatic with  $1 \ge 10^5$  PFU per mouse of MHV-A59, and treated with vehicle
- 1072 (control), ATV006 (50,10, 2 mg/kg, IG, twice daily). (B, C and D) Viral titers from
- 1073 liver tissue were harvested at 3 dpi and analyzed by plaque assay and qRT-PCR. (E
- 1074 and F) ALT and AST analysis of serum from vehicle group and ATV006 (50 mg/kg,
- 1075 IG, twice daily). (G) Histopathology analysis of liver from vehicle group and
- 1076 ATV006 (50 mg/kg, IG, twice daily). (H) Representative chemokine and cytokine
- 1077 assessment of the liver tissues of the indicated groups, as detected in lung tissue
- 1078 homogenate at 3 dpi. \*p values  $\leq 0.05$ ; \*\*p values  $\leq 0.005$ ; \*\*\*p values  $\leq 0.0005$ ; \*

1079 \*\*\*p values  $\leq 0.0001$ .

#### 1080 Figure S7. Analysis of RdRp mutation of SARS-CoV-2 and its variants.

The black triangles represent the key sites where remdesivir binds to RdRp. The red triangles represent the enzyme activity amino acid residues of RdRp. The red dots represent the amino acid sites whose mutation rate is greater than one percent and less than eighty percent compared with the original strain. The blue dots represent the amino acid sites whose mutation rate is greater than eighty percent compared with the original strain.

#### 1087 Figure S8. Coronavirus RdRp Conservation Analysis.

1088 Coronavirus RdRp amino acid sequence alignment, (A) Alpha-coronavirus and (B)

1089 beta-coronavirus. The black dots represent the key sites where remdesivir binds to

1090 RdRp. The red triangles represent the enzyme activity amino acid residues of RdRp.

1091

#### **Tables**

#### 

### Table 1. Anti-SARS-CoV-2 activity and cytotoxicity of the adenosine analogue prodrugs in comparison with remdesivir and 69-0

	EC <sub>50</sub> (µM) in Vero E6 cells			
-	SARS-	SARS-CoV-2	SARS-CoV-2	_
Compound	CoV-2	(Beta,	(Delta,	$CC_{50}$ ( $\mu$ M)
	(B.1)	B.1.351)	B.1.617.2)	
Remdesivir	2.279	1.780	1.645	>50
69-0	1.709	1.354	0.957	>50
ATV006	1.360	1.127	0.349	128.00
ATV009	1.329	1.484	0.492	>50
ATV010	0.696	1.002	0.457	44.62
ATV011	2.117	2.302	0.408	>50
ATV013	2.262	2.434	0.965	>50
ATV017	2.188	2.847	0.428	>50

**Table 2.** Pharmacokinetic profile of ATV006 in Sprague-Dawley (SD) rat and

1100 cynomolgus monkey <sup>a</sup>

	rat		monkey	
parameters	IV (4mg/kg)	IG (20mg/kg)	IV (5mg/kg)	IG (10mg/kg)
$AUC_{last} (\mu g/L*h)$	2347±354	11445±813	5960±490	3560±245
T <sub>1/2</sub> (h)	1.30±0.24	3.62±0.61	$1.78 \pm 0.6$	$4.08 \pm 0.94$
$T_{max}(h)$	0.083	$0.83 \pm 0.29$	0.083	$1.50\pm2.20$
$C_{max}(\mu g/L)$	3030+451	4017±359	3730±709	$1080 \pm 651$
F (%)		98%		30.08%

1101	<sup>a</sup> n=3, ATV006 was administrated using the indicated route, and the parameters were
1102	calculated based on the LC-MS-MS analysis of parent nucleoside 69-0. IV,
1103	intravenous; IG, intragastric.

**Table S1.** Permeability and efflux ratio determination of 69-0, ATV006, ATV019 and ATV020 in Case 2 cells

1107	A I V020 in Caco-2 cells		
	Compound	Caco-2 AB/BA (Papp (10	

Compound	Caco-2 AB/BA (Papp (10 <sup>-6</sup> cm/s)) <sup>a</sup>	Efflux ratio
69-0	1.22/1.20	0.98
ATV006	0.51/0.87	1.7

ATV019	0.28/0.68	2.47
ATV020	0.17/0.22	1.28

<sup>a</sup> Papp (A to B) < 2, low permeability; 2 < Papp (A to B) < 10, moderate permeability;

1110 Papp (A to B) > 10, high permeability.

#### **Table S2.** Animal coronaviruses and cells used in this study

Virus	Cell line	Concentrations for ATV006	Virus strain
Canine coronavirus (CCV)	Feline kidney cells (CRFK cells)	ranging from 0.1μM to 50 μM	ATCC-VR-2068
Porcine epidemic diarrhea virus (PEDV)	African green monkey kidney cells (Vero E6 cells)	ranging from 0.1 μM to 50 μM	isolated by our lab
Transmissible gastroenteritis virus (TGEV)	swine testicle cells (ST cells)	ranging from 0.5 μM to 100 μM	ATCC-VR-1740
Swine acute diarrhea syndrome coronavirus (SADS-CoV)	African green monkey kidney cells (Vero E6 cells)	ranging from 0.5 μM to 100 μM	SADS- CoV/CN/GDST/2017
Feline infectious peritonitis virus (FIPV)	Feline kidney cells (CRFK cells)	ranging from 0.1 μM to 100 μM	ATCC-VR-2128
Mouse hepatitis virus (MHV)	Rat lung epithelial cells (L2 cells)	ranging from 0.01 μM to 10 μM	were kindly provided by Rong Ye (Shanghai Medical School of Fudan University)

### 

**Table S3.** qPCR primers used for detection of viral genomes and various genes

Gene		Sequence(5'-3')
DA'AN	FP	AAGAAATTCAACTCCAGGCAGC
SARS-COV-2-N	RP	GCTGGTTCAATCTGTCAAGCAG
	Prb	TCACCGCCATTGCCAGCCA
SARS-COV-2	FP	CCAGGTAACAAACCAACAA
sgN	RP	TGAGTGAGAGCGGTGAACCAA
MHV-A59-N	FP	GGAACTTCTCGTTGGGCATTATACT

	DD	
	RP	ACCACAAGATTATCATTTTCACAACATA
	Prb	ACATGCTACGGCTCGTGTAACCGAACTGT
MHV-A59- sgN	FP	TATAAGAGTGATTGGCGTCC
	RP	GAGTAATGGGGAACCACACT
	Prb	ACATGCTACGGCTCGTGTAACCGAACTGT
m-GAPDH	FP	AGAACATCATCCCTGCATCC
	RP	CACATTGGGGGGTAGGAACAC
m-IL-6	FP	AACCAAGAGATAAGCTGGAGTCAC
	RP	AACGCACTAGGTTTGCCGAG
m-IL1β	FP	TGCCACCTTTTGACAGTGATGA
	RP	ATCAGGACAGCCCAGGTCAA
m-CXCL-10	FP	TGCAGGATGATGGTCAAGCC
	RP	CCACTTGAGCGAGGACTCAG
m-IFN-γ	FP	CAGCAAGGCGAAAAAGGATGC
	RP	CTTCCTGAGGCTGGATTCCG
m-IFN-β	FP	GTGGGAGATGTCCTCAACTGC
	RP	TCTCTGCTCGGACCACCATC
m-CCL2	FP	TGGGCCTGTTGTTCACAGT
	RP	TTCTCCAGCCGACTCATTG
FIPV	FP	AGCAACTACTGCCACRGGAT
	RP	GGAAGGTTCATCTCCCCAGT
	Prb	AATGGCCACACAGGGACAACGC
CCoV,	FP	CAGTCTAGAAATAGATCTCAATC
	RP	GCTTGTTCTACACTGTCA
	Prb	CCTTCTTGTTATTGGATTGTTGCCTTC
PEDV	FP	CGCAAAGACTGAACCCACTAATTT
	RP	TTGCCTCTGTTGTTACTTGGAGAT
	Prb	TGTTGCCATTGCCACGACTCCTGC
TGEV	FP	GCAGGTAAAGGTGATGTGACAA
	RP	ACATTCAGCCAGTTGTGGGTAA
	Prb	TGGCACTGCTGGGATTGGCAACGA-
SADS	FP	CTGACTGTTGTTGAGGTTAC
	RP	TCTGCCAAAGCTTGTTTAAC
	Prb	TCACAGTCTCGTTCTCGCAATCA

1117

### 

#### 

#### **Table S4.** Viral genome sequences and accession numbers

Viruses	Accession Number
Feline Infectious Peritonitis Virus (FIPV)	YP_004070193.2
Bat coronavirus 1B (BCoV-1B)	ACA52156.1
bat coronavirus 512 (Sc-BatCoV-512)	YP_001351683.1
bat coronavirus HKU2 (BCHV2)	ABB77027.1
bat coronavirus HKU6 (BatCoVHKU6)	ABB77038.1
bat coronavirus HKU7 (BatCoVHKU7)	ABB77040.1
bat coronavirus HKU8 (BatCoVHKU8)	YP_001718610.1
Canine coronavirus (CCoV)	AEQ61967.2
Ferret coronavirus (FRCoV)	AKG92638.1
Human coronavirus 229E (HCoV-229E)	NP_073549.1
Human coronavirus NL63 (HCoV-NL63)	YP_003766.2
Porcine epidemic diarrhea virus (PEDV)	NP_598309.2
Transmissible gastroenteritis virus (TGEV)	NP_058422.1
Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)	YP_009724389.1
Severe acute respiratory syndrome coronavirus (SARS)	NP_828849.7
Swine acute diarrhea syndrome coronavirus (SADS-CoV)	QJF53984.1
bat coronavirus HKU5(BatCoVHKU5)	YP_001039961.1
Bovine coronavirus (BCoV)	AAL57305.1
Human coronavirus OC43 (HCoV-OC43)	YP_009555238.1
Rousettus bat coronavirus HKU9 (BatCoVHKU9)	YP_009924393.1
Human coronavirus HKU1(HCHV1)	YP_173236.1
Middle East respiratory syndrome-related coronavirus (MERS)	YP_009047202.1
Mouse hepatitis virus (MHV)	AAB86818.1
Mouse hepatitis virus (MHV)	AAB86818.1

Lucheng Rn rat coronavirus (LRNV)

QDL88264.1

1121 1122 1123

### Figures

### Figure 1. The chemical structure and synthesis of 69-0 prodrugs.

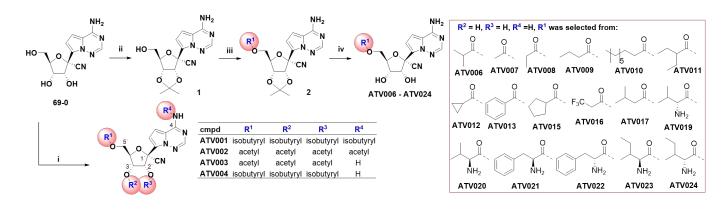
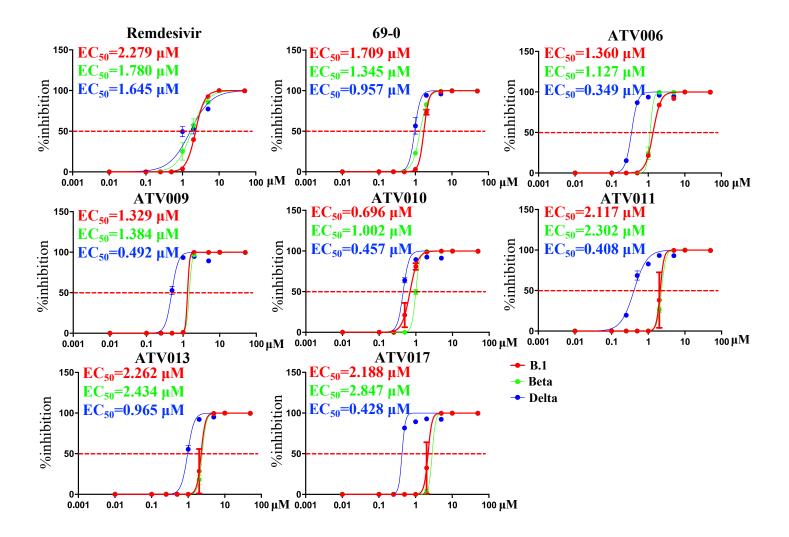
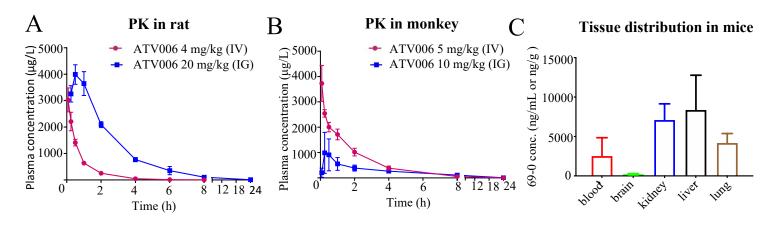


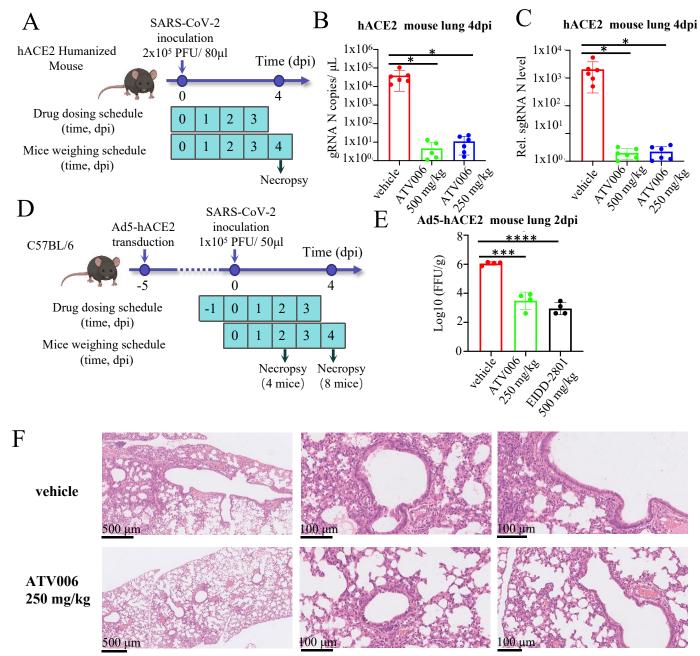
Figure 2. Antiviral activity of the compounds against SARS-CoV-2 variants (B.1, Beta and Delta) in Vero E6 cells.



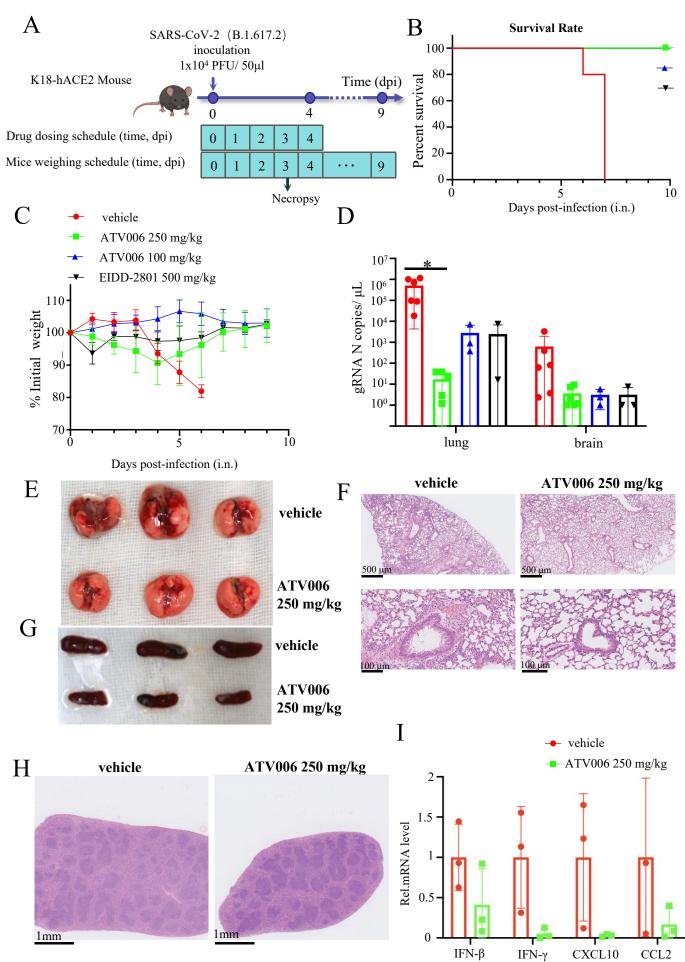
# Figure 3. Pharmacokinetic profile of ATV006 in SD rats and cynomolgus monkeys.



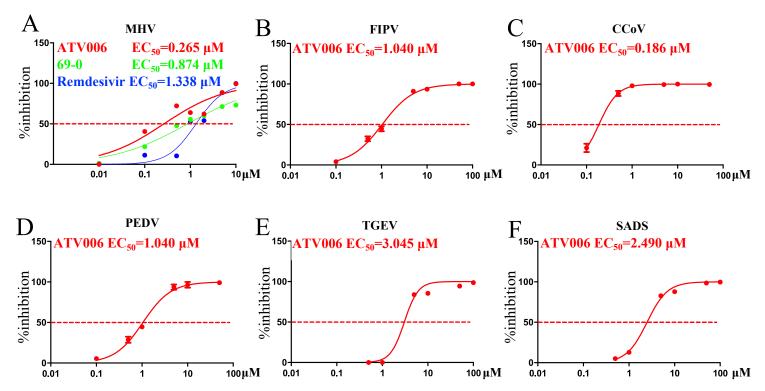
# Figure 4. Anti-SARS-CoV-2 efficacy of ATV006 in hACE2 humanized and Ad5-hACE2 mouse model.



# Figure 5. Anti-SARS-CoV-2 Delta variant efficacy of ATV006 in K18 hACE2 mouse model.



# Figure 6. ATV006 has broad-spectrum antiviral activity among different coronaviruses.



### Figure 7. Anti-viral efficacy of ATV006 against MHV in vivo.

