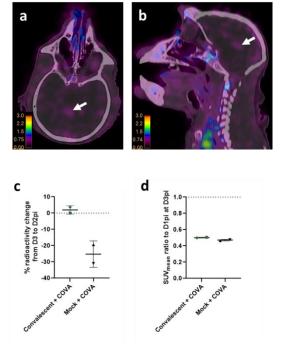


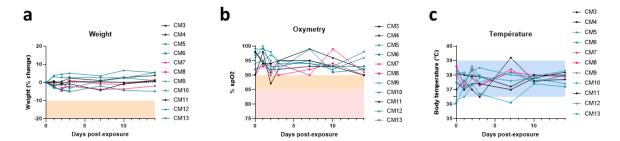
 $Supplementary\ figure\ 1.\ Size\ exclusion\ chromatography\ chromatograms$ 

(a) SEC chromatogram of COVA 1-27; rt = 25.5 min; (b) SEC chromatogram of IgG1 kappa; rt = 23.1 min; (c) SEC chromatogram of COVA 1-27-DFO conjugate; rt = 24.4, 25.2 min; (d) SEC chromatogram of IgG1 kappa; rt = 23.1 min; (e) SEC chromatogram of radiolabelled COVA 1-27;  $\gamma$  detection, rt = 19.2, 22.8 min; (f) SEC chromatogram of radiolabelled IgG1 kappa;  $\gamma$  detection, rt = 34.8, 42.5 min; rt = 23.1 min; (g) SEC chromatogram of radiolabelled COVA 1-27; UV detection, rt = 21.6, 22.6 min; (h) SEC chromatogram of radiolabelled IgG1 kappa; UV detection, rt = 35.9, 42.2 min



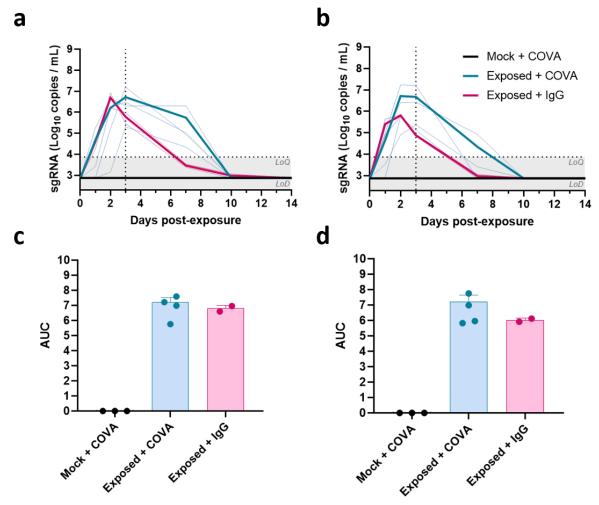
## Supplementary figure 2. COVA1-27 uptake in the brain of convalescent animals

(a, b) [ $^{89}$ Zr]COVA1-27-DFO PET signal in the brain indicated by the white arrows, sagittal and axial view respectively at D3pi. (CM1). (c, d) Corresponding quantification of the maximum (c) and mean (d) [ $^{89}$ Zr]COVA1-27-DFO PET signal in the brain of all animals at D3pi with ratio to D2pi and D1pi respectively. Data are represented as individual values with mean and SD (n = 2 animals per group).

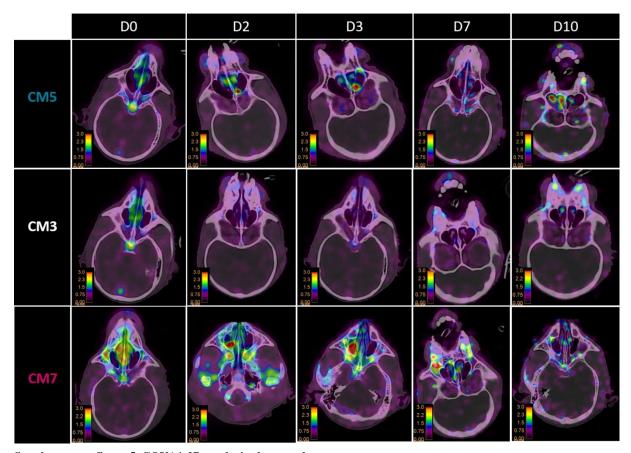


## **Supplementary figure 3. Clinical observations**

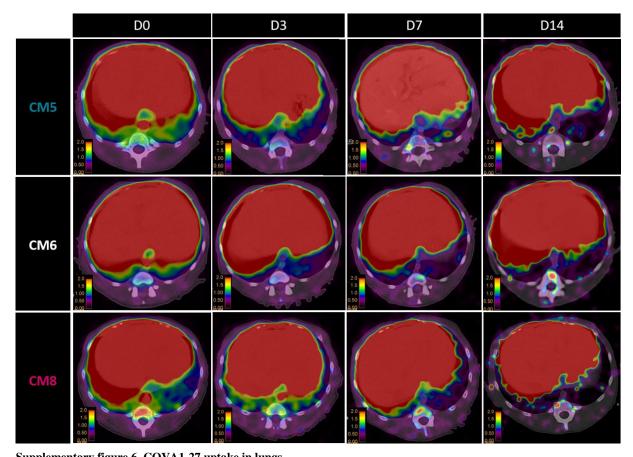
(a) Weight, (b) oxymetry and (c) temperature evolution after exposure, mock+COVA group is represented in black (circle, pyramid, triangle), exposed +COVA group is in turquoise (pyramid, triangle, circle, square, diamond, hexagon) and exposed+IgG in pink (circle, square).



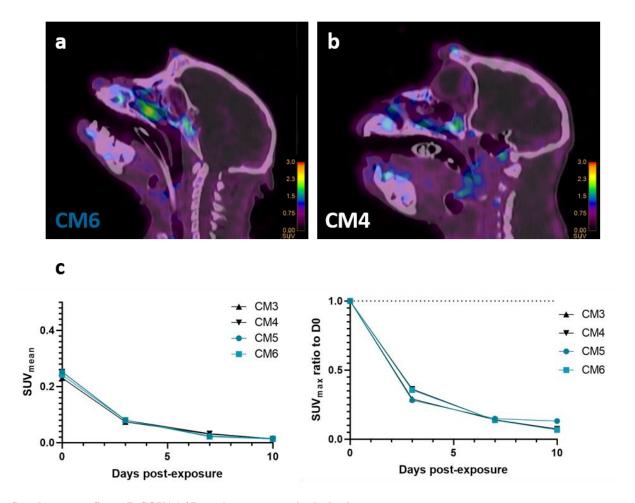
Supplementary figure 4. SARS-CoV-2 subgenomic RNA (sgRNA) titres over time in nasopharynx and trachea Evolution of the of the SARS-CoV-2 sgRNA titres ( $Log_{10}$  copies/mL) over two weeks in the nasopharynx (a) and the trachea; (b) The individual follow up is represented by the thin lines and the average of each group in bold; Area under the curve (AUC) for each group in the nasopharynx (c) and the trachea (d) mock+COVA group is represented in black (n = 3animals), exposed +COVA group is in turquoise (n = 4 animals) and exposed+IgG in pink (n = 2 animals), individual values are represented by circles



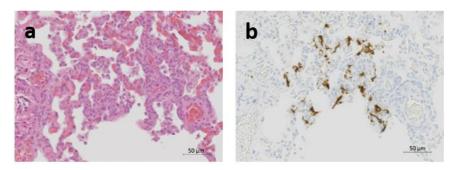
Supplementary figure 5. COVA1-27 uptake in the nasopharynx
Representative images of the nasal cavity at day 0, 2, 3, 7, and 10 post-exposure (pe) for each group: CM5 (turquoise, exposed +COVA), CM3 (white, mock+COVA), and CM7 (pink, exposed+IgG), color scale 0-3 SUV.



Supplementary figure 6. COVA1-27 uptake in lungs
Representative images of the lungs at day 0, 3, 7, and 14 post-exposure (pe) for each group: CM6 (turquoise, exposed +COVA),
CM3 (white, mock+COVA), and CM7 (pink, exposed+IgG), color scale 0-2 SUV.

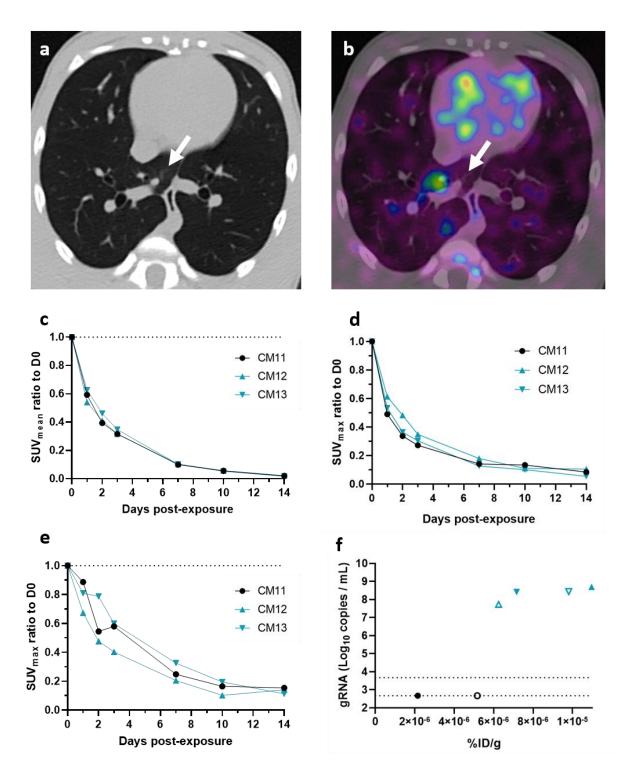


Supplementary figure 7. COVA1-27 uptake assessment in the brain (a-b) Representative images of the brain at day 3 post-exposure (pe) of animals injected with [89Zr]COVA1-27-DFO for each group: CM6 (turquoise, exposed +COVA), CM4 (white, mock+COVA), color scale 0-3 SUV. (c-d) Average (SUVmean) or maximum (SUVmax) brain [89Zr]COVA1-27-DFO uptake over time in [89Zr]COVA1-27-DFO injected animals exposed (turquoise) or not (black) to SARS-CoV-2



## Supplementary figure 8. Representative viral distribution in the lung parenchyma of $\ensuremath{\mathrm{CM10}}$

(a) Histological characterisation using Hematoxylin Eosin (HE) staining in the caudal right lobe of the lungs of CM10 and corresponding (b) cellular-scale evaluation of the virus distribution (brown signal) using in situ hybridisation (ISH) in the lungs (caudal left lobe)



Supplementary figure 9. Low dose COVA1-27 uptake assessment in lungs and nasal cavity (a) Representative CT lung lesion indicated by the white arrow and (b) its associated PET signal indicated by a white arrow, color scale 0-3 SUV; (c) Longitudinal evaluation of the normalized PET signal SUV $_{mean}$  and (d) SUV $_{max}$  to D0 post-exposure (0 dpe) in the lungs of animals; (e) Longitudinal evaluation of the normalized PET signal SUV $_{max}$  to 0 dpe in the nasal cavity of animals; (f) Evolution from 2 (hollow symbol) to 3 dpe (filled symbol) of the radioactivity in the nasopharyngeal swabs correlated with the SARS-CoV-2 gRNA titres in the same sample

Organ samples list	Included in the analysis			
Lung – cranial right lobe	X			
Lung – cranial left lobe	X			
Lung – caudal right lobe	X			
Lung – caudal left lobe	X			
Lung –right lingula lobe	X			
Lung –left lingula lobe	X			
Lung – accessory lobe	X			
Trachea	X			
Spleen	X			
Axillary lymph node	X			
Inguinal lymph node	X			
Tracheobronchial lymph node	X			
Heart	X			
Kidney -Right	X			
Kidney -Left	X			
Liver				

 $\label{eq:constraints} \textbf{Supplementary table 1. Organ sampling at necropsy}$ 

Animal ID	Probe	Tracer dose	Exposition	Group	Follow up time	Nasal rhinorea (Y/N)	Nasal SUVmax ratio at D3	Nasal swab radioactivity at D3 (%ID/g)	Nasal swab RNA titers at D3 (RNA copies/mL)
CM1	COVA1-27	2,5mg	Convalescents SARS-CoV-2 (3months)	Convalescent + COVA	1 week	No	0,44	5,42E-06	0,00E+00
CM2	COVA1-27	2,5mg	Convalescents SARS-CoV-2 (3months)	Convalescent + COVA	1 week	No	0,41	4,62E-05	0,00E+00
CM3	COVA1-27	2,5mg	PBS	Mock + COVA	2 weeks	No	0,45	3,15E-06	0,00E+00
CM4	COVA1-27	2,5mg	PBS	Mock + COVA	2 weeks	No	0,49	1,97E-06	0,00E+00
CM5	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA	2 weeks	Yes	1,00	8,14E-06	2,15E+07
CM6	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA	2 weeks	No	0,51	8,95E-06	2,74E+08
CM7	lgG1	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + IgG	2 weeks	Yes	0,63	2,23E-06	1,11E+08
CM8	lgG1	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + IgG	2 weeks	No	0,46	4,01E-06	1,52E+08
CM9	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (eutha D3)	3 days	No	0,47	1,91E-05	/ (sampling issue)
CM10	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (eutha D3)	3 days	No	0,58	1,08E-05	3,45E+07
CM11	COVA1-27	150µg	PBS	Mock + COVA (low dose)	2 weeks	No	0,58	2,15E-06	0,00E+00
CM12	COVA1-27	150µg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (low dose)	2 weeks	No	0,40	1,10E-05	5,09E+08
CM13	COVA1-27	150µg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (low dose)	2 weeks	No	0,60	7,17E-06	2,64E+08

Animal ID	Probe	Tracer dose	Exposition	Group	Follow up time	Lung lesions (Y/N)	Type of lung lesions (number)	Lung lesion ROI SUVmax ratio at D7	Brain SUVmax ratio at D3
CM1	COVA1-27	2,5mg	Convalescents SARS-CoV-2 (3months)	Convalescent + COVA	1 week	No	/	/	0,80
CM2	COVA1-27	2,5mg	Convalescents SARS-CoV-2 (3months)	Convalescent + COVA	1 week	Yes	GGO (n=1)	0,28	0,64
CM3	COVA1-27	2,5mg	PBS	Mock + COVA	2 weeks	No	/	/	0,43
CM4	COVA1-27	2,5mg	PBS	Mock + COVA	2 weeks	No	/	/	0,48
CM5	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA	2 weeks	Yes	GGO (n=3)	0,22	0,50
CM6	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA	2 weeks	Yes	GGO (n=2)	0,34	0,47
CM7	lgG1	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + IgG	2 weeks	Yes	GGO (n=3)	0,22	not assessed
CM8	IgG1	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + IgG	2 weeks	Yes	GGO (n=1)	0,17	not assessed
CM9	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (eutha D3)	3 days	Yes	GGO (n=1)	not assessed (eutha D3)	0,46
CM10	COVA1-27	2,5mg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (eutha D3)	3 days	Yes	GGO (n=2)	not assessed (eutha D3)	0,40
CM11	COVA1-27	150µg	PBS	Mock + COVA (low dose)	2 weeks	No	/	/	not assessed
CM12	COVA1-27	150µg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (low dose)	2 weeks	Yes	GGO (n=3)	0,18	not assessed
CM13	COVA1-27	150µg	SARS-CoV-2 (B.1.617.2)	Exposed + COVA (low dose)	2 weeks	Yes	GGO (n=2)	0,13	not assessed

Supplementary table 2. Individual data summary