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Reporting Summary

Nature Research 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 Research policies, see <u>Authors & Referees</u> and the <u>Editorial Policy Checklist</u>.

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Statistics For all statistical and	lyses confirm that the following items are present in the figure legand, table legand, main text, or Methods section			
I	or all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
	ample size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
	t 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	A description of all covariates tested			
A description	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 hyp	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 Bayesia	n analysis, information on the choice of priors and Markov chain Monte Carlo settings			
For hierarc	hical and complex designs, identification of the appropriate level for tests and full reporting of outcomes			
Estimates of	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 a	bout <u>availability of computer code</u>			
Data collection	no software is used.			
Data analysis	GraphPad Prism 7.0			
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/review. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.				
Data				
Policy information a	bout <u>availability of data</u>			
All manuscripts mu - Accession codes, - A list of figures th	st include a <u>data availability statement</u> . This statement should provide the following information, where applicable: unique identifiers, or web links for publicly available datasets lat have associated raw data unique identifiers, or data availability			
The data supporting the findings of this study are available from the corresponding authors upon written request.				
Field-spe	cific reporting			
Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.				
Life sciences	Behavioural & social sciences Ecological, evolutionary & environmental sciences			

For a reference copy of the document with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.		
Sample size	No experiment presented in this study requires sample size to be determined	
Data exclusions	no excluded data	
Replication	All infection experiments were repeated at least two independent times. All attempts at replication were successful. Electron Microscopy was performed once, whereas there are at least 5 images for each cell type. The multiplex PCR array was performed one time in triplicate.	
Randomization	No experiment presented in this study requires randomization.	
Blinding	The results are either presented in quantitative data or with a negative control. Blinding is not required.	

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

Materials & experimental systems		Methods	
n/a	Involved in the study	n/a	Involved in the study
		\boxtimes	ChIP-seq
	Eukaryotic cell lines	\boxtimes	Flow cytometry
\boxtimes	Palaeontology	\boxtimes	MRI-based neuroimaging
	Animals and other organisms		•
	Human research participants		
\times	Clinical data		

Antibodies

Antibodies used

Antibodies Company/Supplier Catalog number LOT number Dilution ACE2 R&D Systems AF933 HOK0420021 1:100

TMPRSS2 Invitrogen PA5-14264 SI2433682G 1:100

Villin Abcam ab130751 GR3229609-9 1:100

Goat anti-Mouse 488 Invitrogen A-11001 2090562 1:500

Goat anti-Rabbit 488 Invitrogen A-11034 1885241 1:500

Goat anti-Rabbit 594 Invitrogen A-11037 1608397 1:500

Donkey anti-Goat 488 Invitrogen A-11055 1771339 1:500

mouse anti-viral NP in-house made n.a n.a 1:10000

rabbit anti-viral NP in-house made n.a n.a 1:10000

Validation

- 1) the application of commercial antibodies for IF staining has been validated by the providers.
- 2) the validation result of in-house generated antibodies will be provided upon request.

Eukaryotic cell lines

Policy information about <u>cell lines</u>	
Cell line source(s)	Vero cells from ATCC (https://www.atcc.org/products/all/CCL-81.aspx)
Authentication	The cells are purchased from ATCC.
Mycoplasma contamination	we verified the cells free of mycoplasma contamination by using the mycoplasma screening service provided by HKU core facilities.
Commonly misidentified lines (See ICLAC register)	No commonly misidentified lines were used.

Animals and other organisms

Wild animals

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

Laboratory animals no lab animal involved.

Three female and male horseshoe bats (Rhinolophus sinicus) were captured in roosting caves with hand-nets on long pole by bat experts of Agriculture, Fisheries and Conservation Department, the Government of Hong Kong Special Administrative Region. The captured bats were kept in cloth bags individually and were transferred by the technical staff to HKU laboratory right after the capture. The organs were applied to establishing organoid culture after the bats were euthanized in biosafety hood by intraperitoneal injecting overdosed pentobarbitone sodium (100-150mg/kg). We have obtained the ethnic approval for this study.

no field-collected sample involved. Field-collected samples

Ethics oversight Approved by Agriculture, Fisheries and Conservation Department, the Government of Hong Kong Special Administrative Region,

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

Human research participants

Population characteristics

Policy information about studies involving human research participants

Four COVID-19 patients who were confirmed by laboratory test. This study only used the patients' respiratory and fecal specimens for virus isolation and analysis. Patients themselves were not involved in any experiments.

Recruitment The patients are confirmed cases of SARS-CoV-2 infection in two public hospital in Hong Kong.

Institutional Review Board of the University of Hong Kong/Hospital Authority Hong Kong West Cluster (UW 13-372) Ethics oversight

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