# nature portfolio

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## **Reporting Summary**

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

Statistics			
For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
/a Confirmed			
The exact sample size $(n)$ for each experimental group/condition, given as a discrete number and unit of measurement			
A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly			
The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.			
A description of all covariates tested			
A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons			
A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)			
For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted Give <i>P</i> values as exact values whenever suitable.			
For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings			
For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes			
$\square$ Estimates of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated			
Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.			
Software and code			
Policy information about <u>availability of computer code</u>			
Data collection N/A			
Data analysis  Bulk RNA seq and single cell RNA seq is analyzed by R. Body weights, viral load, pathology is analyzed by Graphpad Prism. Fluorescence and inflammation level is analyzed by HALO.			
For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.			

#### Data

Policy information about <u>availability of data</u>

All manuscripts must include a <u>data availability statement</u>. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

Provide your data availability statement here.

## Research involving human participants, their data, or biological material

Policy information and sexual orientat		vith <u>human participants or human data</u> . See also policy information about <u>sex, gender (identity/presentation),</u> thnicity and racism.	
Reporting on sex	and gender	One female patient and two male patients were included as human data. We designed to included both sex in study design.  Sex/ gender was determined by self-reporting or assigned. Human case summary is reported as supplemental table 12.	
Reporting on race other socially rele groupings		The socially constructed or socially relevant categorization variables are not used in the manuscript.	
Population chara	cteristics	The detailed information is summarized in supplemental table 12.	
Recruitment		Participants are recruited by the disease severity. Severe COVID patients showing pulmonary disease are further included in the study.	
Ethics oversight		Tulane University approved the study protocol.	
	_	oval of the study protocol must also be provided in the manuscript.	
Field-spe	ecific re	porting	
	ne below that is	the best fit for your research. If you are not sure, read the appropriate sections before making your selection.	
☐ Behavioural & social sciences ☐ Ecological, evolutionary & environmental sciences			
For a reference copy of t	the document with a	all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>	
Life scier	nces stu	udy design	
All studies must dis	close on these	points even when the disclosure is negative.	
Sample size	In our previous published paper (doi: 10.7150/thno.61810. eCollection 2021.). We performed a comprehensive test on both male and female K18 mice infected with SARS-CoV-2. Based on these data, we didn't do statistical methods to predetermine sample size.		
Data exclusions	No data were ex	e excluded from the analysis.	
Replication	facility, it is not	nce our focus is long-term covid, we collected the tissue as long as 21 days and 45 days post infection. Together with the limitation of BSL3 cility, it is not practical to replicate the experiment for several times. To increase the vadility and reproductivity, we tested the samples from ultiple time points such as 7, 14, 21, and 45 days post infection.	
Randomization	Randomization	domization is not relevant to our study. To increase the validity, the virus is diluted from the same stock.	
Blinding	The pathologist performed the histology analysis is blinded to the animal grouping.		
Reportin	g for sp	pecific materials, systems and methods	
		about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.	
Materials & exp	perimental sy	ystems Methods	
n/a Involved in the study			
Clinical dat	a esearch of concer	n	
Plants	54. 5.1 6. 66.16611		

## **Antibodies**

Antibodies used Anti-SARS Anti-SARS-CoV 1: 1000 BEI Resources # NR-10361

Anti-CD206 Anti-mouse 1: 50 R & D # AF2535 Anti-SMA Anti-mouse 1: 100 Abcam # ab5694-100 Anti-Krt-5 Anti-mouse 1: 1000 BioLegend # 905901 Anti-SMA Anti-human 1: 100 Agilent # M085129-2 Anti-Krt-5 Anti-human 1: 50 Abcam # ab17130

Anti-pro SPC Anti-mouse 1: 500 Seven Hills Bioreagents # WRAB-9337 Anti-trp63 Anti-mouse 1: 200 Cell Signaling Technology # 13109

Anti-NGFR Anti-mouse 1: 100 Abcam # ab52987

Validation The validation of each primary antibody is performed by manufacturers.

 $https://www.beiresources.org/Catalog/BEIPolyclonalAntiserum/NR-10361.aspx \\ https://www.rndsystems.com/products/mouse-mmr-cd206-antibody_af2535$ 

https://www.abcam.com/products/primary-antibodies/alpha-smooth-muscle-actin-antibody-ab5694.html https://www.biolegend.com/fr-fr/explore-new-products/keratin-5-polyclonal-chicken-antibody-purified-10957

https://www.agilent.com/en/product/immunohistochemistry/antibodies-controls/primary-antibodies/actin-%28smooth-muscle%29-

%28concentrate%29-76542

https://www.abcam.com/products/primary-antibodies/cytokeratin-5-antibody-xm26-ab17130.html

https://www.sevenhillsbioreagents.com/products/anti-pro-sp-c-rabbit-n-terminal

https://www.cellsignal.com/products/primary-antibodies/p63-a-d2k8x-xp-rabbit-mab/13109

https://www.abcam.com/products/primary-antibodies/p75-ngf-receptor-antibody-ep1039y-bsa-and-azide-free-ab256584.html

## Animals and other research organisms

Policy information about <u>studies involving animals</u>; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Laboratory animals Strain Sex Sample Size Age/ week Inoculum Infection Dose Timepoint/DPI

K18-hACE2 +/- Male and Female 3 28 N/A N/A 0

K18-hACE2 +/- Male 1 16 SARS-CoV-2(WA1/2020) 1.0X 10^4/ TCID 50 3

K18-hACE2 +/- Male and Female 3 6-10 SARS-CoV-2(WA1/2020) 2.0X 10^5/ TCID 50 7 K18-hACE2 +/- Male and Female 3 12-16 SARS-CoV-2(WA1/2020) 1.0X 10^4/ TCID 50 7 K18-hACE2 +/- Male and Female 3 12-16 SARS-CoV-2(WA1/2020) 1.0X 10^4/ TCID 50 14

K18-hACE2 +/- Male and Female 3 10 SARS-CoV-2(WA1/2020) 1.0X 10^4/ TCID 50 21 K18-hACE2 +/- Female 2 6-10 SARS-CoV-2(WA1/2020) 1.0X 10^4/ TCID 50 45

C57BL/6J Male 3 6-8 H1N1 A/PR/8/34 (PR8) 50 PFU 7 C57BL/6J Male 4 6-8 H1N1 A/PR/8/34 (PR8) 50 PFU 14 C57BL/6J Male 5 6-8 H1N1 A/PR/8/34 (PR8) 50 PFU 21

K18-hACE2 +/- Female 4 8 N/A N/A 0

K18-hACE2 +/- Female 3 6-10 SARS-CoV-2(WA1/2020) 2.0X 10^5/ TCID 50 4 K18-hACE2 +/- Female 3 6-10 SARS-CoV-2(WA1/2020) 2.0X 10^5/ TCID 50 6

K18-hACE2 +/- Female 3 8-10 H1N1 A/PR/8/34 (PR8) 50 PFU 4 K18-hACE2 +/- Female 3 8-10 H1N1 A/PR/8/34 (PR8) 50 PFU 6

K18-hACE2 +/- Female 1 12 SARS-CoV-2(WA1/2020) 2.0X 10^5/ TCID 50 4

K18-hACE2 +/- Female 1 12 H1N1 A/PR/8/34 (PR8) 50 PFU 4

Wild animals No wild animals are included in the study

Reporting on sex The finding applys to both sex. No sex-based analysis is performed.

Field-collected samples No field-collected samples are included in the study.

Ethics oversight Tulane University

Note that full information on the approval of the study protocol must also be provided in the manuscript.

### Clinical data

Policy information about clinical studies

All manuscripts should comply with the ICMJE guidelines for publication of clinical research and a completed CONSORT checklist must be included with all submissions.

Clinical trial registration N

N/A

Study protocol

N/A

April 2023

Data collection N/A
Outcomes N/A