

Article

Not peer-reviewed version

A Soluble ACE2 Protein Improves Survival and Lowers Viral Titers in a Lethal Mouse Model of SARS-CoV-2 Infection with the Delta Variant

Cosimo Cianfarini , Luise Hassler , [Jan Wysocki](#) , Abdelsabour Hassan , Vlad Nicolaescu , Derek Elli , Haley Gula , Amany M. Ibrahim , [Glenn Randall](#) , Jack Henkin , [Daniel Batlle](#) *

Posted Date: 25 December 2023

doi: 10.20944/preprints202312.1799.v1

Keywords: SARS-CoV-2, COVID-19, Angiotensin-Converting-Enzyme 2, ACE2



Preprints.org is a free multidiscipline platform providing preprint service that is dedicated to making early versions of research outputs permanently available and citable. Preprints posted at Preprints.org appear in Web of Science, Crossref, Google Scholar, Scilit, Europe PMC.

Copyright: This is an open access article distributed under the Creative Commons Attribution License which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Article

A Soluble ACE2 Protein Improves Survival and Lowers Viral Titers in a Lethal Mouse Model of SARS-CoV-2 Infection with the Delta Variant

Cosimo Cianfarini ^{1,2}, Luise Hassler ¹, Jan Wysocki ¹, Abdelsabour Hassan ¹, Vlad Nicolaescu ³, Derek Elli ³, Haley Gula ³, Amany M. Ibrahim ³, Glenn Randall ³, Jack Henkin ⁴ and Daniel Batlle ^{1,*}

¹ Division of Nephrology/Hypertension, Department of Medicine, Northwestern University, Feinberg School of Medicine, Chicago, Illinois

² Charité Universitätsmedizin Berlin, Berlin, Germany

³ Department of Microbiology, Howard Taylor Ricketts Laboratory, The University of Chicago, Lemont, Illinois

⁴ Center for Developmental Therapeutics, Northwestern University, Evanston, Illinois

* Correspondence: Daniel Batlle MD, Earle, del Greco Levin Professor of Nephrology/Hypertension, Northwestern University Feinberg School of Medicine, Division of Nephrology/Hypertension, 710 North Fairbanks Court, Chicago, IL 60611, USA, Phone: +1 312 908 8342, Email: d-batlle@northwestern.edu

Abstract: Severe acute respiratory syndrome coronavirus type 2 (SARS-CoV-2) utilizes Angiotensin Converting Enzyme 2 (ACE2) as its main receptor for cell entry. We have bioengineered a soluble ACE2 protein termed ACE2 618-DDC-ABD that has increased binding to SARS-CoV-2 and prolonged duration of action. Here we investigated the protective effect of this protein when administered intranasally to k18-hACE2 mice infected with the aggressive SARS-CoV-2 delta variant. k18-hACE2 mice were infected with the SARS-CoV-2 delta variant by inoculation of a lethal dose (2×10^4 PFU) and followed for up to 14 days. ACE2-618-DDC-ABD (10mg/kg) or PBS was administered intranasally six hours prior and 24 and 48 hours post viral inoculation. All animals in the PBS-control group had succumbed to the disease on day 7 post infection (0% survival) whereas, by contrast, there was only one casualty in the group that received ACE2-618-DDC-ABD (90% survival). Mice in the ACE2-618-DDC-ABD group had minimal disease as assessed by a clinical score and stable weight and both brain and lung viral titers were markedly reduced. These findings demonstrate the efficacy of a bioengineered soluble ACE2 decoy with extended duration of action in protecting against the aggressive delta SARS-CoV-2 variant. Together with previous work these findings demonstrate the universal protective potential against current and future emerging SARS-CoV-2 variants.

Keywords: SARS-CoV-2; COVID-19; Angiotensin-Converting-Enzyme 2; ACE2

1. Introduction

Since its first appearance in Wuhan, China in December 2019, severe acute respiratory syndrome coronavirus type 2 (SARS-CoV-2) spread worldwide posing a threat to public health [1,2]. Four years later and despite massive improvements largely because of the vaccination efforts, new cases are frequently reported [3,4]. Omicron variants currently are the most commonly variants seen worldwide [5–7]. SARS-CoV-2, like other coronaviruses including its predecessor SARS-CoV-1, uses membrane bound Angiotensin Converting Enzyme 2 (ACE2) as its primary cell entry receptor [8–11]. The complex multistep process of SARS-CoV-2 cell entry starts with interaction of the receptor binding domain (RBD) of the viral S1 spike protein with ACE2 on the cell surface [12–15]. Conformational changes then lead to priming of the viral spike protein by proteases like transmembrane protease serine 2 (TMPRSS2) which results in fusion pore formation and ultimately cell infection [9,16–18].

At the beginning of the COVID-19 pandemic in early 2020 our group proposed the use of soluble ACE2 proteins to intercept the binding of SARS-CoV-2 to its receptor, membrane bound ACE2, via

the so called “decoy effect” [19,20]. We had bioengineered a human soluble ACE2 protein that was shortened from 740 to 618 amino acids and fused with an albumin binding domain (ABD) to increase its duration of action [21]. Later a dodecapeptide (DDC) motif that contains four cysteine residues was introduced to enhance binding affinity for SARS-CoV-2 via formation of a disulfide bonded dimeric protein [21,22]. In human kidney organoids and the k18-hACE2 mouse model of lethal SARS-CoV-2 infection we demonstrated the potential of our bioengineered soluble ACE2 proteins to neutralize SARS-CoV-2, markedly improve survival and provide *in-vivo* lung, brain and kidney protection [21,22]. We established preventative, intranasal delivery, starting one hour before viral inoculation followed by two additional doses at 24 and 48 hours post viral inoculation, as an effective mode of ACE2-618-DDC-ABD delivery achieving optimal protection in terms of survival and organ injury in the k18-hACE2 mouse model infected with ancestral SARS-CoV-2 [23].

As new variants of SARS-CoV-2 emerge, mutational escape and resistance to treatment are critical issues and both antibodies and antiviral drugs have been shown to be associated with the development of mutational escape of SARS-CoV-2 [24,25]. Mutations of SARS-CoV-2 that would decrease ACE2 decoy affinity while simultaneously maintaining efficacious binding of SARS-CoV-2 to the membrane bound ACE2 receptor have not been described and are unlikely to occur [26,27]. This highlights an advantage of soluble ACE2 proteins over monoclonal antibodies [28]. Studies in permissive cells have shown that soluble ACE2 proteins moreover are effective against different SARS-CoV-2 variants [10,21–23,29–31]. Here we investigated intranasal administration of ACE2-618-DDC-ABD to k18-hACE2 mice infected with the SARS-CoV-2 delta variant which causes severe disease in humans to further demonstrate the universal efficacy of our unique soluble ACE2 protein. Moreover, ACE2-618-DDC-ABD was applied as early as six hours prior viral inoculation to further determine its lasting protective effect and extended duration of action.

2. Materials and Methods

2.1. *In-vivo* infectivity studies

All *in-vivo* infectivity studies were performed at the BSL-3 facility of the Ricketts Regional Biocontainment Laboratory in accordance with a protocol approved by the Institutional Animal Care and Use Committees of Northwestern University (IS00004795) and the University of Chicago (72642).

k18-hACE2 mice were purchased from The Jackson Laboratory at 6-9 weeks of age. k18-hACE2 mice express full-length human ACE2 (hACE2) with an estimated hACE2 gene copy number of 8 and are susceptible to infection with SARS-CoV-2 [22,23,32–37]. Animals were infected by intranasal inoculation of 2×10^4 plaque-forming units (PFU) SARS-CoV-2 delta variant (Isolate hCoV-19/USA/MD-HP05647/2021 Lineage B.1.617.2; Delta variant) in a volume of 20 μ l per animal. Disease severity in k18-hACE2 mice infected with the SARS-CoV-2 delta variant has been reported to be similar to or greater than infection with the ancestral virus, where animals typically succumb to disease by days 5-9 post viral inoculation [22,23,32–35,38,39].

2.2. Administration of ACE2-618-DDC-ABD

For the following experiments two groups of k18-hACE2 mice (n=10 per group, five male and five female each) administered with either ACE2-618-DDC-ABD or phosphate buffered saline (PBS) as vehicle control were investigated. ACE2 618-DDC-ABD was administered intranasally (10 mg/kg BW in a volume of 30 μ l per animal) for a total of three doses. The first dose was given six hours before inoculation of the animal with the SARS-CoV-2 delta variant followed by the same dose 24 and 48 hours after viral inoculation. PBS as vehicle control was administered following the same protocol at the same time points.

2.3. Evaluation of body weight and clinical score

Following inoculation of the animals with the SARS-CoV-2 delta variant all mice were monitored for health two to three times a day using a clinical scoring system as previously reported [22,23]. Additionally, animals were weighed once daily throughout the 14-day study period. Mice

that lost more than 20% of their initial body weight and did not regain it within two days or had a severely worsened clinical score were humanely euthanized. This was considered a fatal event as per study protocol.

To compare viral titers at similar time points, randomly selected animals of each sex from both groups were euthanized on day six post viral inoculation together with the animals that were euthanized because of disease severity. All remaining mice were monitored for up to 14 days at the BSL-3 facility of the Ricketts Regional Biocontainment Laboratory and euthanized at day 14 at the end of the study protocol if not previously deceased. Mice were euthanized by CO₂ forced inhalation. After the last breathing movement, cervical dislocation was performed and lung, brain and kidney tissue were collected for plaque assay analysis (see below).

2.4. Plaque Assay

Lung, brain, and kidney tissue was removed from all euthanized animals and used for viral load measurements by plaque assay. Organs from one mouse in the PBS-control group could not be collected, and the animal was therefore not included in the plaque assay analysis.

Tissue samples for plaque assay analysis were collected in Dulbecco's Modified Eagle Medium (DMEM) with 2% Fetal Bovine Serum (FBS). Samples were then homogenized and centrifuged at 1000g for five minutes. The supernatant was serially diluted 10-fold and used to infect Vero E6 cells for one hour. Inoculum was removed and 1.25% methylcellulose DMEM solution was added to the cells followed by incubation for three days. Plates were then fixed in 1:10 formalin for one hour, stained with crystal violet for one hour, and counted to determine PFU expressed as PFU/ml.

2.5. Statistical Analysis

Statistical analysis and figure preparation were performed using GraphPad Prism v. 9.5.1. Normal distribution was tested using the Shapiro-Wilk test. When data was normally distributed, differences between two groups were analyzed by unpaired t-test. Differences between groups with non-normally distributed data were analyzed by Mann-Whitney test. Differences in survival were analyzed by log-rank (Mantel-Cox) test. A p-value < 0.05 was considered significant.

3. Results:

3.1. Survival of k18-hACE2 mice infected with SARS-CoV-2 delta variant

All infected k18-hACE2 mice that received PBS either died or had to be humanely euthanized by day seven post viral inoculation as per the study protocol (**Figure 1, black**). In contrast, survival of the mice that received ACE2-618-DDC-ABD was 90% on day six post viral inoculation. Only one mouse in this group had to be humanely euthanized on day six post viral inoculation (**Figure 1, purple**). To compare viral titers at similar time points, five healthy mice from the ACE2 618-DDC-ABD treated group were sacrificed on day six post viral inoculation. The remaining four mice in this group all survived until day 14, the end of the study period. This marked difference in survival was statistically significant ($P=0.0004$, by log-rank (Mantel-Cox) test).

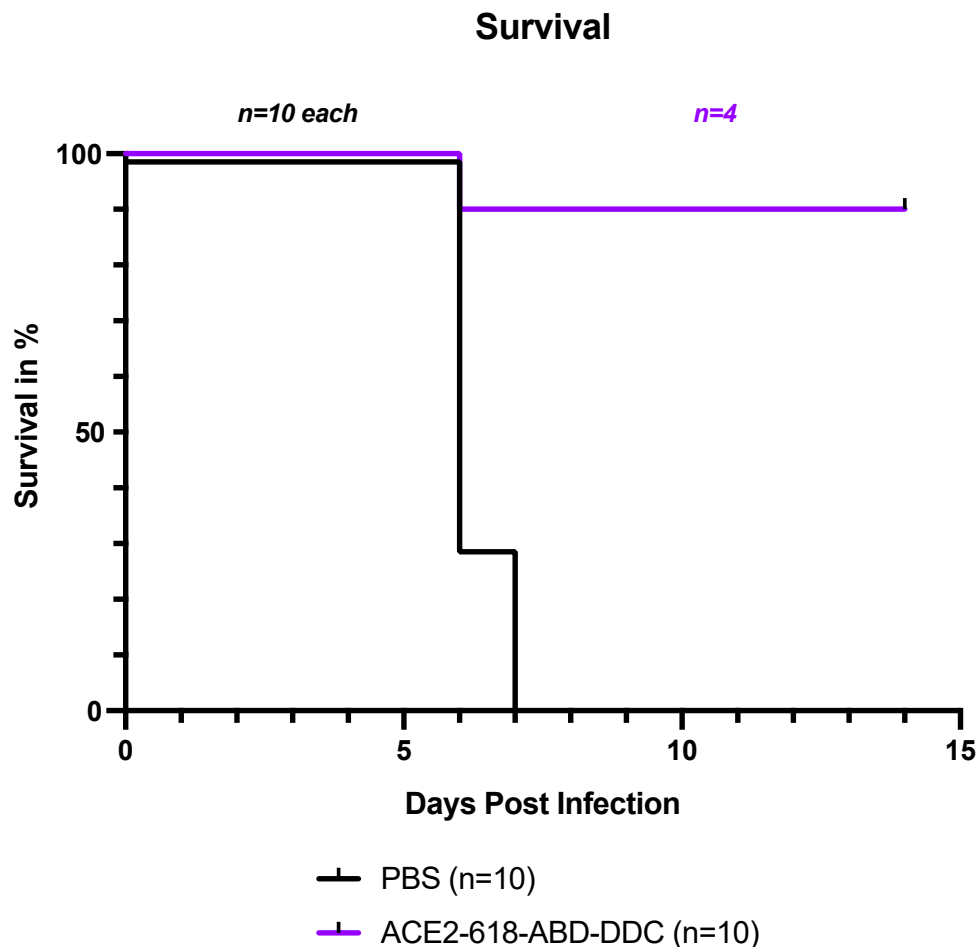


Figure 1. Survival of k18-hACE2 mice infected with the SARS-CoV-2 delta variant that received either ACE2-618-DDC-ABD or PBS. Infected animals in the PBS-control group (black) showed uniform lethality and survival by day seven post viral inoculation was 0%. By contrast, in the ACE2-618-DDC-ABD treated group (purple), survival was 90% by day six post viral inoculation with only one mouse having to be humanely euthanized on day six. Five mice from the ACE2-618-DDC-ABD treated group that were healthy by clinical score and weight were sacrificed on day six post viral inoculation for comparison of viral titers at similar time points. The remaining four mice in the ACE2 618-DDC-ABD treated group all survived until the end of the study protocol at day 14 post viral inoculation. The difference in survival was statistically significant ($P=0.0004$ by log-rank (Mantel-Cox) test).

3.2. Clinical score of k18-hACE2 mice infected with SARS-CoV-2 delta variant

All infected animals were evaluated for health by a clinical score two to three times a day (see material and methods section). Infected k18-hACE2 mice that received PBS as vehicle control developed severe disease and clinical scores increased until day six and seven post viral inoculation (**Figure 2A, black**). At that point all animals in the PBS-vehicle group had either died or had to be humanely euthanized as per the study protocol (**Figure 1**). Animals that received ACE2-618-DDC-ABD developed less severe disease and clinical scores were lower (**Figure 2A, purple**). By the end of the 14-day study period all remaining mice in the ACE2-618-DDC-ABD group had recovered.

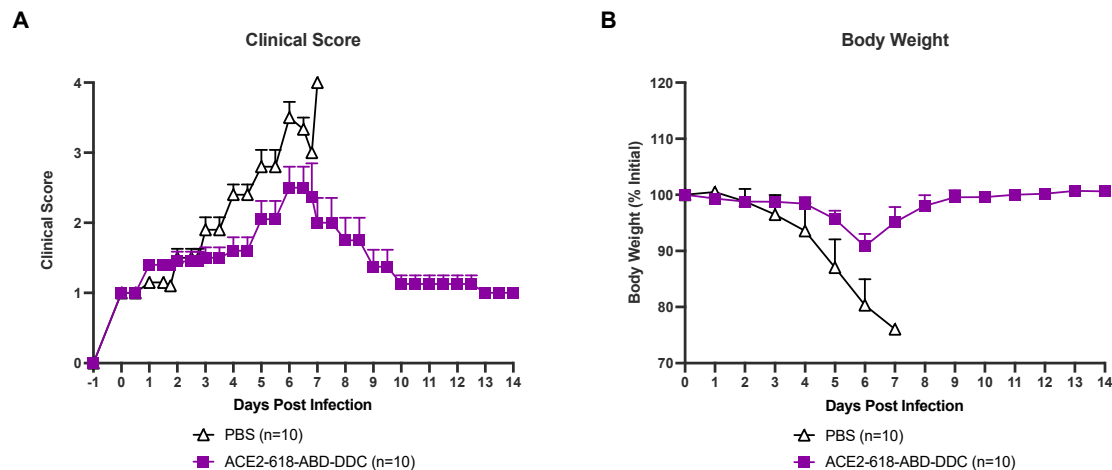


Figure 2. Clinical score and body weight of k18-hACE2 mice infected with the SARS-CoV-2 delta variant that received either ACE2-618-DDC-ABD or PBS. Panel A. Infected animals in the PBS-control group (black) developed severe disease and clinical scores increased accordingly until days six and seven post viral inoculation by which time point all mice had either died or had to be humanely euthanized. By contrast, animals in the ACE2-618-DDC-ABD treated group (purple) developed less severe disease and the clinical score was lower as compared to the PBS-control group. All remaining animals from the ACE2-618-DDC-ABD treated group had recovered by the end of the 14-day study period. Panel B. Infected animals in the PBS-control group (black) lost up to ~20% of their initial bodyweight until days six and seven post viral inoculation by which time point all mice had either died or had to be humanely euthanized. By contrast, animals in the ACE2-618-DDC-ABD treated group (purple) lost on average less than ~10% of their initial body weight by days six and seven post viral inoculation. All remaining animals from the ACE2-618-DDC-ABD treated group recovered stable body weight by the end of the 14-day study period.

3.3. Weight loss of k18-hACE2 mice infected with SARS-CoV-2 delta variant

All infected mice were weighed once daily throughout the 14-day study period. Infected mice that received PBS lost up to ~20% of their initial bodyweight until days six and seven post viral inoculation. At this point all mice from the PBS-control group either died or had to be humanely euthanized (**Figure 2B, black**). Mice that received ACE2-618-DDC-ABD lost on average less than ~10% of their initial body weight by day six post viral inoculation and the remaining mice in this group recovered stable body weight towards the end of the 14-day study period (**Figure 2B, purple**).

3.4. Lung viral titers of k18-hACE2 mice infected with SARS-CoV-2 delta variant

Lung viral titers were assessed in all ten mice that received ACE2-618-DDC-ABD and in nine mice that received PBS. Lung viral titers in infected animals that received ACE2-618-DDC-ABD were reduced as compared to infected mice in the PBS-control group ($1.5 \times 10^2 \pm 0.7 \times 10^2$ vs. $7.8 \times 10^3 \pm 4.7 \times 10^3$ PFU/ml, $P=0.04$) (**Figure 3A**). When only comparing lung viral titers in animals sacrificed at the same time point on day six post viral inoculation ($n=6$ per group), a marked reduction in the ACE2-618-DDC-ABD treated group was seen as well ($2.5 \times 10^2 \pm 0.9 \times 10^2$ vs. $1.18 \times 10^4 \pm 6.7 \times 10^3$ PFU/ml, $P=0.004$). Of note, lung viral titers were completely undetectable in all remaining animals of the ACE2-618-DDC-ABD treated group at the end of the 14-day study period.

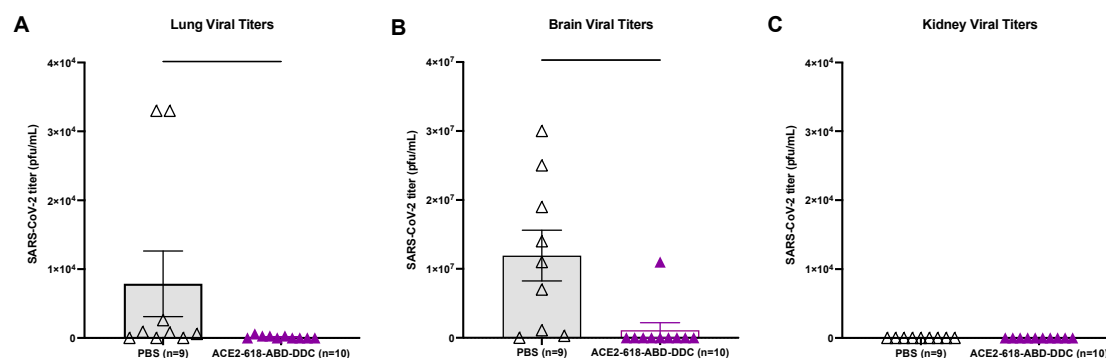


Figure 3. Lung, brain and kidney SARS-CoV-2 titers of k18-hACE2 mice infected with the SARS-CoV-2 delta variant that received either ACE2-618-DDC-ABD or PBS. Panel A. Lung viral titers in infected animals from the ACE2-618-DDC-ABD treated group (purple) were markedly reduced as compared to lung viral titers in infected animals that received PBS (black) ($1.5 \times 10^2 \pm 0.7 \times 10^2$ vs. $7.8 \times 10^3 \pm 4.7 \times 10^3$ PFU/ml, $P=0.04$). Panel B. Brain viral titers were also markedly reduced in mice that received ACE2-618-DDC-ABD (purple) as compared to brain viral titers in the PBS-control group (black) ($1.1 \times 10^6 \pm 1.1 \times 10^6$ vs. $1.2 \times 10^7 \pm 3.7 \times 10^6$ PFU/ml, $P=0.0007$). Panel C. Kidney viral titers could not be detected in any of the animals from either the ACE2-618-DDC-ABD treated (purple) or the PBS-control group (black).

3.5. Brain viral titers of k18-hACE2 mice infected with SARS-CoV-2 delta variant

Brain viral titers were assessed in all ten mice that received ACE2-618-DDC-ABD and nine mice that received PBS. Infected animals in the PBS-control group had very high brain viral titers, whereas brain viral titers were markedly reduced in infected mice that received ACE2-618-DDC-ABD ($1.1 \times 10^6 \pm 1.1 \times 10^6$ vs. $1.2 \times 10^7 \pm 3.7 \times 10^6$ PFU/ml, $P=0.0007$) (**Figure 3B**). When only comparing brain viral titers in animals sacrificed at the same time point on day six post viral inoculation ($n=6$ per group), a marked reduction in the ACE2-618-DDC-ABD treated group was seen as well ($1.8 \times 10^6 \pm 1.8 \times 10^6$ vs. $1.4 \times 10^7 \pm 5.0 \times 10^6$ PFU/ml, $P=0.008$). Similar to the data for lung viral titers, brain viral titers were completely undetectable in all remaining animals of the ACE2-618-DDC-ABD treated group at the end of the 14-day study period.

Of note, when directly comparing viral titers in brains to lungs from the PBS-control group, brain viral titers were about 1,000-fold higher than lung viral titers ($1.2 \times 10^7 \pm 3.7 \times 10^6$ vs. $7.8 \times 10^3 \pm 4.7 \times 10^3$ PFU/ml, $P=0.004$). In fact, several of the infected, untreated mice had only slightly increased lung viral titers (see Figure 3A).

3.6. Kidney viral titers of k18-hACE2 mice infected with SARS-CoV-2 delta variant

Kidney viral titers were assessed in all ten mice that received ACE2-618-DDC-ABD and in nine mice that received PBS. No kidney viral titers could be detected in either the PBS-control or the ACE2-618-DDC-ABD treated group (**Figure 3C**). The limit of detection for plaque assay is 10^2 PFU/ml.

4. Discussion

This study demonstrates a dramatic protective effect of the soluble ACE2 decoy protein termed ACE2-618-DDC-ABD in the lethal k18-hACE2 mouse model of SARS-CoV-2 infection caused by the SARS-CoV-2 delta variant. Animals that received ACE2-618-DDC-ABD six hours prior to as well as 24 and 48 hours after viral inoculation had 90% survival as compared to uniform lethality in the PBS-control group. Disease severity as evaluated by a clinical score and weight loss were markedly improved in the ACE2 618-DDC-ABD treated group. Lung and brain viral titers moreover were markedly reduced in animals receiving ACE2-618-DDC-ABD as compared to the PBS-control group. At the end of the study period on day 14 post viral inoculation all remaining animals in the ACE2-618-DDC-ABD treated group had no detectable lung and brain viral titers Taken together with our

previous studies in the same mouse model infected with the ancestral SARS-CoV-2 variant (Washington isolate) [22,23] our data demonstrates remarkable efficacy of ACE2-618-DDC-ABD in preventing mortality and organ infectivity in the aggressive delta variant. Furthermore, we also show that intranasal administration of ACE2-618-DDC-ABD prior to viral inoculation results in a protective effect when administered at least six hours prior to viral inoculation which expands our previous work when ACE2-618-DDC-ABD was administered close to viral inoculation with the ancestral SARS-CoV-2 variant (one hour before).

The SARS-CoV-2 delta variant first emerged in October 2020 and was shown to cause severe disease in humans [40–43]. Thirteen amino acid mutations have been found in the viral spike protein of the SARS-CoV-2 delta variant as compared to the ancestral virus genome [38]. Of particular interest are the mutations L452R, T478K, P681R, and D614G, which have been reported to promote infectivity and immune escape by increasing the stability of the spike protein and RBD-ACE2 interaction, as well as facilitating cleavage of the spike precursor protein and promoting viral replication and transmission [44–52]. Infection of k18-hACE2 mice with the SARS-CoV-2 delta variant induces distinct pathogenic patterns of respiratory disease [38,39]. As compared to infection with the SARS-CoV-2 alpha variant significantly more inflammation, increased production of antiviral cytokines and a greater number of activated immunological pathways have been demonstrated which underlines the uniqueness of the host response against different SARS-CoV-2 variants [39]. Of note, we found that in k18-hACE2 mice infected with the SARS-CoV-2 delta variant brain viral titers were markedly higher than lung viral titers. This is in agreement with our findings in the same model infected with the ancestral virus variant and strongly suggests that the high lethality is due to brain viral invasion [23].

We have previously reported that ACE2-618-DDC-ABD, administered to k18-hACE2 mice both intranasally and systemically one hour before as well as 24 and 48 hours after viral inoculation with the wildtype SARS-CoV-2 variant markedly improved survival and mitigated disease severity as assessed by clinical score and weight [22]. ACE2-618-DDC-ABD also provided *in-vivo* lung and kidney protection and reduced lung and brain viral titers [22]. As recently demonstrated by Hassler et al. the best protective effect can be achieved when administering ACE2-618-DDC-ABD intranasally to k18-hACE2 mice both before and after viral inoculation which is superior to systemic administration at the same time-points [23]. Administration of ACE2 618-DDC-ABD only post viral inoculation provided only partial protection as compared to the untreated control group and was markedly less efficacious than administration of ACE2 618-DDC-ABD both before and after viral inoculation [23]. Specifically, intranasal administration of ACE2-618-DDC-ABD before and after viral inoculation resulted in 90% survival on day 6 post viral inoculation, brain viral titers were undetectable in all mice, and lung and brain histopathology were essentially normal [23]. In the present study that followed the same protocol of intranasal ACE2-618-DDC-ABD administration brain viral titers were similarly undetectable in the ACE2-618-DDC-ABD treated group except for the one animal that had to be humanly euthanized and considered a fatality on day six post viral inoculation. This further supports the idea that brain invasion is likely the main cause of lethality in this model.

The present study was aimed at evaluating the protective effect of ACE2-618-DDC-ABD in k18-hACE2 mice infected with the SARS-CoV-2 delta variant to establish the universality of the *in-vivo* protective effect of this ACE2 decoy protein. In previous *in-vitro* studies ACE2-618-DDC-ABD exerted a neutralizing effect on infection of Vero E6 cells by the SARS-CoV-2 wildtype, delta and gamma variants at high concentrations [10,22,23]. Infection of A549 cells with the SARS-CoV-2 omicron BA.1 variant was neutralized by 20-fold lower concentrations of ACE2 618-DDC-ABD than those required to neutralize the wildtype variant [23]. Some SARS-CoV-2 variants, like omicron BA.1, may therefore be neutralized by markedly lower concentrations of ACE2 618-ABD-DDC.

The previous *in-vitro* cell studies together with the *in-vivo* data in this study show a universal protective effect of ACE2-618-DDC-ABD against SARS-CoV-2 variants. Mutational changes during viral evolution may lead to the development of resistance to current treatment approaches [24]. Strategies that have been employed to combat SARS-CoV-2 include monoclonal antibodies and

specific antiviral drugs [53–55]. For those approaches applied in clinical practice the development of mutations that compromise treatment efficacy has been described. *In-vitro* studies have demonstrated the development of resistance towards the antiviral drugs Remdesivir and Paxlovid [56–58]. The emergence of resistant mutations has moreover been shown when *in-vitro* passing the virus in the presence of SARS-CoV-2 specific monoclonal antibodies highlighting the potential of rapidly occurring mutational changes [28,54,59]. Further studies have shown the development of resistant mutations in patients infected with the SARS-CoV-2 delta and omicron variants that received the monoclonal antibody Sotrovimab [60–62]. Mutations causing resistance against the antibody cocktails Casirivimab/Imdevimab (REGN-CoV-2) and Bamlanivimab/Etesevimab (LY-CoV555/LY-CoV016), moreover, have also been shown [63,64]. These drugs consists of two monoclonal antibodies targeting distinct epitopes [65–67], which is particularly interesting as combination therapies are typically less prone to mutational escape and decreased treatment efficiency [68–71]. The possibility of mutational escape has also been described for polyclonal antibodies, although unlike monoclonal antibodies a variety of different epitopes is being targeted [72,73].

In contrast to monoclonal antibodies, *in-vitro* passaging of SARS-CoV-2 in the presence of a soluble ACE2 decoy protein did not lead to the development of resistant mutations [28]. Importantly, saturation mutagenesis of the SARS-CoV-2 RBD followed by *in vitro* selection with full-length ACE2 and an engineered ACE2 decoy protein did not show any RBD mutations that discriminated in favor of full-length ACE2 as compared to the decoy protein [26]. Although the development of mutations that compromise the affinity for ACE2 decoy binding is theoretically possible, ACE2 decoy proteins clearly have less susceptibility to resistance caused by SARS-CoV-2 mutations [26,27]. Mutations that decrease decoy affinity would simultaneously decrease affinity for full-length membrane bound ACE2 which is essential for cell entry and infection of all currently known SARS-CoV-2 variants [8–10]. In fact, mutational escape from the neutralizing effect of ACE2 decoy proteins would therefore come at the expense of diminished viral infectivity and virulence [26,27].

During the viral evolution different mutations have been described that increase affinity of the SARS-CoV-2 RBD for ACE2 receptor binding as for example the amino acid exchanges N501Y in the SARS-CoV-2 alpha and beta variants as well as L452R in the SARS-CoV-2 delta variant [44,74,75]. The currently dominant circulating SARS-CoV-2 omicron variants also harbor various spike protein mutations leading to increased ACE2 receptor binding while efficiently evading neutralizing antibodies [76–78]. Consistent with this, as we have previously shown and noted above, concentrations of ACE2-618-DDC-ABD required to neutralize the SARS-CoV-2 omicron BA.1 variant *in-vitro* are significantly lower than those needed to neutralize the wild-type variant [23]. Of note, while this might be explained by higher binding affinity of the SARS-CoV-2 RBD to ACE2, *in-vitro* studies with SARS-CoV-2 omicron variants have also shown decreased fusion and syncytium formation which may in turn result in lower concentrations being effective for neutralization of the SARS-CoV-2 omicron variant than those needed for other SARS-CoV-2 variants [79–81].

The present study also shows the absence of kidney SARS-CoV-2 titers in k18-hACE2 mice infected with the SARS-CoV-2 delta variant. While renal involvement has been recognized as a frequent complication of COVID-19 with increased occurrence of acute kidney injury and associated higher mortality in hospitalized COVID-19 patients, the contributing molecular mechanisms remain unclear [82–85]. Direct SARS-CoV-2 infection and viral replication in the kidney parenchyma has been discussed as ACE2 is highly expressed in many cell types present in the kidney like proximal tubular cells, podocytes, mesangial and endothelial cells [86–88]. SARS-CoV-2 has been shown to directly infect human kidney organoids and the presence of SARS-CoV-2 in kidney parenchyma of patients with COVID-19 has been demonstrated by methods like immunohistochemistry (IHC), immunofluorescence (IF), real-time PCR, and single cell RNA sequencing [89]. This data, however, originates mainly from autopsies where the postmortem interval from death to autopsy ranges from several hours to days during which SARS-CoV-2 could spread to and within the kidney post-mortem [89]. Many studies have shown negative findings regarding the presence of SARS-CoV-2 in the kidney including a large series of kidney biopsy samples (n=284) of patients with COVID-19 where no direct kidney SARS-CoV-2 infection could be demonstrated [89,90]. Moreover, results from our

group in the k18-hACE2 mouse model infected with the ancestral SARS-CoV-2 variant also showed lack of evidence for kidney invasion [22,91]. Kidney tissue of k18hACE2 mice infected with the ancestral SARS-CoV-2 variant showed no detectable viral titers by plaque assay as well as no evidence for viral spike and nucleoprotein by IHC and IF staining [22,91]. Further analysis using single molecule fluorescence in situ hybridization was also negative for SARS-CoV-2 RNA [91]. The absence of kidney viral titers in k18-hACE2 mice infected with the delta SARS-CoV-2 variant supports the findings from our previous study. Considering conflicting data throughout the literature, the question whether direct kidney invasion by SARS-CoV-2 occurs remains not fully elucidated. Multiple factors, such as timing between confirmation of SARS-CoV-2 infection and obtaining of the kidney sample, the type of sample obtained (biopsy or autopsy) as well as sensitivity of the methods to detect potential presence of SARS-CoV-2 may influence the detection of SARS-CoV-2 in the kidney [89,92].

In conclusion, the present study demonstrates the efficacy of ACE2-618-DDC-ABD to protect k18-hACE2 mice from lethal infection with the aggressive SARS CoV-2 delta variant. Animals that received ACE2-618-DDC-ABD before and after viral inoculation had significantly improved survival, mitigated weight loss and improved clinical scores. Lung and brain SARS-CoV-2 titers, moreover, were significantly reduced in the ACE2 618-DDC-ABD treated group. Importantly, brain viral titers were much higher than lung viral titers and appear much more likely to be the cause of lethality in the k18-hACE2 model. Taken together with previous *in-vivo* experiments with wildtype SARS-CoV-2 and *in-vitro* data on the neutralizing effect on various variants including wildtype, gamma, delta, and omicron BA.1 [22,23], this study demonstrates the universality of the protective effect of ACE2-618-DDC-ABD against infection with SARS-CoV-2 variants. Its therapeutic and preventive potential in humans should be investigated for current and future emerging coronaviruses that use ACE2 as their main receptor for cell entry.

Author Contributions: Grant support to D Battle from NIH (1R21 AI166940-01) and a gift from the Joseph and Bessie Feinberg Foundation. L Hassler and C Cianfarini were supported by the Biomedical Education Program during a part of their stay in Chicago. A/BSL3 research management at the Howard T. Ricketts Laboratory is supported by NIAID award UC7AI180312 under the leadership of Dr. D. Missiakas. Isolate hCoV-19/USA/MD-HP05647/2021 (Lineage B.1.617.2; Delta variant), NR-55672, was obtained through BEI Resources, NIAID, NIH: SARS-Related Coronavirus 2, and contributed by Dr. Andrew S. Pekos.

Acknowledgments: Grant support to D Battle from NIH (1R21 AI166940-01) and a gift from the Joseph and Bessie Feinberg Foundation. L Hassler and C Cianfarini were supported by the Biomedical Education Program during a part of their stay in Chicago. A/BSL3 research management at the Howard T. Ricketts Laboratory is supported by NIAID award UC7AI180312 under the leadership of Dr. D. Missiakas. Isolate hCoV-19/USA/MD-HP05647/2021 (Lineage B.1.617.2; Delta variant), NR-55672, was obtained through BEI Resources, NIAID, NIH: SARS-Related Coronavirus 2, and contributed by Dr. Andrew S. Pekos.

Conflicts of Interest: D Battle and J Wysocki are coinventors of patents entitled “Active Low Molecular Weight Variants of Angiotensin Converting Enzyme 2 (ACE2),” and “Soluble ACE2 Variants and Uses therefor.” D Battle is founder of Angiotensin Therapeutics Inc. D Battle has received consulting fees from Advicenne unrelated to this work and received unrelated research support from a grant from AstraZeneca; G Randall reports consultancy agreements with Optikira. J Wysocki and J Henkin report scientific advisor capacity for Angiotensin Therapeutics Inc. All remaining authors have nothing to disclose related to this publication.

References

1. Zhou, P.; Yang, X.L.; Wang, X.G.; Hu, B.; Zhang, L.; Zhang, W.; Si, H.R.; Zhu, Y.; Li, B.; Huang, C.L.; et al. A pneumonia outbreak associated with a new coronavirus of probable bat origin. *Nature* **2020**, *579*, 270-273, doi:10.1038/s41586-020-2012-7.
2. Hu, B.; Guo, H.; Zhou, P.; Shi, Z.L. Characteristics of SARS-CoV-2 and COVID-19. *Nat Rev Microbiol* **2021**, *19*, 141-154, doi:10.1038/s41579-020-00459-7.
3. Barouch, D.H. Covid-19 Vaccines - Immunity, Variants, Boosters. *N Engl J Med* **2022**, *387*, 1011-1020, doi:10.1056/NEJMr2206573.

4. Tregoning, J.S.; Flight, K.E.; Higham, S.L.; Wang, Z.; Pierce, B.F. Progress of the COVID-19 vaccine effort: viruses, vaccines and variants versus efficacy, effectiveness and escape. *Nat Rev Immunol* **2021**, *21*, 626-636, doi:10.1038/s41577-021-00592-1.
5. Viana, R.; Moyo, S.; Amoako, D.G.; Tegally, H.; Scheepers, C.; Althaus, C.L.; Anyaneji, U.J.; Bester, P.A.; Boni, M.F.; Chand, M.; et al. Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa. *Nature* **2022**, *603*, 679-686, doi:10.1038/s41586-022-04411-y.
6. Karim, S.S.A.; Karim, Q.A. Omicron SARS-CoV-2 variant: a new chapter in the COVID-19 pandemic. *Lancet* **2021**, *398*, 2126-2128, doi:10.1016/S0140-6736(21)02758-6.
7. Tegally, H.; Moir, M.; Everatt, J.; Giovanetti, M.; Scheepers, C.; Wilkinson, E.; Subramoney, K.; Makatini, Z.; Moyo, S.; Amoako, D.G.; et al. Emergence of SARS-CoV-2 Omicron lineages BA.4 and BA.5 in South Africa. *Nat Med* **2022**, *28*, 1785-1790, doi:10.1038/s41591-022-01911-2.
8. Wan, Y.; Shang, J.; Graham, R.; Baric, R.S.; Li, F. Receptor recognition by the novel coronavirus from Wuhan: an analysis based on decade-long structural studies of SARS coronavirus. *Journal of virology* **2020**, *94*, e00127-00120.
9. Hoffmann, M.; Kleine-Weber, H.; Schroeder, S.; Kruger, N.; Herrler, T.; Erichsen, S.; Schiergens, T.S.; Herrler, G.; Wu, N.H.; Nitsche, A.; et al. SARS-CoV-2 Cell Entry Depends on ACE2 and TMPRSS2 and Is Blocked by a Clinically Proven Protease Inhibitor. *Cell* **2020**, *181*, 271-280 e278, doi:10.1016/j.cell.2020.02.052.
10. Batlle, D.; Monteil, V.; Garreta, E.; Hassler, L.; Wysocki, J.; Chandar, V.; Schwartz, R.E.; Mirazimi, A.; Montserrat, N.; Bader, M.; Penninger, J.M. Evidence in favor of the essentiality of human cell membrane-bound ACE2 and against soluble ACE2 for SARS-CoV-2 infectivity. *Cell* **2022**, *185*, 1837-1839, doi:10.1016/j.cell.2022.05.004.
11. Li, W.; Moore, M.J.; Vasilieva, N.; Sui, J.; Wong, S.K.; Berne, M.A.; Somasundaran, M.; Sullivan, J.L.; Luzuriaga, K.; Greenough, T.C.; et al. Angiotensin-converting enzyme 2 is a functional receptor for the SARS coronavirus. *Nature* **2003**, *426*, 450-454, doi:10.1038/nature02145.
12. Jackson, C.B.; Farzan, M.; Chen, B.; Choe, H. Mechanisms of SARS-CoV-2 entry into cells. *Nat Rev Mol Cell Biol* **2022**, *23*, 3-20, doi:10.1038/s41580-021-00418-x.
13. Shang, J.; Ye, G.; Shi, K.; Wan, Y.; Luo, C.; Aihara, H.; Geng, Q.; Auerbach, A.; Li, F. Structural basis of receptor recognition by SARS-CoV-2. *Nature* **2020**, *581*, 221-224, doi:10.1038/s41586-020-2179-y.
14. Lan, J.; Ge, J.; Yu, J.; Shan, S.; Zhou, H.; Fan, S.; Zhang, Q.; Shi, X.; Wang, Q.; Zhang, L.; Wang, X. Structure of the SARS-CoV-2 spike receptor-binding domain bound to the ACE2 receptor. *Nature* **2020**, *581*, 215-220, doi:10.1038/s41586-020-2180-5.
15. Walls, A.C.; Park, Y.J.; Tortorici, M.A.; Wall, A.; McGuire, A.T.; Veesler, D. Structure, Function, and Antigenicity of the SARS-CoV-2 Spike Glycoprotein. *Cell* **2020**, *181*, 281-292.e286, doi:10.1016/j.cell.2020.02.058.
16. Glowacka, I.; Bertram, S.; Muller, M.A.; Allen, P.; Soilleux, E.; Pfefferle, S.; Steffen, I.; Tsegaye, T.S.; He, Y.; Gnirss, K.; et al. Evidence that TMPRSS2 activates the severe acute respiratory syndrome coronavirus spike protein for membrane fusion and reduces viral control by the humoral immune response. *J Virol* **2011**, *85*, 4122-4134, doi:10.1128/JVI.02232-10.
17. Matsuyama, S.; Nagata, N.; Shirato, K.; Kawase, M.; Takeda, M.; Taguchi, F. Efficient activation of the severe acute respiratory syndrome coronavirus spike protein by the transmembrane protease TMPRSS2. *J Virol* **2010**, *84*, 12658-12664, doi:10.1128/JVI.01542-10.
18. Shulla, A.; Heald-Sargent, T.; Subramanya, G.; Zhao, J.; Perlman, S.; Gallagher, T. A transmembrane serine protease is linked to the severe acute respiratory syndrome coronavirus receptor and activates virus entry. *J Virol* **2011**, *85*, 873-882, doi:10.1128/JVI.02062-10.
19. Batlle, D.; Wysocki, J.; Satchell, K. Soluble angiotensin-converting enzyme 2: a potential approach for coronavirus infection therapy? *Clinical science* **2020**, *134*, 543-545.
20. Davidson, A.M.; Wysocki, J.; Batlle, D. Interaction of SARS-CoV-2 and Other Coronavirus With ACE (Angiotensin-Converting Enzyme)-2 as Their Main Receptor: Therapeutic Implications. *Hypertension* **2020**, *76*, 1339-1349, doi:10.1161/HYPERTENSIONAHA.120.15256.
21. Wysocki, J.; Ye, M.; Hassler, L.; Gupta, A.K.; Wang, Y.; Nicoleascu, V.; Randall, G.; Wertheim, J.A.; Batlle, D. A novel soluble ACE2 variant with prolonged duration of action neutralizes SARS-CoV-2 infection in human kidney organoids. *Journal of the American Society of Nephrology* **2021**, *32*, 795-803.

22. Hassler, L.; Wysocki, J.; Gelarden, I.; Sharma, I.; Tomatsidou, A.; Ye, M.; Gula, H.; Nicoleascul, V.; Randall, G.; Pshenychnyi, S.; et al. A Novel Soluble ACE2 Protein Provides Lung and Kidney Protection in Mice Susceptible to Lethal SARS-CoV-2 Infection. *J Am Soc Nephrol* **2022**, doi:10.1681/ASN.2021091209.
23. Hassler, L.; Wysocki, J.; Ahrendsen, J.T.; Ye, M.; Gelarden, I.; Nicolaescu, V.; Tomatsidou, A.; Gula, H.; Cianfarini, C.; Forster, P.; et al. Intranasal soluble ACE2 improves survival and prevents brain SARS-CoV-2 infection. *Life Sci Alliance* **2023**, 6, doi:10.26508/lsa.202301969.
24. Markov, P.V.; Ghafari, M.; Beer, M.; Lythgoe, K.; Simmonds, P.; Stilianakis, N.I.; Katzourakis, A. The evolution of SARS-CoV-2. *Nat Rev Microbiol* **2023**, 21, 361-379, doi:10.1038/s41579-023-00878-2.
25. Carabelli, A.M.; Peacock, T.P.; Thorne, L.G.; Harvey, W.T.; Hughes, J.; Consortium, C.-G.U.; Peacock, S.J.; Barclay, W.S.; de Silva, T.I.; Towers, G.J.; Robertson, D.L. SARS-CoV-2 variant biology: immune escape, transmission and fitness. *Nat Rev Microbiol* **2023**, 21, 162-177, doi:10.1038/s41579-022-00841-7.
26. Chan, K.K.; Tan, T.J.C.; Narayanan, K.K.; Procko, E. An engineered decoy receptor for SARS-CoV-2 broadly binds protein S sequence variants. *Sci Adv* **2021**, 7, doi:10.1126/sciadv.abf1738.
27. Li, G.; Qian, K.; Zhang, S.; Fu, W.; Zhao, J.; Lei, C.; Hu, S. Engineered soluble ACE2 receptor: Responding to change with change. *Front Immunol* **2022**, 13, 1084331, doi:10.3389/fimmu.2022.1084331.
28. Higuchi, Y.; Suzuki, T.; Arimori, T.; Ikemura, N.; Mihara, E.; Kiritani, Y.; Ohgitani, E.; Mazda, O.; Motooka, D.; Nakamura, S.; et al. Engineered ACE2 receptor therapy overcomes mutational escape of SARS-CoV-2. *Nat Commun* **2021**, 12, 3802, doi:10.1038/s41467-021-24013-y.
29. Hofmann, H.; Geier, M.; Marzi, A.; Krumbiegel, M.; Peipp, M.; Fey, G.H.; Gramberg, T.; Pohlmann, S. Susceptibility to SARS coronavirus S protein-driven infection correlates with expression of angiotensin converting enzyme 2 and infection can be blocked by soluble receptor. *Biochem Biophys Res Commun* **2004**, 319, 1216-1221, doi:10.1016/j.bbrc.2004.05.114.
30. Lei, C.; Qian, K.; Li, T.; Zhang, S.; Fu, W.; Ding, M.; Hu, S. Neutralization of SARS-CoV-2 spike pseudotyped virus by recombinant ACE2-Ig. *Nat Commun* **2020**, 11, 2070, doi:10.1038/s41467-020-16048-4.
31. Montell, V.; Kwon, H.; Prado, P.; Hagelkruys, A.; Wimmer, R.A.; Stahl, M.; Leopoldi, A.; Garreta, E.; Hurtado Del Pozo, C.; Prosper, F.; et al. Inhibition of SARS-CoV-2 Infections in Engineered Human Tissues Using Clinical-Grade Soluble Human ACE2. *Cell* **2020**, 181, 905-913 e907, doi:10.1016/j.cell.2020.04.004.
32. Winkler, E.S.; Bailey, A.L.; Kafai, N.M.; Nair, S.; McCune, B.T.; Yu, J.; Fox, J.M.; Chen, R.E.; Earnest, J.T.; Keeler, S.P.; et al. SARS-CoV-2 infection of human ACE2-transgenic mice causes severe lung inflammation and impaired function. *Nat Immunol* **2020**, 21, 1327-1335, doi:10.1038/s41590-020-0778-2.
33. Zheng, J.; Wong, L.R.; Li, K.; Verma, A.K.; Ortiz, M.E.; Wohlford-Lenane, C.; Leidinger, M.R.; Knudson, C.M.; Meyerholz, D.K.; McCray, P.B., Jr.; Perlman, S. COVID-19 treatments and pathogenesis including anosmia in K18-hACE2 mice. *Nature* **2021**, 589, 603-607, doi:10.1038/s41586-020-2943-z.
34. Golden, J.W.; Cline, C.R.; Zeng, X.; Garrison, A.R.; Carey, B.D.; Mucker, E.M.; White, L.E.; Shamblin, J.D.; Brocato, R.L.; Liu, J.; et al. Human angiotensin-converting enzyme 2 transgenic mice infected with SARS-CoV-2 develop severe and fatal respiratory disease. *JCI Insight* **2020**, 5, doi:10.1172/jci.insight.142032.
35. Oladunni, F.S.; Park, J.G.; Pino, P.A.; Gonzalez, O.; Akhter, A.; Allue-Guardia, A.; Olmo-Fontanez, A.; Gautam, S.; Garcia-Vilanova, A.; Ye, C.; et al. Lethality of SARS-CoV-2 infection in K18 human angiotensin-converting enzyme 2 transgenic mice. *Nat Commun* **2020**, 11, 6122, doi:10.1038/s41467-020-19891-7.
36. Moreau, G.B.; Burgess, S.L.; Sturek, J.M.; Donlan, A.N.; Petri Jr, W.A.; Mann, B.J. Evaluation of K18-hACE2 mice as a model of SARS-CoV-2 infection. *The American Journal of Tropical Medicine and Hygiene* **2020**, 103, 1215-1219.
37. Rathnasinghe, R.; Strohmeier, S.; Amanat, F.; Gillespie, V.L.; Krammer, F.; García-Sastre, A.; Coughlan, L.; Schotsaert, M.; Uccellini, M.B. Comparison of transgenic and adenovirus hACE2 mouse models for SARS-CoV-2 infection. *Emerging microbes & infections* **2020**, 9, 2433-2445.
38. Liu, X.; Mostafavi, H.; Ng, W.H.; Freitas, J.R.; King, N.J.C.; Zaid, A.; Taylor, A.; Mahalingam, S. The Delta SARS-CoV-2 Variant of Concern Induces Distinct Pathogenic Patterns of Respiratory Disease in K18-hACE2 Transgenic Mice Compared to the Ancestral Strain from Wuhan. *mBio* **2022**, 13, e0068322, doi:10.1128/mbio.00683-22.
39. Lee, K.S.; Wong, T.Y.; Russ, B.P.; Horspool, A.M.; Miller, O.A.; Rader, N.A.; Givi, J.P.; Winters, M.T.; Wong, Z.Y.A.; Cyphert, H.A.; et al. SARS-CoV-2 Delta variant induces enhanced pathology and inflammatory responses in K18-hACE2 mice. *PLoS One* **2022**, 17, e0273430, doi:10.1371/journal.pone.0273430.
40. Singh, J.; Rahman, S.A.; Ehtesham, N.Z.; Hira, S.; Hasnain, S.E. SARS-CoV-2 variants of concern are emerging in India. *Nat Med* **2021**, 27, 1131-1133, doi:10.1038/s41591-021-01397-4.

41. McCrone, J.T.; Hill, V.; Bajaj, S.; Pena, R.E.; Lambert, B.C.; Inward, R.; Bhatt, S.; Volz, E.; Ruis, C.; Dellicour, S.; et al. Context-specific emergence and growth of the SARS-CoV-2 Delta variant. *Nature* **2022**, *610*, 154-160, doi:10.1038/s41586-022-05200-3.
42. Fisman, D.N.; Tuite, A.R. Evaluation of the relative virulence of novel SARS-CoV-2 variants: a retrospective cohort study in Ontario, Canada. *CMAJ* **2021**, *193*, E1619-E1625, doi:10.1503/cmaj.211248.
43. Sheikh, A.; McMennamin, J.; Taylor, B.; Robertson, C.; Public Health, S.; the, E.I.I.C. SARS-CoV-2 Delta VOC in Scotland: demographics, risk of hospital admission, and vaccine effectiveness. *Lancet* **2021**, *397*, 2461-2462, doi:10.1016/S0140-6736(21)01358-1.
44. Motozono, C.; Toyoda, M.; Zahradnik, J.; Saito, A.; Nasser, H.; Tan, T.S.; Ngare, I.; Kimura, I.; Uriu, K.; Kosugi, Y.; et al. SARS-CoV-2 spike L452R variant evades cellular immunity and increases infectivity. *Cell Host Microbe* **2021**, *29*, 1124-1136 e1111, doi:10.1016/j.chom.2021.06.006.
45. Liu, Z.; VanBlargan, L.A.; Bloyet, L.M.; Rothlauf, P.W.; Chen, R.E.; Stumpf, S.; Zhao, H.; Errico, J.M.; Theel, E.S.; Liebeskind, M.J.; et al. Identification of SARS-CoV-2 spike mutations that attenuate monoclonal and serum antibody neutralization. *Cell Host Microbe* **2021**, *29*, 477-488 e474, doi:10.1016/j.chom.2021.01.014.
46. Li, Q.; Wu, J.; Nie, J.; Zhang, L.; Hao, H.; Liu, S.; Zhao, C.; Zhang, Q.; Liu, H.; Nie, L.; et al. The Impact of Mutations in SARS-CoV-2 Spike on Viral Infectivity and Antigenicity. *Cell* **2020**, *182*, 1284-1294 e1289, doi:10.1016/j.cell.2020.07.012.
47. Callaway, E. The mutation that helps Delta spread like wildfire. *Nature* **2021**, *596*, 472-473, doi:10.1038/d41586-021-02275-2.
48. Di Giacomo, S.; Mercatelli, D.; Rakhimov, A.; Giorgi, F.M. Preliminary report on severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) Spike mutation T478K. *J Med Virol* **2021**, *93*, 5638-5643, doi:10.1002/jmv.27062.
49. Hou, Y.J.; Chiba, S.; Halfmann, P.; Ehre, C.; Kuroda, M.; Dinnon, K.H., 3rd; Leist, S.R.; Schafer, A.; Nakajima, N.; Takahashi, K.; et al. SARS-CoV-2 D614G variant exhibits efficient replication ex vivo and transmission in vivo. *Science* **2020**, *370*, 1464-1468, doi:10.1126/science.abe8499.
50. Korber, B.; Fischer, W.M.; Gnanakaran, S.; Yoon, H.; Theiler, J.; Abfalterer, W.; Hengartner, N.; Giorgi, E.E.; Bhattacharya, T.; Foley, B.; et al. Tracking Changes in SARS-CoV-2 Spike: Evidence that D614G Increases Infectivity of the COVID-19 Virus. *Cell* **2020**, *182*, 812-827 e819, doi:10.1016/j.cell.2020.06.043.
51. Gobeil, S.M.; Janowska, K.; McDowell, S.; Mansouri, K.; Parks, R.; Manne, K.; Stalls, V.; Kopp, M.F.; Henderson, R.; Edwards, R.J.; et al. D614G Mutation Alters SARS-CoV-2 Spike Conformation and Enhances Protease Cleavage at the S1/S2 Junction. *Cell Rep* **2021**, *34*, 108630, doi:10.1016/j.celrep.2020.108630.
52. Plante, J.A.; Liu, Y.; Liu, J.; Xia, H.; Johnson, B.A.; Lokugamage, K.G.; Zhang, X.; Muruato, A.E.; Zou, J.; Fontes-Garfias, C.R.; et al. Spike mutation D614G alters SARS-CoV-2 fitness. *Nature* **2021**, *592*, 116-121, doi:10.1038/s41586-020-2895-3.
53. Murakami, N.; Hayden, R.; Hills, T.; Al-Samkari, H.; Casey, J.; Del Sorbo, L.; Lawler, P.R.; Sise, M.E.; Leaf, D.E. Therapeutic advances in COVID-19. *Nat Rev Nephrol* **2023**, *19*, 38-52, doi:10.1038/s41581-022-00642-4.
54. Focosi, D.; McConnell, S.; Casadevall, A.; Cappello, E.; Valdiserra, G.; Tuccori, M. Monoclonal antibody therapies against SARS-CoV-2. *Lancet Infect Dis* **2022**, *22*, e311-e326, doi:10.1016/S1473-3099(22)00311-5.
55. Tao, K.; Tzou, P.L.; Nouhin, J.; Bonilla, H.; Jagannathan, P.; Shafer, R.W. SARS-CoV-2 Antiviral Therapy. *Clin Microbiol Rev* **2021**, *34*, e0010921, doi:10.1128/CMR.00109-21.
56. Szemiel, A.M.; Merits, A.; Orton, R.J.; MacLean, O.A.; Pinto, R.M.; Wickenhagen, A.; Lieber, G.; Turnbull, M.L.; Wang, S.; Furnon, W.; et al. In vitro selection of Remdesivir resistance suggests evolutionary predictability of SARS-CoV-2. *PLoS Pathog* **2021**, *17*, e1009929, doi:10.1371/journal.ppat.1009929.
57. Stevens, L.J.; Pruijssers, A.J.; Lee, H.W.; Gordon, C.J.; Tchesnokov, E.P.; Gribble, J.; George, A.S.; Hughes, T.M.; Lu, X.; Li, J.; et al. Mutations in the SARS-CoV-2 RNA-dependent RNA polymerase confer resistance to remdesivir by distinct mechanisms. *Sci Transl Med* **2022**, *14*, eabo0718, doi:10.1126/scitranslmed.abo0718.
58. Zhou, Y.; Gammelfoft, K.A.; Ryberg, L.A.; Pham, L.V.; Tjornelund, H.D.; Binderup, A.; Duarte Hernandez, C.R.; Fernandez-Antunez, C.; Offersgaard, A.; Fahnoe, U.; et al. Nirmatrelvir-resistant SARS-CoV-2 variants with high fitness in an infectious cell culture system. *Sci Adv* **2022**, *8*, eadd7197, doi:10.1126/sciadv.add7197.
59. Hoffmann, M.; Arora, P.; Gross, R.; Seidel, A.; Hornich, B.F.; Hahn, A.S.; Kruger, N.; Graichen, L.; Hofmann-Winkler, H.; Kempf, A.; et al. SARS-CoV-2 variants B.1.351 and P.1 escape from neutralizing antibodies. *Cell* **2021**, *184*, 2384-2393 e2312, doi:10.1016/j.cell.2021.03.036.

60. Birnie, E.; Biemond, J.J.; Appelman, B.; de Bree, G.J.; Jonges, M.; Welkers, M.R.A.; Wiersinga, W.J. Development of Resistance-Associated Mutations After Sotrovimab Administration in High-risk Individuals Infected With the SARS-CoV-2 Omicron Variant. *JAMA* **2022**, *328*, 1104-1107, doi:10.1001/jama.2022.13854.
61. Rockett, R.; Basile, K.; Maddocks, S.; Fong, W.; Agius, J.E.; Johnson-Mackinnon, J.; Arnott, A.; Chandra, S.; Gall, M.; Draper, J.; et al. Resistance Mutations in SARS-CoV-2 Delta Variant after Sotrovimab Use. *N Engl J Med* **2022**, *386*, 1477-1479, doi:10.1056/NEJMc2120219.
62. Vellas, C.; Tremieux, P.; Del Bello, A.; Latour, J.; Jeanne, N.; Ranger, N.; Danet, C.; Martin-Blondel, G.; Delobel, P.; Kamar, N.; Izopet, J. Resistance mutations in SARS-CoV-2 omicron variant in patients treated with sotrovimab. *Clin Microbiol Infect* **2022**, *28*, 1297-1299, doi:10.1016/j.cmi.2022.05.002.
63. Starr, T.N.; Greaney, A.J.; Addetia, A.; Hannon, W.W.; Choudhary, M.C.; Dings, A.S.; Li, J.Z.; Bloom, J.D. Prospective mapping of viral mutations that escape antibodies used to treat COVID-19. *Science* **2021**, *371*, 850-854, doi:10.1126/science.abf9302.
64. Starr, T.N.; Greaney, A.J.; Dings, A.S.; Bloom, J.D. Complete map of SARS-CoV-2 RBD mutations that escape the monoclonal antibody LY-CoV555 and its cocktail with LY-CoV016. *Cell Rep Med* **2021**, *2*, 100255, doi:10.1016/j.xcrm.2021.100255.
65. Baum, A.; Fulton, B.O.; Wloga, E.; Copin, R.; Pascal, K.E.; Russo, V.; Giordano, S.; Lanza, K.; Negron, N.; Ni, M.; et al. Antibody cocktail to SARS-CoV-2 spike protein prevents rapid mutational escape seen with individual antibodies. *Science* **2020**, *369*, 1014-1018, doi:10.1126/science.abd0831.
66. Hansen, J.; Baum, A.; Pascal, K.E.; Russo, V.; Giordano, S.; Wloga, E.; Fulton, B.O.; Yan, Y.; Koon, K.; Patel, K.; et al. Studies in humanized mice and convalescent humans yield a SARS-CoV-2 antibody cocktail. *Science* **2020**, *369*, 1010-1014, doi:10.1126/science.abd0827.
67. Gottlieb, R.L.; Nirula, A.; Chen, P.; Boscia, J.; Heller, B.; Morris, J.; Huhn, G.; Cardona, J.; Mocherla, B.; Stosor, V.; et al. Effect of Bamlanivimab as Monotherapy or in Combination With Etesevimab on Viral Load in Patients With Mild to Moderate COVID-19: A Randomized Clinical Trial. *JAMA* **2021**, *325*, 632-644, doi:10.1001/jama.2021.0202.
68. Hammond, J.; Leister-Tebbe, H.; Gardner, A.; Abreu, P.; Bao, W.; Wisemandle, W.; Baniecki, M.; Hendrick, V.M.; Damle, B.; Simon-Campos, A.; et al. Oral Nirmatrelvir for High-Risk, Nonhospitalized Adults with Covid-19. *N Engl J Med* **2022**, *386*, 1397-1408, doi:10.1056/NEJMoa2118542.
69. Jeong, J.H.; Chokkakula, S.; Min, S.C.; Kim, B.K.; Choi, W.S.; Oh, S.; Yun, Y.S.; Kang, D.H.; Lee, O.J.; Kim, E.G.; et al. Combination therapy with nirmatrelvir and molnupiravir improves the survival of SARS-CoV-2 infected mice. *Antiviral Res* **2022**, *208*, 105430, doi:10.1016/j.antiviral.2022.105430.
70. Clavel, F.; Hance, A.J. HIV drug resistance. *N Engl J Med* **2004**, *350*, 1023-1035, doi:10.1056/NEJMra025195.
71. Dunning, J.; Baillie, J.K.; Cao, B.; Hayden, F.G.; International Severe Acute, R.; Emerging Infection, C. Antiviral combinations for severe influenza. *Lancet Infect Dis* **2014**, *14*, 1259-1270, doi:10.1016/S1473-3099(14)70821-7.
72. Greaney, A.J.; Starr, T.N.; Barnes, C.O.; Weisblum, Y.; Schmidt, F.; Caskey, M.; Gaebler, C.; Cho, A.; Agudelo, M.; Finkin, S.; et al. Mapping mutations to the SARS-CoV-2 RBD that escape binding by different classes of antibodies. *Nat Commun* **2021**, *12*, 4196, doi:10.1038/s41467-021-24435-8.
73. Chen, R.E.; Zhang, X.; Case, J.B.; Winkler, E.S.; Liu, Y.; VanBlargan, L.A.; Liu, J.; Errico, J.M.; Xie, X.; Suryadevara, N.; et al. Resistance of SARS-CoV-2 variants to neutralization by monoclonal and serum-derived polyclonal antibodies. *Nat Med* **2021**, *27*, 717-726, doi:10.1038/s41591-021-01294-w.
74. Starr, T.N.; Greaney, A.J.; Hilton, S.K.; Ellis, D.; Crawford, K.H.D.; Dings, A.S.; Navarro, M.J.; Bowen, J.E.; Tortorici, M.A.; Walls, A.C.; et al. Deep Mutational Scanning of SARS-CoV-2 Receptor Binding Domain Reveals Constraints on Folding and ACE2 Binding. *Cell* **2020**, *182*, 1295-1310 e1220, doi:10.1016/j.cell.2020.08.012.
75. Zahradnik, J.; Marciano, S.; Shemesh, M.; Zoler, E.; Harari, D.; Chiaravalli, J.; Meyer, B.; Rudich, Y.; Li, C.; Marton, I.; et al. SARS-CoV-2 variant prediction and antiviral drug design are enabled by RBD in vitro evolution. *Nat Microbiol* **2021**, *6*, 1188-1198, doi:10.1038/s41564-021-00954-4.
76. Mannar, D.; Saville, J.W.; Zhu, X.; Srivastava, S.S.; Berezuk, A.M.; Tuttle, K.S.; Marquez, A.C.; Sekirov, I.; Subramaniam, S. SARS-CoV-2 Omicron variant: Antibody evasion and cryo-EM structure of spike protein-ACE2 complex. *Science* **2022**, *375*, 760-764, doi:10.1126/science.abn7760.
77. Hoffmann, M.; Zhang, L.; Pohlmann, S. Omicron: Master of immune evasion maintains robust ACE2 binding. *Signal Transduct Target Ther* **2022**, *7*, 118, doi:10.1038/s41392-022-00965-5.

78. Meng, B.; Abdullahi, A.; Ferreira, I.; Goonawardane, N.; Saito, A.; Kimura, I.; Yamasoba, D.; Gerber, P.P.; Fatihi, S.; Rathore, S.; et al. Altered TMPRSS2 usage by SARS-CoV-2 Omicron impacts infectivity and fusogenicity. *Nature* **2022**, *603*, 706-714, doi:10.1038/s41586-022-04474-x.
79. Meng, B.; Abdullahi, A.; Ferreira, I.A.; Goonawardane, N.; Saito, A.; Kimura, I.; Yamasoba, D.; Gerber, P.P.; Fatihi, S.; Rathore, S. Altered TMPRSS2 usage by SARS-CoV-2 Omicron impacts infectivity and fusogenicity. *Nature* **2022**, *603*, 706-714.
80. Suzuki, R.; Yamasoba, D.; Kimura, I.; Wang, L.; Kishimoto, M.; Ito, J.; Morioka, Y.; Nao, N.; Nasser, H.; Uriu, K. Attenuated fusogenicity and pathogenicity of SARS-CoV-2 Omicron variant. *Nature* **2022**, *603*, 700-705.
81. Sun, C.; Wang, H.; Yang, J.; Kong, D.; Chen, Y.; Wang, H.; Sun, L.; Lu, J.; Teng, M.; Xie, L. Mutation N856K in spike reduces fusogenicity and infectivity of Omicron BA. 1. *Signal Transduction and Targeted Therapy* **2023**, *8*, 75.
82. Battle, D.; Soler, M.J.; Sparks, M.A.; Hiremath, S.; South, A.M.; Welling, P.A.; Swaminathan, S.; Covid; Ace2 in Cardiovascular, L.; Kidney Working, G. Acute Kidney Injury in COVID-19: Emerging Evidence of a Distinct Pathophysiology. *J Am Soc Nephrol* **2020**, *31*, 1380-1383, doi:10.1681/ASN.2020040419.
83. Kunutsor, S.K.; Laukkanen, J.A. Renal complications in COVID-19: a systematic review and meta-analysis. *Ann Med* **2020**, *52*, 345-353, doi:10.1080/07853890.2020.1790643.
84. Hirsch, J.S.; Ng, J.H.; Ross, D.W.; Sharma, P.; Shah, H.H.; Barnett, R.L.; Hazzan, A.D.; Fishbane, S.; Jhaveri, K.D.; Northwell, C.-R.C.; Northwell Nephrology, C.-R.C. Acute kidney injury in patients hospitalized with COVID-19. *Kidney Int* **2020**, *98*, 209-218, doi:10.1016/j.kint.2020.05.006.
85. Cheng, Y.; Luo, R.; Wang, K.; Zhang, M.; Wang, Z.; Dong, L.; Li, J.; Yao, Y.; Ge, S.; Xu, G. Kidney disease is associated with in-hospital death of patients with COVID-19. *Kidney Int* **2020**, *97*, 829-838, doi:10.1016/j.kint.2020.03.005.
86. Hikmet, F.; Mear, L.; Edvinsson, A.; Micke, P.; Uhlen, M.; Lindskog, C. The protein expression profile of ACE2 in human tissues. *Mol Syst Biol* **2020**, *16*, e9610, doi:10.15252/msb.20209610.
87. Ye, M.; Wysocki, J.; William, J.; Soler, M.J.; Cokic, I.; Battle, D. Glomerular localization and expression of Angiotensin-converting enzyme 2 and Angiotensin-converting enzyme: implications for albuminuria in diabetes. *J Am Soc Nephrol* **2006**, *17*, 3067-3075, doi:10.1681/asn.2006050423.
88. Hamming, I.; Timens, W.; Bulthuis, M.L.; Lely, A.T.; Navis, G.; van Goor, H. Tissue distribution of ACE2 protein, the functional receptor for SARS coronavirus. A first step in understanding SARS pathogenesis. *J Pathol* **2004**, *203*, 631-637, doi:10.1002/path.1570.
89. Hassler, L.; Reyes, F.; Sparks, M.A.; Welling, P.; Battle, D. Evidence For and Against Direct Kidney Infection by SARS-CoV-2 in Patients with COVID-19. *Clin J Am Soc Nephrol* **2021**, *16*, 1755-1765, doi:10.2215/cjn.04560421.
90. May, R.M.; Cassol, C.; Hannoudi, A.; Larsen, C.P.; Lerma, E.V.; Haun, R.S.; Braga, J.R.; Hassen, S.I.; Wilson, J.; VanBeek, C.; et al. A multi-center retrospective cohort study defines the spectrum of kidney pathology in Coronavirus 2019 Disease (COVID-19). *Kidney Int* **2021**, *100*, 1303-1315, doi:10.1016/j.kint.2021.07.015.
91. Syed, M.; Cianfarini, C.; Hassler, L.; Ye, M.; Wysocki, J.; Kanwar, Y.; Missiakas, D.; Randall, G.; Battle, D. Lack of Evidence for Kidney Invasion by SARS-CoV-2 in a Lethal Mouse Model. *J Am Soc Nephrol* **2022**, *331*.
92. Hassler, L.; Battle, D. Potential SARS-CoV-2 kidney infection and paths to injury. *Nat Rev Nephrol* **2022**, *18*, 275-276, doi:10.1038/s41581-022-00551-6.

Disclaimer/Publisher's Note: The statements, opinions and data contained in all publications are solely those of the individual author(s) and contributor(s) and not of MDPI and/or the editor(s). MDPI and/or the editor(s) disclaim responsibility for any injury to people or property resulting from any ideas, methods, instructions or products referred to in the content.