## **RESEARCH ARTICLE**

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# A single intranasal administration of AdCOVID protects against SARS-CoV-2 infection in the upper and lower respiratory tracts

Michael D. Schultz (1)<sup>a\*</sup>, John J. Suschak (1)<sup>b\*</sup>, Davide Botta (1)<sup>a</sup>, Aaron Silva-Sanchez (1)<sup>c</sup>, R. Glenn King (1)<sup>a</sup>, Thomas W. Detchemendy<sup>d</sup>, Chetan D. Meshram (1)<sup>e</sup>, Jeremy B. Foote (1)<sup>a</sup>, Fen Zhou<sup>a</sup>, Jennifer L. Tipper<sup>e</sup>, Jianfeng Zhang<sup>b</sup>, Kevin S. Harrod (1)<sup>e</sup>, Sixto M. Leal Jr<sup>d</sup>, Troy D. Randall (1)<sup>c</sup>, M. Scot Roberts<sup>b</sup>, Bertrand Georges<sup>b</sup>, and Frances E. Lund (1)<sup>a</sup>

<sup>a</sup>Department of Microbiology, The University of Alabama at Birmingham, Birmingham, AL, USA; <sup>b</sup>Altimmune Inc., Gaithersburg, MD, USA; <sup>c</sup>Department of Medicine, Division of Clinical Immunology and Rheumatology, The University of Alabama at Birmingham, Birmingham, AL, USA; <sup>d</sup>Department of Pathology, Division of Laboratory Medicine, The University of Alabama at Birmingham, Birmingham, AL, USA; <sup>e</sup>Department of Anesthesiology and Perioperative Medicine, The University of Alabama at Birmingham, AL, USA; <sup>e</sup>Department of Anesthesiology and Perioperative Medicine, The University of Alabama at Birmingham, AL, USA; <sup>e</sup>Department of Anesthesiology and Perioperative Medicine, The University of Alabama at Birmingham, AL, USA

#### ABSTRACT

The coronavirus disease 2019 (COVID-19) pandemic has illustrated the critical need for effective prophylactic vaccination to prevent the spread of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). Intranasal vaccination is an attractive approach for preventing COVID-19 as the nasal mucosa is the site of initial SARS-CoV-2 entry and viral replication prior to aspiration into the lungs. We previously demonstrated that a single intranasal administration of a candidate adenovirus type 5-vectored vaccine encoding the receptor-binding domain of the SARS-CoV-2 spike protein (AdCOVID) induced robust immunity in both the airway mucosa and periphery, and completely protected K18-hACE2 mice from lethal SARS-CoV -2 challenge. Here we show that a single intranasal administration of AdCOVID limits viral replication in the nasal cavity of K18-hACE2 mice. AdCOVID also induces sterilizing immunity in the lungs of mice as reflected by the absence of infectious virus. Finally, AdCOVID prevents SARS-CoV-2 induced pathological damage in the lungs of mice. These data show that AdCOVID not only limits viral replication in the respiratory tract, but it also prevents virus-induced inflammation and immunopathology following SARS-CoV-2 infection.

#### Introduction

Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) emerged in late 2019 and has resulted in a global pandemic of an acute respiratory disease, named "coronavirus disease 2019" (COVID-19), which threatens human health and public safety. The infective dose for SARS-CoV-2 remains undetermined, but it is known that airborne transmission occurs via inhalation of respiratory droplets expelled from infected hosts.<sup>1,2</sup> SARS-CoV-2 entry into host cells is dependent on engagement of the receptor-binding domain (RBD) of the spike protein to the viral entry receptor angiotensinconverting enzyme-2 (ACE2), initiating fusion of the virus with the cellular membrane.<sup>3</sup> ACE2 is expressed in the nasal goblet and ciliated epithelial cells lining the respiratory tract and the lungs, and it is hypothesized that the nasal cavity serves as the initial reservoir for seeding of SARS-CoV-2 to the lungs.<sup>4,5</sup> High viral load in the nasopharyngeal space directly correlates with SARS-CoV-2 transmission and mortality.<sup>6-10</sup> The lung is the principle organ affected in COVID-19 patients with hypoxic respiratory failure being the primary cause of COVID-19 mortality.<sup>11</sup> The lungs of COVID-19 patients exhibit distinct histological characteristics including acute endothelial injury, pervasive capillary

inflammation, fibrosis, thrombosis, and apoptotic or pyroptotic cell death.<sup>12,13</sup> These data suggest that the optimal anti-SARS-CoV-2 immune response for preventing disease will contain a mucosal immunity component.

The urgent need for safe and effective SARS-CoV-2 countermeasures has resulted in eleven vaccines being marketed worldwide, with multiple vaccines being granted authorization by the U.S. Food and Drug Administration (USFDA) for adults, adolescents, and children.<sup>14-17</sup> The vast majority of SARS-CoV-2 vaccines are either inactivated whole virus vaccines or encode the full-length spike protein (S) as the antigen, and all are delivered by intramuscular (IM) injection.<sup>18</sup> IM injection elicits systemic immunity but cannot yield potent mucosal immunity.<sup>19,20</sup> Alternatively, a noninvasive administration route such as intranasal delivery may be attractive.<sup>21</sup> Data suggest intranasal vaccination, acting in a manner analogous to viral infection, has the potential to confer sterilizing immunity in the respiratory tract, reducing the morbidity and transmission of COVID-19.22-27 By removing the need for needles, intranasal delivery may also increase vaccine uptake, presenting a more efficient vaccine administration method.<sup>28</sup>

We recently reported that a replication-deficient adenovirus serotype 5 (Ad5)-vectored vaccine candidate encoding only the SARS-CoV-2 RBD was highly immunogenic in mice when

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**CONTACT** Frances E. Lund R flund@uab.edu Department of Microbiology, The University of Alabama at Birmingham, Birmingham, AL, USA. \*These authors contributed equally to this work.

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delivered by the intranasal route.<sup>29</sup> RBD-only vaccines may have multiple advantages over trimeric spike-encoding vaccines. Immunofocusing of the antibody repertoire may yield higher affinity antibodies and neutralizing antibodies directed against conserved subdominant epitopes, allowing for potent cross-variant neutralization.<sup>30–33</sup> Furthermore, the majority of neutralizing antibodies in human convalescent sera are directed against the RBD, highlighting the importance of anti-RBD responses in protective immunity.<sup>34–36</sup> Our vaccine, referred to as AdCOVID, elicited systemic neutralizing antibody responses in the sera of animals that persisted for at least six months.<sup>29</sup> AdCOVID administration also resulted in a robust population of anti-RBD T cells and anti-spike IgA antibodies in the lungs of all vaccinated mice. Importantly, AdCOVID was completely protective against lethal SARS-CoV-2 challenge in K18-hACE2 mice, with no animals exhibiting overt morbidity or mortality.

Here, we further investigated the protective efficacy of AdCOVID in K18-hACE2 mice. We report that a single intranasal administration of AdCOVID significantly decreased the copies of viral RNA in SARS-CoV-2 challenged mice compared to vehicle controls. AdCOVID vaccination also prevented the seeding of infectious virus in the lungs of challenged mice. Finally, AdCOVID vaccination significantly reduced SARS-CoV-2-induced immunopathology in challenged mice.

#### Methods

#### Ethics statement and mice

K18-hACE2 breeding pairs were originally purchased from Jackson Laboratory (Strain #034860) and subsequent generations were bred at the University of Alabama at Birmingham (UAB). Animal procedures performed at UAB were conducted in accordance with Public Health Service Policy on the Humane Care and Use of Laboratory Animals and Guide for the Care and Use of Laboratory Animals. Studies were performed in the Southeastern Biosafety Lab (SEBLAB) ABSL3 located at UAB. Studies were performed under UAB's Institutional Animal Care and Use Committee (IACUC: D16-00162 A3255-01) Protocol 22065 approved on 27 April 2020.

### Vaccine candidate

The vaccine candidate evaluated in this study was based on a replication-deficient, E1- and E3-deleted Ad5 vector platform and expresses a human codon-optimized gene for the RBD domain (residues 302 to 543) of SARS-CoV-2 spike protein (accession number QHD43416.1).<sup>37</sup> The Ad5-vectored RBD transgene includes a human tissue plasminogen activator leader sequence and is expressed under the control of the cytomegalovirus immediate early promoter/enhancer.<sup>29</sup> An initial seed stock was obtained from transfection of recombinant vector plasmid into E1-complementing PER.C6 cells as previously described.<sup>29</sup> Vaccine titers were measured in HEK 293 cells as previously described.<sup>29</sup>

# **Propagation of SARS-CoV-2**

The original SARS-CoV-2 isolate USA-WA1/2020 was obtained from BEI resources (#NR-52281) and propagated in Vero E6 cells (ATCC, #CRL-1586). Confluent cells in a 25 CM<sup>2</sup> flask were inoculated with virus (0.015 MOI) in serum free Eagle's MEM (Life Technologies, #11430–030) with added supplements, and incubated with rocking at 35°C (5% CO<sub>2</sub>) for 2 hours. The inoculum was then removed, cells were overlaid with 2% FBS in media, and virus growth continued until obvious cytopathic effect (CPE) was observed in 2 to 3 days post infection. A large-scale working stock was obtained through 2–3 more passages of virus in cells in 150 CM<sup>2</sup> flasks until significant viral titer was achieved.

#### K18-hACE2 vaccination and challenge

Mixed sex K18-hACE2 mice of at least 6 weeks of age were randomly allocated into vaccination groups for immunogenicity experiments. 2.2E + 09 infectious units (ifu) of replicationdeficient Ad5 vector encoding the RBD (AdCOVID) from the SARS-CoV-2 spike protein were administered intranasally at the described doses in a volume of 50  $\mu$ L (25  $\mu$ L/nostril). Briefly, mice were sedated with isoflurane and placed in the supine position. AdCOVID was delivered into the nasal mucosa by pipette during normal inhalation, and mice were allowed to recover naturally. The control group received 50  $\mu$ L of vehicle alone by intranasal administration. On study day 32, all mice were challenged by the intranasal route with LD<sub>50</sub> doses of either 2.0E + 04 or 0.5E + 04 plaque forming units (PFU) SARS-CoV-2 strain USA/WA-1/2020. Mice were euthanized either 3- or 4-days post-infection (DPI) as indicated.

#### Serum collection

Blood samples were collected from the submandibular vein of vaccinated mice into BD Microtainer blood collection tubes (BD Biosciences, #365963). The samples were centrifuged at 13,000 rpm at RT for 10 minutes and the serum was collected, aliquoted and frozen at  $-80^{\circ}$ C until analyzed.

#### **Recombinant SARS-CoV-2 protein production**

To produce recombinant SARS-CoV-2 spike ectodomain protein, two human codon-optimized constructs were generated with linear sequence order encoding: a human IgG leader sequence, the SARS-CoV-2 spike ectodomain (amino acids 14–1211), a GGSG linker, T4 fibritin foldon sequence, a GS linker, and finally an AviTag (construct 1) or 6X-HisTag (construct 2). Each construct was engineered with two sets of mutations to stabilize the protein in a pre-fusion conformation as previously described.<sup>29</sup> IgG standards were generated as previously described.<sup>29</sup>

# SARS-CoV-2 spike cytometric bead array

To generate the spike cytometric bead array (CBA), recombinant SARS-CoV-2 ectodomain trimers were passively absorbed onto streptavidin functionalized fluorescent microparticles (3.6 µm, Spherotech, # CPAK-3567-4K). 500 µg of biotinylated SARS2-CoV-2 was incubated with 2E + 07 streptavidin functionalized fluorescent microparticles in 400 µL of 1% BSA in PBS. Following coupling, the SARS-CoV-2 spike conjugated beads were washed twice in 1 mL of 1% BSA, PBS, 0.05% NaN3, resuspended at 1E + 08 beads/mL and stored at 4°C. The loading of recombinant SARS2-CoV-2 spike onto the beads was evaluated by staining 1E + 05 beads with dilutions ranging from 1 µg/mL to 2 ng/mL of the recombinant anti-SARS spike antibody CR3022 (IDT Biologika) and visualized with an anti-human IgG secondary antibody.

# CBA measurement of spike-specific IgG responses

The quantification of SARS-CoV-2 spike IgG was performed in serum samples obtained from all animals using the spike CBA described above.  $5 \,\mu\text{L}$  of a suspension containing 5E + 05 of each SARS-CoV-2 spike and anti-IgG beads was added to the diluted samples and CBA measurement was conducted as previously described.<sup>29</sup>

#### Nasal swipe collection

Nasal swipes were collected immediately prior to intranasal challenge to establish baseline viral load, and then 1-hour postinfection. Nasal swipes were then collected daily through study termination (DPI +4). Awake mice were scruffed and the exterior surface of their nose was swiped for 5–10 seconds with a polyester-tipped swab (ThermoFisher, #22-029-574) that had been dipped in a screw-cap tube containing 300  $\mu$ L Viral Transport Medium (VTM; HBSS (+Ca<sup>2+</sup> +Mg<sup>2+</sup>) containing 2% FBS, 100  $\mu$ g/mL Gentamicin and 0.5  $\mu$ g/mL AmphotericinB). The tip of the swab was cut and dropped inside the VTM-containing tube. Once all swab samples were collected, the tubes were capped and vortexed, and then the swab tips were removed and discarded.

### Nasal wash collection

The upper respiratory tract was washed postmortem with 400  $\mu$ L VTM instilled into the trachea with a blunt 19-gauge needle and flushed out of the nose. 200–300  $\mu$ L of the nasal wash collected was used for quantification of viral RNA by qRT-PCR. The left-over sample volumes were used for antibody titer quantification.

#### Tissue processing and single cell isolation

The lungs were excised at euthanasia (without pre-collection of the bronchoalveolar lavage (BAL) fluid) and placed in a 2 mL Lysing Matrix M tube (MP Biomedicals, #116923050-CF) containing 1 mL VTM. The lungs were homogenized using a FastPrep-24 classic bead beating grinder lysis system (MP Biomedicals, #116004500). The total lung homogenate was centrifuged at 10,000 rpm for 10 minutes at room temperature.  $300 \,\mu$ L of the homogenate supernatant was used for quantification of viral load by qRT-PCR. The remaining homogenate supernatants were used for quantification of viral load by plaque assay. Brains were excised and placed in a 2-mL Lysing Matrix M tube containing  $800 \mu$ L VTM and homogenized using a FastPrep-24 classic bead beating grinder lysis system. The homogenate was centrifuged at 10,000 rpm for 10 minutes at room temperature.  $300 \mu$ L of the homogenate supernatant was used for quantification of viral load by qRT-PCR. The remaining homogenate supernatants were used for quantification of viral load by plaque assay.

# qRT-PCR

The quantification of SARS-CoV-2 viral RNA was performed from nasal swipe (300 µL), nasal wash (300 µL), total lung (tissue + BAL) homogenate (300 µL), and brain homogenates (300 µL) isolated from individual mice. Each sample was mixed 1:1 with lysis buffer containing 10% proteinase K from the Maxwell RSC Viral Total Nucleic Acid Purification kit (Promega, #AS1330). The samples were mixed and incubated at 56°C for 10 minutes. RNA was extracted utilizing the Maxwell RSC 48 Instrument (Promega, Madison, WI) and qRT-PCR performed utilizing QuantStudio<sup>™</sup> 5 Real-Time PCR instrument (ThermoFisher, Waltham, MA). The genomic RNA PCR primers (forward 5'-GAC CCC AAA ATC AGC GAA AT-3' and reverse 5'-TCT GGT TAC TAC TGC CAG TTG AAT CTG-3') and probe (/SFAM/ACC CCG CAT TAC GTT TGG TGG ACC/BHQ\_1) targeted the nucleocapsid (N) gene and were purchased from Integrated DNA Technologies, Inc. (IDT, Coralville, Iowa). AccuPlex<sup>™</sup> SARS-CoV-2 Reference Material (SeraCare Life Sciences, Inc, #0505-0126) was extracted and amplified in parallel to generate a standard curve enabling viral quantitation. The lowest limit of quantification for this assay is 25 viral RNA copies/mL. The subgenomic RNA PCR assay utilized a forward primer targeting the leader sequence (5'-CGA TCT CTT GTA GAT CTG TTC TC-3') and the same reverse primer targeting the N gene with RT-PCR analysis utilizing the Power SYBR™ Green RNA-to-CT™ 1-Step Kit (ThermoFisher, #4391178).

#### Plaque assay

Six-well plates were seeded with Vero E6 cells (4E + 05/well)16 hours before the start of the assay. Samples were thawed at 37°C and centrifuged at 10,000 rpm for 5 minutes to separate the supernatant from any residual tissue debris. The sample supernatants were serially diluted ten-fold in 1x PBS containing 1% FBS. To obtain the lower limit of detection, a 1:2 dilution was also included. The plated Vero E6 cells were infected with each virus dilution and incubated for 1 hour at 37°C with shaking every 10 minutes. Post-incubation, the virus solution was aspirated, and the cells were overlayed with 0.6% Avicel solution supplemented with 3% FBS and 1x MEM. The plates were incubated in a CO<sub>2</sub> incubator for 72 hours. The Avicel overlay was aspirated, and the cell monolayer was fixed with 10% neutral-buffered formalin for 1 hour. The cells were stained with 1% crystal violet solution and plaques were counted for each dilution. The virus titers were determined by multiplying the number of plaques by the dilution factor. The lowest limit of quantification for this assay was determined by the dilution of the lung tissue plated.

# Histology and lung pathology assessment

Lung tissues were collected and fixed for seven days in 10% neutral-buffered formalin and then embedded dorsal side down in paraffin using standard procedures. Samples were sectioned at 5 µm, and resulting slides were stained with hematoxylin and eosin (H&E). All tissue slides were evaluated by light microscopy by a board-certified veterinary pathologist blinded to study group allocations. Representative photo images were collected using a Nikon Eclipse Ci microscope (Nikon Inc., Melville, NY) and analyzed with NIS-Elements software (Nikon Inc., Melville, NY). To assess lung histopathologic lesions in mice, we used an algorithm developed at UC Davis.<sup>38</sup> Additional scoring of airway, alveolar, and vascular pathology in affected areas was scored as previously indicated.<sup>39–41</sup> Briefly, to quantify total airway score, the extent of airway involvement, including the (i) extent of inflammation, (ii) bronchiolar epithelial hyperplasia and (iii) necrosis were individually scored (scale 0-4) and summed. To generate a total alveolar score, (i) alveolar inflammation, (ii) alveolar damage and (iii) type II pneumocyte hyperplasia were individually assessed (scored 0-4) and summed. To determine the vascular score, (i) vascular involvement and inflammation were assessed (scale 0-4) with the presence of perivascular hemorrhage/edema, fibrosis, and fibrin individually measured (absent = 0, present = 1). The airway, alveolar, and vascular scores were added together to generate a total lung pathology score.

### Quantification and statistical analysis

Statistical significance was assigned when P values were <0.05 using Prism Version 9.3.1 (GraphPad Software, San Diego, CA). Data was converted to a log scale prior to graphing and statistical analysis. All tests and values are indicated in the relevant figure legends.

## Results

# Intranasal vaccination with AdCOVID elicits SARS-CoV-2 spike-specific antibody responses

AdCOVID is a replication-deficient, E1- and E3-deleted Ad5 vector platform expressing a human codon-optimized gene for the RBD (residues 302 to 543) from the spike antigen of the Wuhan-1 strain of SARS-CoV-2 (accession number QHD43416).<sup>29,37</sup> We previously showed that AdCOVID, administered intranasally as a single dose, was highly immunogenic in both inbred C57BL/6J and outbred CD-1 mice through the induction of mucosal IgA, serum neutralizing antibodies, and CD4<sup>+</sup> and CD8<sup>+</sup> T cells.<sup>29</sup> Here, we vaccinated K18-hACE2 mice with a single dose of AdCOVID (2.2E + 09)ifu) via the intranasal route. At 21 days post-vaccination, anti-SARS-CoV-2 spike IgG antibodies were measured in sera samples using a spike cytometric bead array (CBA). All but 1 mouse (7/8) receiving AdCOVID seroconverted, generating SARS-CoV-2-specific IgG (Figure 1a). Vehicle control mice did not develop anti-spike IgG antibodies. On day 32, all mice were intranasally challenged with 1 x LD<sub>50</sub> dose of 2.0E + 4 PFU of SARS-CoV-2 strain USA/WA-1/2020. Following challenge, mice were weighed daily until day post-infection (DPI) +4. As expected, AdCOVID vaccinated animals did not lose weight, and weight loss in the control group was minimal over the first four days of infection (Figure 1b). On DPI +4, all mice were euthanized, and post-challenge spike-specific antibodies were quantified by CBA. All AdCOVID vaccinated mice had anti-spike serum IgG antibodies at DPI +4, while vehicle control mice failed to generate antigen-specific antibodies at this timepoint (Figure 1c). The presence of anti-spike IgG in all AdCOVID vaccinated mice, and the absence of anti-spike IgG antibodies in all vehicle control mice on DPI +4, suggests that the single AdCOVID mouse without detectable antibody levels prior to challenge had been successfully immunized. We next investigated the ability of intranasal AdCOVID to elicit immunity within the upper respiratory tract. Nasal washes were collected from a subset of mice at euthanasia (n = 4 or 6)group) and anti-spike antibodies were quantified by CBA. Vehicle control mice did not have quantifiable levels of antispike IgG in nasal washes. Conversely, AdCOVID vaccinated mice had high levels of anti-spike IgG present in the nasal cavity (Figure 1d). To confirm that viral challenge causes morbidity in our model, we performed a separate study wherein K18-hACE2 mice were intranasally challenged with an infectious dose of 0.5E + 4 PFU of SARS-CoV-2 strain USA/ WA-1/2020. Following challenge, mice were weighed daily until DPI +13. As was seen in the above experiment, AdCOVID vaccinated animals did not lose weight while the non-vaccinated control group had weight loss starting at DPI +3 and peaking between DPI +7-9 (Figure S1).

# Intranasal vaccination with AdCOVID limits SARS-CoV-2 replication in the upper respiratory tract

Intranasal vaccination has the potential to reduce viral shedding within the nasal cavity of individuals experiencing breakthrough infection. We therefore measured the viral load in the nasal cavity of the AdCOVID-vaccinated and vehicle control K18-hACE2 mice challenged with 2.0E + 04 ifu SARS-CoV-2. Nasal swipes were taken daily through DPI +4. Both vehicle control and AdCOVID vaccinated mice had detectable levels of viral RNA (Figure 2a) and subgenomic nucleocapsid (N) RNA indicative of replicated virus (Figure 2b) after twenty-four hours of infection. Importantly, AdCOVID mice had a significant reduction in viral RNA on DPI +3. Similarly, AdCOVID vaccinated mice had subgenomic N RNA values that trended lower than those measured in vehicle control mice throughout the course of the study. At study termination on DPI +4, viral load in the nasal cavity was measured by collecting nasal wash from a subset of animals from each vaccination group. AdCOVID vaccinated mice exhibited an 2-fold decrease in viral RNA (Figure 2c) and an 2-fold decrease in replicated subgenomic RNA (Figure 2d) compared to vehicle control mice.

Since the upper respiratory tract represents the first site of infection with SARS-CoV-2 and the nasal swab data indicated that the peak viral shedding from the nasal passage of unvaccinated animals was at DPI +3, we analyzed a separate cohort of AdCOVID and vehicle control mice (Figure 2e–g). As before, we observed that the AdCOVID-vaccinated animals exhibited



**Figure 1.** Spike-specific serum IgG responses following single dose intranasal vaccination with AdCOVID. K18-hACE2 mice (8 mice/group) were intranasally vaccinated with 2.2E + 09 ifu AdCOVID or vehicle control on day 0. All mice were challenged on day 32 with 2.0E + 04 PFU of SARS-CoV-2 strain USA/WA-1/2020 by the intranasal route. (a) Sera were collected on day 21 and analyzed individually for quantification of spike-specific IgG by CBA. (b) Body weight was measured daily. Four days post-challenge (DPI +4), spike-specific IgG in the (c) serum and (d) nasal wash was quantified by CBA. IgG results are expressed in either µg/ml or ng/ml as indicated. Bars represent mean ± SEM. Body weights are presented as the mean ± SEM. Lowest limit of quantification (LLOQ). Statistical analyses were performed with unpaired Student's *t* tests. \*\*\*, *P* < .001; \*\*\*\*, *P* < .0001.

high titers of anti-spike IgG antibodies in serum 21 days postvaccination (Figure 2e). All mice were challenged on day 32 (0.5E + 04 PFU) and euthanized on DPI +3. On DPI +3, we detected spike specific IgA antibodies in both the nasal wash and bronchial alveolar lavage (BAL) derived from the AdCOVID vaccinated mice (Figure 2f), which was not detected in the non-vaccinated animals. Consistent with our prior experiment showing decreased viral RNA in the nasal secretions of AdCOVID vaccinated mice on DPI +3, we observed a highly significant decrease in both total viral RNA (Figure 2g) and replicating virus subgenomic RNA (Figure 2h) in the nasal wash of the DPI +3 AdCOVID vaccinated mice. Indeed, subgenomic viral RNA was only detected in the nasal wash from 1 of 4 of the vaccinated animals compared to 5/5 of the vehicle control mice. These data show that intranasal vaccination with AdCOVID promotes humoral immune responses in the mucosa and can significantly decrease replicating viral load in the nasal passage at early timepoints following exposure to SARS-CoV-2.

# Intranasal vaccination with AdCOVID limits SARS-CoV-2 replication in the lower respiratory tract

Since early replication of SARS-CoV-2 was diminished in the upper respiratory tract of the AdCOVID-vaccinated animals, we next tested the ability of AdCOVID to decrease SARS-CoV -2 replication in the lungs. On DPI +4 we observed that the lung tissue of vehicle control mice had high levels of SARS-

CoV-2 RNA (Figure 3a). Consistent with this result, viral plaque assay using lung tissue from non-vaccinated mice revealed high levels of infectious virus (Figure 3b). Conversely, AdCOVID vaccinated mice had significantly reduced viral RNA levels (375-fold decrease) in the lungs, with only two mice rising above the lowest limit of quantification (LLOQ) (Figure 3a). Importantly, we were unable to recover infectious virus from the lungs of AdCOVID vaccinated mice on DPI +4 (Figure 3b). The same outcome was observed in an independent experiment when we measured total viral RNA (Figure 3c) and infectious virus (Figure 3d) in the lungs of AdCOVID-vaccinated and non-vaccinated animals on DPI +3. Therefore, intranasal AdCOVID vaccination decreased infectious virus in the lungs to undetectable levels within 3 days of exposure to SARS-CoV-2.

# Intranasal vaccination with AdCOVID limits lung inflammation following SARS-CoV-2 exposure

As AdCOVID vaccinated mice exhibited significantly reduced viral load in the lungs on DPI +3 and DPI +4, we used histology to assess inflammation and damage within the lung airways, the alveoli, and the lung vasculature (Table S1), and determined a total lung pathology score that encompassed each of these measurements. Consistent with our findings showing lower viral load in the lungs of AdCOVID-vaccinated mice, we observed a 4-fold reduction in total lung pathology scores in AdCOVID-vaccinated mice compared to vehicle control mice



**Figure 2.** AdCOVID limits viral load in the nasal cavity following intranasal SARS-CoV-2 challenge. K18-hACE2 mice were intranasally vaccinated with 2.2E + 09 ifu AdCOVID or vehicle control on day 0, challenged on day 32 with (a-d) 2.0E + 04 PFU SARS-CoV-2 strain USA/WA-1/2020 and analyzed between 0–4 days post-infection or (E-H) challenged on day 32 with 0.5E + 04 PFU SARS-CoV-2 strain USA/WA-1/2020 and analyzed on DPI +3. (a-d) Nasal swipes (n = 8 mice/group) were taken daily for qRT-PCR quantification of (A) total viral RNA and (b) subgenomic N RNA. On DPI +4, animals (n = 4–6 mice/group) were euthanized, and nasal wash was collected for qRT-PCR quantification of (c) total viral N RNA and (d) subgenomic N RNA. On DPI +4, animals (n = 4–6 mice/group) were euthanized, n = 11 mice/group) on day 21 post-vaccination. Spike-specific IgA antibodies measured in BAL and nasal wash on DPI +3 (panel F, n = 4–5 mice/group). qRT-PCR quantification of (g) total viral N RNA in nasal wash on DPI +3 (n = 4–5 mice/group). Viral quantitation presented as RNA copies. Bars represent mean ± SEM. Lowest limit of quantification (LLOQ). Statistical analyses were performed with unpaired Student's *t* tests. \*\*, P < .001; \*\*\*\*, P < .0001;



**Figure 3.** AdCOVID limits viral replication in the lungs following intranasal SARS-CoV-2 challenge. AdCOVID-vaccinated and control K18-hACE2 mice were challenged on day 32 with either (a, b) 2.0E + 04 PFU or (c, d) 0.5E + 04 PFU SARS-CoV-2 strain USA/WA-1/2020. On (a-b) DPI +4 or (c, d) DPI +3, animals were euthanized, and both lung lobes were harvested. (A, C) Total viral N RNA was quantified by qRT-PCR. Results are expressed as RNA copies. (b, d) Infectious virus was quantified in the lungs by plaque assay. Bars represent mean  $\pm$  SEM. Lowest limit of quantification (LLOQ). Statistical analyses were performed with unpaired Student's *t* tests. \*, *P* < .05; \*\*, *P* < .01; \*\*\*\*, *P* < .001; \*\*\*\*, *P* < .001.

on DPI +4 (Figure 4a). Notably, mice receiving vehicle control showed evidence of viral pneumonia following SARS-CoV-2 challenge (Figure 4b, left panel) as characterized by immune (polymorphonuclear and mononuclear) cell accumulation in perivascular, intimal (Figure 4b, middle panel), and alveolar interstitium (Figure 4b, right panel) as well as evidence of pneumocyte and bronchial epithelial hyperplasia (Table S1). In contrast, AdCOVID-vaccinated mice showed a marked reduction in alveolar inflammation (Figure 4c, left panel) with the exception of the accumulation of mononuclear cells around medium to large caliber pulmonary vessels (Figure 4c, middle panel), and no evidence of alveolar interstitial inflammation (Figure 4c, right panel). Thus, intranasal vaccination with AdCOVID greatly reduces early virus-elicited airway, alveolar, and vascular inflammation, possibly preventing alveolar damage and hyperplasia of Type II pneumocytes and bronchial epithelial cells.

# Intranasal AdCOVID vaccination prevents SARS-CoV-2 replication in the periphery of K18-hACE2 mice

It was previously reported that intranasal SARS-CoV-2 infection can cause encephalitis and, in some cases, death in a fraction of infected K18-hACE2 mice.<sup>42</sup> To determine whether AdCOVID vaccination can prevent neuroinvasion in the K18-hACE2 animals, we quantified viral RNA and infectious virus in the brain of the SARS-CoV-2 challenged mice. Consistent with other reports,<sup>43</sup> 3/4 SARS-CoV-2 infected control mice had detectable viral RNA in the brain on DPI +4 (Figure 5a) and infectious virus was

recovered by plaque assay from the brains of two of these animals (Figure 5b). In contrast, neither viral RNA (Figure 5a) nor infectious virus (Figure 5b) was recovered from the brains of any of the AdCOVID vaccinated-SARS-CoV-2 infected animals. Collectively, these data suggest that a single vaccination with AdCOVID was sufficient to decrease viral load in the nasal epithelium and lungs of SARS-CoV-2 challenged mice at early timepoints postinfection and was able to minimize the risk of viral spread to the brain.

# Discussion

Preclinical and clinical studies imply that intranasal vaccination has the potential to be efficacious against respiratory viruses.44-47 Intranasal administration bypasses preexisting adenovector immunity and elicits comparable immunity to intramuscular injection.<sup>22,46</sup> Consistent with these reports, we previously showed that AdCOVID, an intranasal Ad5vectored vaccine encoding the RBD of the SARS-CoV-2 spike protein, elicited long-lived, protective systemic and mucosal immunity following a single administration in mice.<sup>29</sup> Here, we further expanded upon these findings. A single administration of AdCOVID limited viral replication in the upper and lower respiratory tracts following intranasal SARS-CoV-2 challenge. AdCOVID vaccinated K18-hACE2 mice had significantly reduced levels of viral RNA in both the nasal wash and nasal swipes. AdCOVID vaccination also markedly impaired viral replication in the lungs of challenged mice, as evidenced by the reduction in



**Figure 4.** AdCOVID protects mice from SARS-CoV-2 induced lung inflammation. AdCOVID and control vaccinated K18-hACE2 mice were challenged with 2.0E + 04 PFU SARS-CoV-2 strain USA/WA-1/2020 on day 32. Mice were euthanized on DPI +4 (n = 4/group) and evaluated for lung inflammation and damage. (a) Total lung pathology score representing airway, alveolar, and vascular changes associated with inflammation, cellular hyperplasia and damage. Individual points represent the total pathology score/mouse (See Table S1 for individual scores per mouse/measurement). (b, c) Representative lung sections stained with hematoxylin and eosin and imaged at 40X (left; scale bar, 500 µm) and 400X (middle & right; scale bar, 50 µm). Bars in panel a represent mean ± SEM. Statistical analysis (n = 4 mice/group) was performed with an unpaired Student's *t* test. \*\*\*, *P* < .001.

subgenomic viral RNA and the absence of infectious virus. Importantly, AdCOVID vaccinated mice exhibited little evidence of viral pneumonia, with a near absence of alveolar inflammation.

It is becoming increasingly apparent that the nasal cavity serves as the principal point of viral entry.<sup>48</sup> The nasal epithelium has a high concentration of ACE2, and ciliated cells in the nasal cavity are the primary targets for SARS-CoV-2 replication during the early stages of infection.<sup>4,49</sup> Studies utilizing the ferret model of influenza indicate that viral seeding of lower airways is heavily dependent on viral replication and shedding in the nasal cavity.<sup>50</sup> Our findings suggest that intranasal vaccination has the potential to substantially limit viral replication within the nasal cavity, thereby preventing productive infection of the lower respiratory tract. AdCOVID vaccinated mice had significant reductions in the level of viral RNA within the upper respiratory tract. Reduced viral seeding, combined with AdCOVID-elicited anti-RBD IgA and anti-RBD T cell populations in the lungs, most likely prevented productive SARS-CoV -2 infection of alveolar Type II cells and histopathological damage.<sup>29,51</sup> Additionally, the lack of lung inflammation implies protection from immunopathology,<sup>52,53</sup> and may reduce the possibility of long-term COVID disease. Similar pathology results have been reported using other intranasal vaccine modalities.<sup>27,54</sup>

Of further importance, it is thought that intranasal vaccines have the potential to block transmission between vaccinated hosts and antigenically naïve bystanders.<sup>55,56</sup> It was recently reported that intranasal vaccination with a parainfluenza virus type 5 (PIV5)-based SARS-CoV-2 vaccine prevented viral transmission from vaccinated ferrets to co-housed, naïve ferrets.<sup>57</sup> AdCOVID's ability to block SARS-CoV-2 transmission remains to be confirmed, but the reduced viral burden in the respiratory tract suggests that such a beneficial effect is feasible.

While effective at preventing severe COVID-19 disease, evidence suggests that IM vaccination does not elicit longlasting, sterilizing immunity within the respiratory tract.<sup>20</sup> IM vaccination of rhesus macaques with an adenoviral vectored vaccine candidate prevented viral pneumonia, but viral replication in the upper respiratory tract was unaffected.<sup>58</sup> It has also been reported that IM vaccination with CoronaVac, an inactivated virus vaccine, failed to produce antibody responses in the nasal epithelial lining fluid (NELF) of human vaccinees.<sup>19</sup> This same study showed that the Comirnaty mRNA vaccine elicited low levels of neutralizing antibodies in the NELF which waned within two months of vaccination.<sup>19</sup> Pre-clinical studies suggest that intranasal SARS-CoV-2 vaccination is capable of inducing sustained mucosal immunity,<sup>29</sup> but it remains to be seen if intranasal vaccines are sufficiently



**Figure 5.** AdCOVID prevents neuroinvasion following intranasal SARS-CoV-2 challenge. K18-hACE2 mice were challenged with 2.0E + 04 PFU SARS-CoV-2 strain USA/WA-1/2020 on day 32. On DPI +4, mice (n = 4/group) were euthanized, and the brains were harvested. (a) Total viral N RNA was quantified by qRT-PCR. Results are expressed as RNA copies. (b) Infectious virus was quantified in the whole brain by plaque assay. Bars represent mean  $\pm$  SEM. Lowest limit of quantification (LLOQ). Statistical analyses were performed with unpaired Student's *t* tests.

immunogenic to yield protective immunity in humans without preexisting immunity.<sup>59</sup> The dearth of marketed intranasal vaccines suggests that the path to licensure as a standalone vaccine may be difficult.

Despite ever-increasing numbers of infected persons and extensive deployment of SARS-CoV-2 vaccines, intranasal vaccines may still offer advantages for the control of the COVID-19 pandemic by serving as a priming regimen or a booster intervention. Vaccination regimens consisting of IM priming and intranasal boosting have been well studied and it appears to be an effective approach. Intramuscular prime-Ad5 intranasal boosting resulted in improved peripheral and local humoral immunity in SARS-CoV-2 vaccinated mice compared to two doses delivered by either route alone.<sup>60</sup> Antibodies in the lungs had broad potency and were capable of neutralizing multiple SARS-CoV-2 viral variants. Heterologous prime-boost also generated significantly higher numbers of lung-resident T cells compared to IM vaccination alone. Furthermore, a heterologous prime-boost strategy completely protected hamsters from COVID-19 disease Syrian and immunopathology.<sup>61</sup> Indeed, similar results were reported in a recent intranasal vaccine Phase 1 trial where heterologous delivery routes yielded improved neutralizing antibody titers while maintaining T cell immunity.<sup>59</sup> These data suggest that heterologous delivery vaccination regimens may result in balanced systemic and mucosal immunity capable of protecting against divergent SARS-CoV-2 strains.

In summary, there remains a need for SARS-CoV-2 vaccines capable of eliciting mucosal immunity. The reduced viral shedding measured in the upper and lower respiratory tracts of pre-clinical challenge models establishes the ability of intranasal vaccines to activate local immune cell populations. How best to deploy intranasal SARS-CoV-2 vaccine regimens remains unclear, but they have the potential to be key weapons in the fight against COVID-19.

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#### **Author contributions**

Conceptualization: J.J.S., R.G.K., T.D.R, B.G., F.E.L., and M.S.R.; Methodology: M.D.S., J.J.S., D.B., A.S-S., R.G.K, C.D.M., J.B.F., K.S.H., S.M.L., T.D.R., M.S.R., F.E.L. and B.G.; Investigation: R.G.K., T.D.R., and F.E.L.; Reagent Development and Data Collection: M.D.S., D.B., A.S-S.,R. G.K., T.W.D., C.D.M. J.B.F., F.Z., J.L.T.; Vaccine candidate design/production: J.Z., B.G., and M.S.R; Writing-Original Draft Preparation: J.J.S., B.G., and M.S.R.; Writing-Review and Editing: All authors; All authors had full access to all the data in the study and take responsibility for integrity of the data and the accuracy of the data analysis.

# ORCID

Michael D. Schultz () http://orcid.org/0000-0002-6741-4115 John J. Suschak () http://orcid.org/0000-0002-2867-8265 Davide Botta () http://orcid.org/0000-0003-3926-0662 Aaron Silva-Sanchez () http://orcid.org/0000-0002-1882-2534 R. Glenn King D http://orcid.org/0000-0001-6522-0053 Chetan D. Meshram D http://orcid.org/0000-0001-5426-2063 Jeremy B. Foote D http://orcid.org/0000-0002-6621-6096 Kevin S. Harrod D http://orcid.org/0000-0003-0780-9470 Troy D. Randall D http://orcid.org/0000-0003-0643-0311 Frances E. Lund D http://orcid.org/0000-0003-3083-1246

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