

# 156. Novel nanoparticles composition

(University of Pennsylvania)



## ▶ Asset Overview

<b>Product Type</b>	Antibody + Nanoparticle
<b>Disease Area</b>	Inflammatory Disease
<b>Indication</b>	Acute Inflammatory Disease
<b>Current Stage</b>	Lead Optimization
<b>Target</b>	Neutrophils in Acute Lung Inflammation (ALI)
<b>MoA</b>	Arrangement of protein in or on the nanoparticles via hydrophobic interactions, crosslinking and electrostatic interactions
<b>Brief Description</b>	<ul style="list-style-type: none"><li>• Acute inflammatory diseases, including sepsis, pneumonia, and acute respiratory distress syndrome (ARDS), account for over 1.5 million hospitalizations and 300,000 deaths per year in the US. These diseases all share a common cellular player known as marginated leukocytes.</li><li>• Marginated leukocytes are white blood cells that accumulate in the blood vessels of inflamed organs and potentiate disease progression by releasing toxins and pro-inflammatory cytokines and induce clotting and further inflammation. Accumulation of marginated leukocytes can ultimately lead to organ dysfunction.</li><li>• Inventors developed a molecular label consisting of an IgG antibody bound to dibenzocyclooctyne (D20 tag). When attached to nanoparticles, the D20 tag almost exclusively localized to marginated leukocytes to alleviate disease symptoms in a small animal model of acute inflammatory injury.</li><li>• Using the D20 tag to target marginated leukocytes with imaging agents or drugs could better enable clinical diagnosis and treatment of acute inflammatory diseases.</li></ul>
<b>Intellectual Property</b>	WO2021/113519A1
<b>Publication</b>	Supramolecular Arrangement of Protein in Nanoparticle Structures Predicts Nanoparticle Tropism for Neutrophils in Acute Lung Inflammation. Nat Nanotechnol. (2022)
<b>Inventors</b>	Priyal PATEL, Jacob Myerson, Jacob BRENNER, Vladimir R. Muzykantov, Landis R. WALSH

## ▶ Highlights

- Provides ~ 50% more protection against cellular leakage than non-tagged nanoparticles following inflammatory injury, which is key for alleviating acute inflammatory disease progression
- Provides ~ 2x specificity to inflamed lung tissue versus heart, liver, spleen, and kidneys following inflammatory lung injury, which is key for accurately diagnosing and effectively treating acute inflammatory diseases
- Compatible with liposomes for small molecule drug or imaging agent delivery and solid lipid nanoparticles for modified mRNA delivery

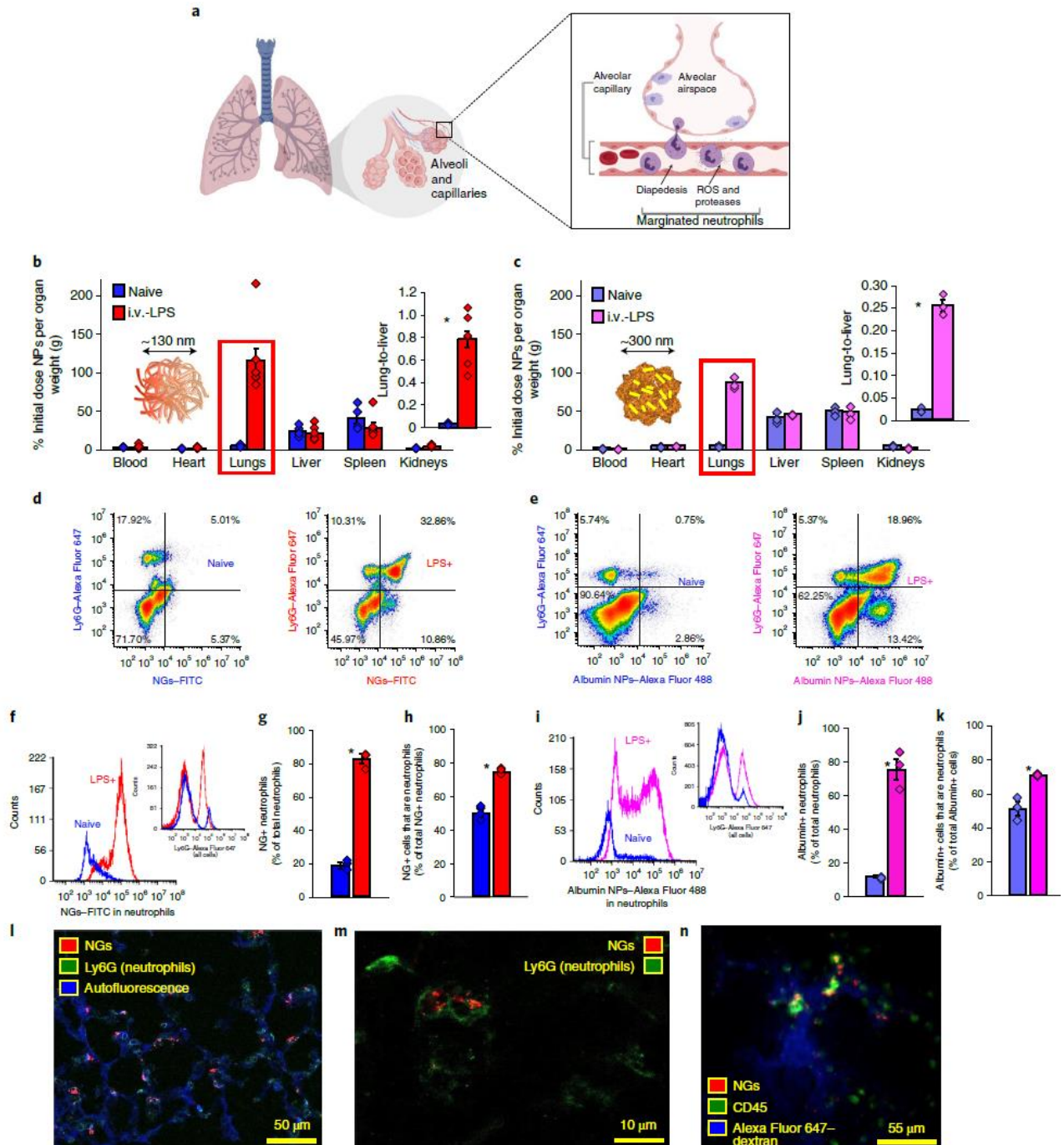
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5<sup>TH</sup> KDDF GLOBAL  
C&D TECH FAIR

## ► Key Data

Lysozyme–dextran NGs and crosslinked albumin NPs accumulate in margined neutrophils in inflamed lungs



To be continued

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## ► Key Data

### Lysozyme–dextran NGs and crosslinked albumin NPs accumulate in margined neutrophils in inflamed lungs

**a**, Schematic of neutrophil margination and extravasation in inflamed lungs (created with [BioRender.com](https://www.biorender.com)). ROS, reactive oxygen species. **b**, Biodistributions of lysozyme–dextran NGs in naive ( $n = 4$  animals) and i.v.-LPS-affected ( $n = 8$  animals) male C57BL/6 mice (red box,  $P < 1 \times 10^{-10}$ ;  $*P = 0.00008$ ). Inset: ratio of nanoparticle uptake in the lungs to nanoparticle uptake in the liver. **c**, Biodistributions of PEG–*N*-hydroxysuccinimide crosslinked human albumin NPs in naive ( $n = 3$  animals) and i.v.-LPS-injured ( $n = 3$  animals) mice (red box,  $P < 1 \times 10^{-10}$ ;  $*P = 0.004$ ). Inset: ratio of nanoparticle uptake in the lungs to nanoparticle uptake in the liver. **d–k**, Flow cytometry characterization of single-cell suspensions prepared from naive and i.v.-LPS-affected mouse lungs. Vertical axis in **d** and **e** indicates Ly6G staining for neutrophils and horizontal axis indicates signal from fluorescent NGs (**d**) or fluorescent albumin NPs (**e**). NG (**f**) and albumin NP (**i**) fluorescent signal from neutrophils in i.v.-LPS-injured mouse lungs (red/pink), compared to naive lungs (blue). Insets in **f** and **i**: flow cytometry data verifying increased neutrophil concentration in i.v.-LPS-injured mouse lungs (red/pink). Fraction of neutrophils positive for NGs (**g**) or albumin NPs (**j**) in naive or i.v.-LPS-injured lungs and fraction of NG-positive (**h**) or albumin NP-positive (**k**) cells that are neutrophils. For **g** and **h**, NGs/naive:  $n = 4$  animals, NGs/LPS:  $n = 4$  animals. For **j** and **k**, albumin NPs/naive:  $n = 3$  animals, albumin NPs/LPS:  $n = 3$  animals.  $*P = 2.6 \times 10^{-7}$  (**g**),  $*P = 1.7 \times 10^{-5}$  (**h**),  $*P = 0.0006$  (**j**),  $*P = 0.007$  (**k**). For **l** and **m**, fluorescence micrographs indicating association of NGs (red) with neutrophils (green, Ly6G stain) in the lungs of an i.v.-LPS-affected mouse (blue, tissue autofluorescence). Data are from histology for two naive mice and two i.v.-LPS-affected mice. **l**, Broad field of view indicating neutrophils and NGs alongside lung anatomy. **m**, Narrow field of view showing two neutrophils containing NGs. **n**, Single frame from real-time intravital imaging of NG (red) uptake in leukocytes (green) in the lungs of one i.v.-LPS-affected mouse (blue, Alexa Fluor 647-dextran). Statistical significance in **b** and **c** is derived from two-way analysis of variance (ANOVA) with Sidak's multiple-comparisons test. Statistical significance in **g,h,j,k** is derived from paired two-tailed  $t$  tests. All error bars indicate mean  $\pm$  s.e.m.

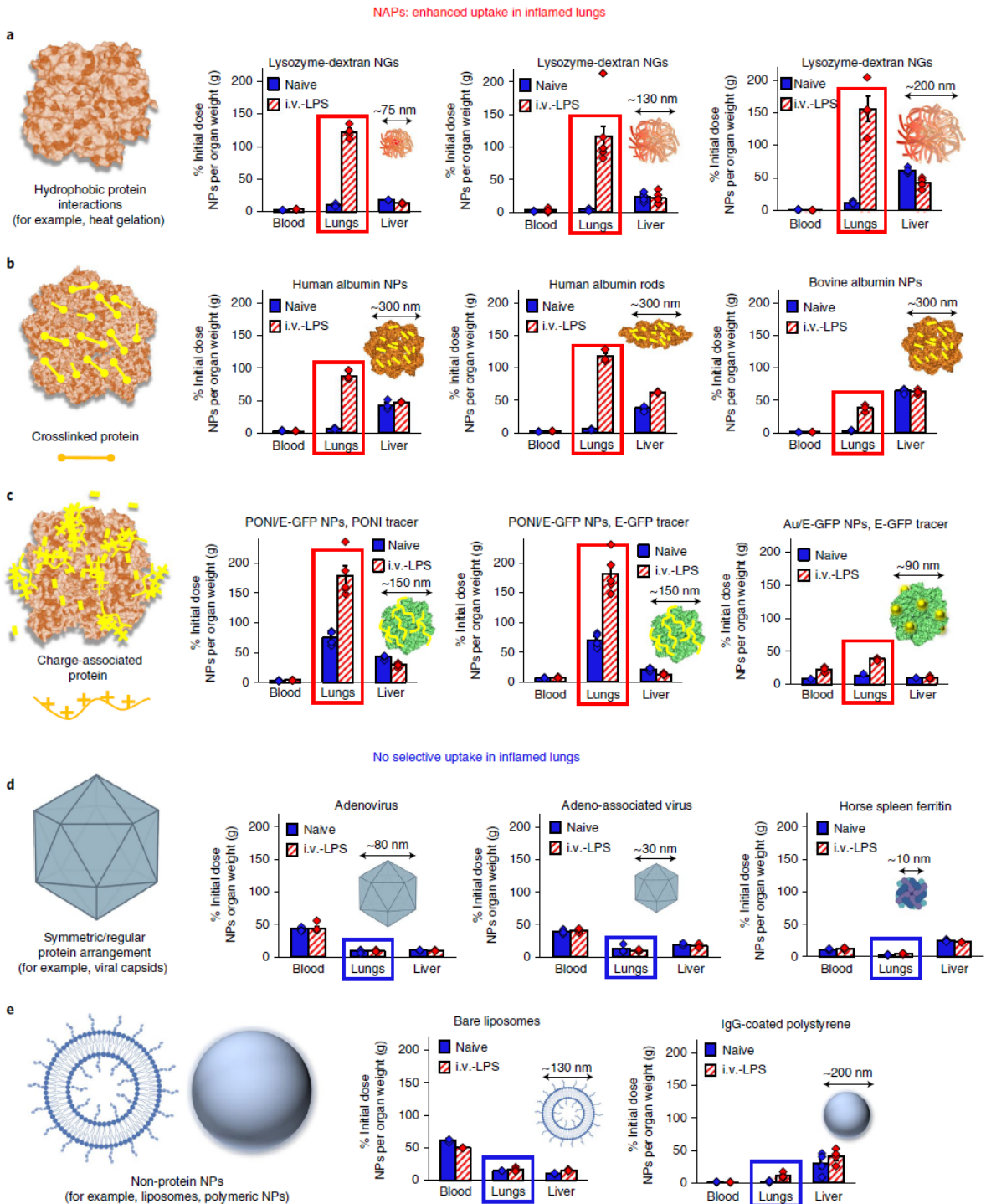
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## Key Data

### Screen of diverse NP biodistributions in naive and i.v.-LPS-affected lungs



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## ► Key Data

### Screen of diverse NP biodistributions in naive and i.v.-LPS-affected lungs

**a–c**, NAPs accumulate in acutely inflamed lungs. **a**, Biodistributions of variant NGs indicating uptake of 75 nm NGs ( $n = 4$  i.v.-LPS animals,  $n = 4$  naive animals, red box:  $P < 1 \times 10^{-10}$ ) and 200 nm NGs ( $n = 5$  i.v.-LPS,  $n = 5$  naive, red box:  $P < 1 \times 10^{-10}$ ) in LPS-injured lungs, but not naive lungs. Data for 130 nm NGs are identical to that presented in Fig. 1b. **b**, Biodistributions of variant crosslinked albumin NPs indicating uptake of albumin nanorods ( $n = 3$  i.v.-LPS animals,  $n = 3$  naive animals; red box,  $P < 1 \times 10^{-10}$ ) and bovine albumin NPs ( $n = 3$  i.v.-LPS animals,  $n = 3$  naive animals; red box,  $P < 1 \times 10^{-10}$ ) in LPS-injured, but not naive lungs. Data for human albumin nanoparticles are identical to that presented in Fig. 1c. **c**, Biodistributions of charge-agglutinated protein NPs, indicating uptake of particles comprised of E-GFP and guanidine-tagged PONI or particles comprised of E-GFP and guanidine-tagged gold nanoparticles in LPS-injured (PONI:  $n = 5$  animals; Au:  $n = 3$  animals), but not naive (PONI:  $n = 4$  animals; Au:  $n = 3$  animals) lungs. PONI/E-GFP data reflect tracing of both  $^{131}\text{I}$ -labelled PONI and  $^{125}\text{I}$ -labelled E-GFP. For PONI tracer data: red box,  $P < 1 \times 10^{-10}$ . For E-GFP tracer data: red box,  $P = 0.0003$ . For Au/E-GFP data: red box,  $P = 1.6 \times 10^{-9}$ . **d**, NPs based on symmetric supramolecular arrangement of protein do not have tropism for inflamed lungs (schematics created with [BioRender.com](https://www.biorender.com)). Biodistributions of adenovirus ( $n = 5$  i.v.-LPS animals,  $n = 5$  naive animals; blue box,  $P = 0.88$ ), adeno-associated virus ( $n = 3$  i.v.-LPS animals,  $n = 3$  naive animals; blue box,  $P = 0.56$ ) and ferritin nanocages ( $n = 5$  i.v.-LPS animals,  $n = 5$  naive animals; blue box,  $P = 0.35$ ) indicating no selectivity for LPS-injured versus naive lungs. **e**, Biodistributions of bare liposomes (schematic created with [BioRender.com](https://www.biorender.com),  $n = 4$  i.v.-LPS animals,  $n = 4$  naive animals) indicating no selectivity for LPS-injured versus naive lungs (blue box,  $P = 0.31$ ). Biodistributions of IgG-coated polystyrene NPs indicating low levels of uptake in both naive ( $n = 4$  animals) and LPS-injured ( $n = 4$  animals) lungs (blue box,  $P = 0.0004$ ). Statistical significance in all panels is derived from two-way ANOVA with Sidak's multiple comparisons test. All error bars indicate mean  $\pm$  s.e.m.

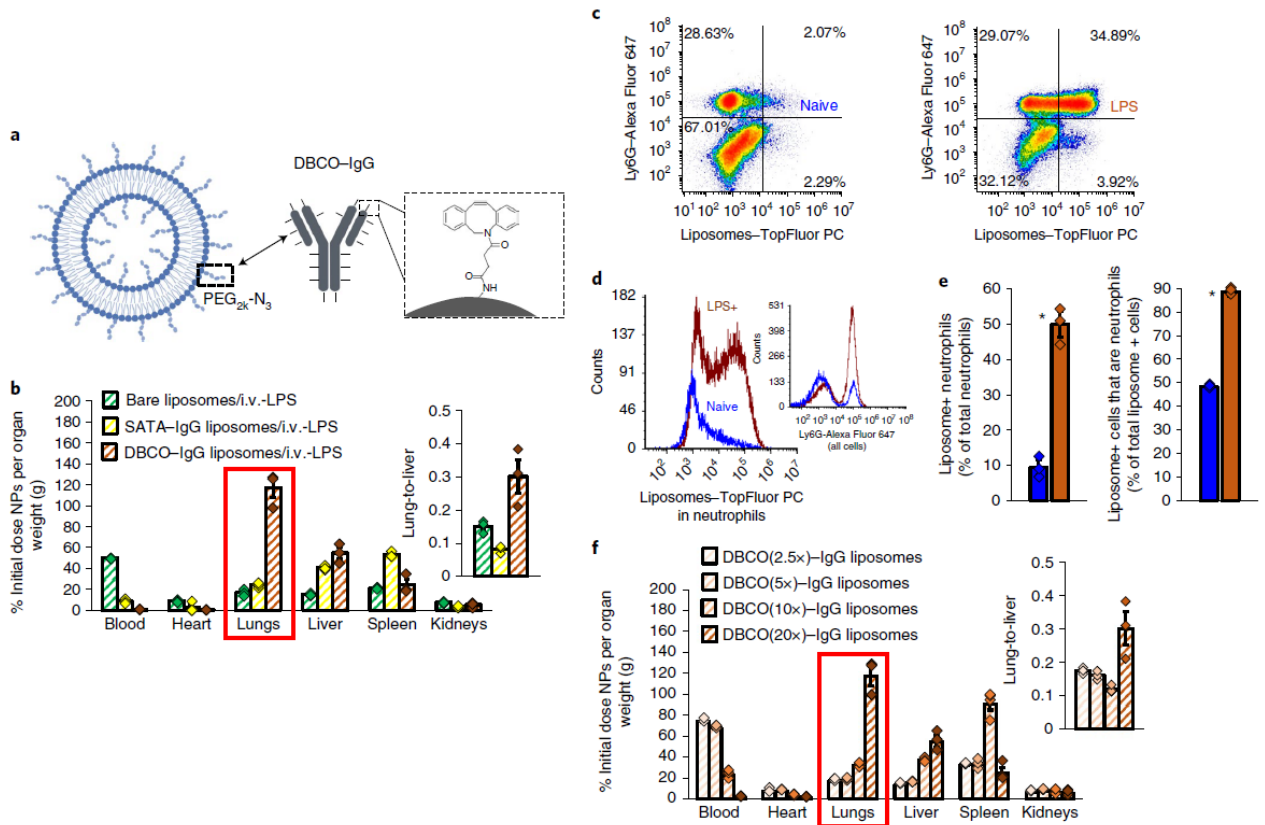
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## ► Key Data

### Engineering of liposome surface chemistry to confer NAP-like behaviour in LPS-inflamed lungs



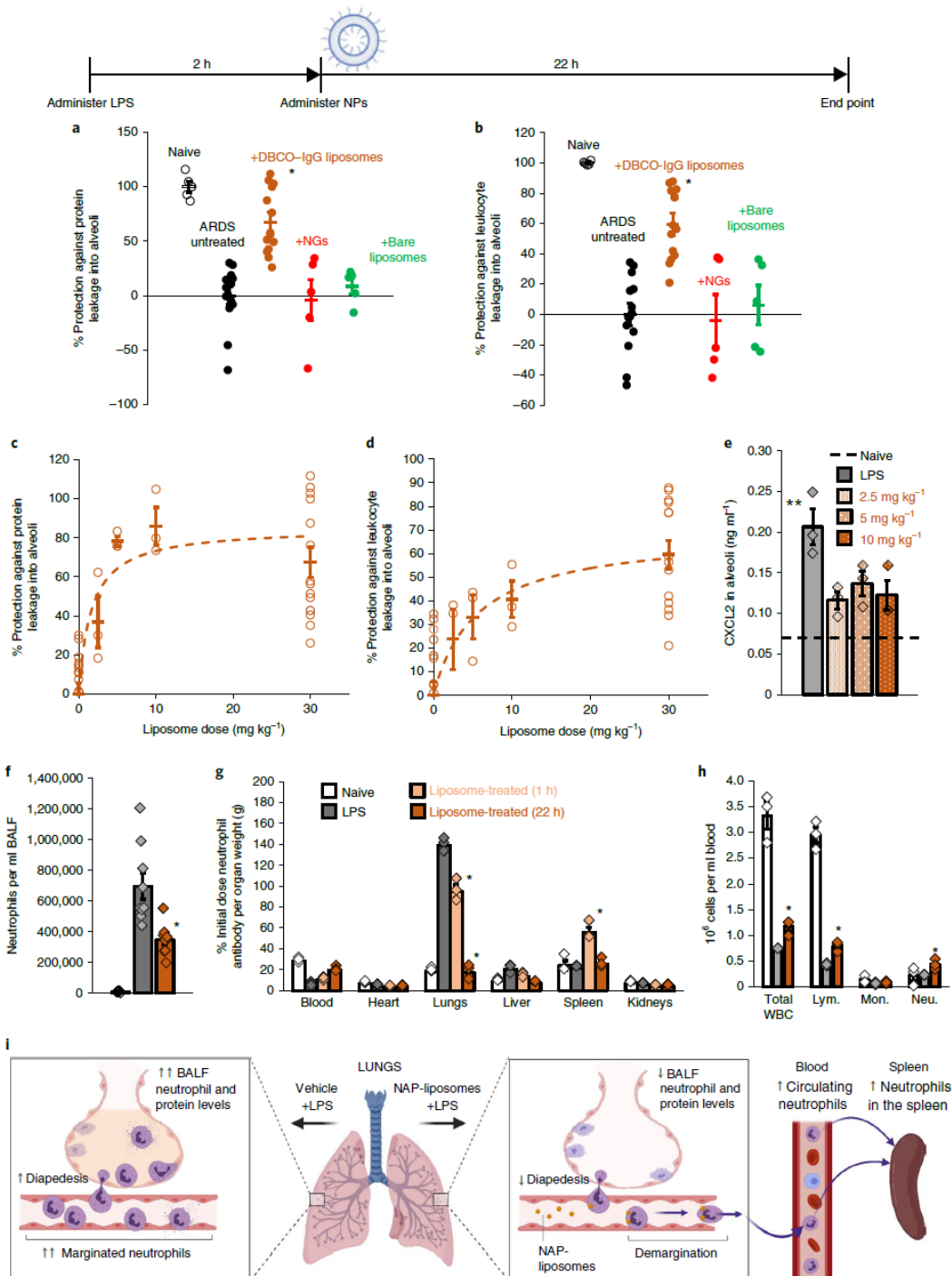
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### Effects of NAPs in model ARDS



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### Effects of NAPs in model ARDS

Timeline: nanoparticles or vehicle were administered as an i.v. bolus 2 h after nebulized LPS administration (liposome schematic created with [BioRender.com](#)). **a,b**, BALF was harvested 22 h after nanoparticle (30 mg kg<sup>-1</sup>) or vehicle administration. **a**, Protein concentration in BALF, reflecting quantity of oedema in naive mice ( $n = 5$  animals), sham-treated mice with model ARDS ( $n = 15$  animals) and mice with model ARDS treated with DBCO-IgG liposomes ( $n = 14$  animals), NGs ( $n = 5$  animals) or bare PEGylated liposomes ( $n = 5$  animals).  $*P = 6.6 \times 10^{-7}$ , 0.0001 and 0.002 for comparison of DBCO-IgG liposome treatment with sham treatment, NG treatment and bare liposome treatment, respectively. **b**, Concentration of leukocytes in BALF for same groups as in **a**.  $*P = 1.1 \times 10^{-6}$ , 0.0001 and 0.002 for comparison of DBCO-IgG liposome treatment with sham treatment, NG treatment and bare liposome treatment, respectively. Quantities in **a** and **b** are represented as degree of protection against infiltration into alveoli, extrapolated from levels in naive mice (100% protection) and untreated mice with model ARDS (0% protection). **c,d**, Dose-response for oedema (**c**) and leukocyte infiltration (**d**) in alveoli of ARDS mice treated with DBCO-IgG liposomes. Data were obtained as in **a** and **b**, but with different liposome doses ( $n = 3$  animals for 2.5, 5 and 10 mg kg<sup>-1</sup> liposome doses). **e**, Chemokine CXCL2 levels in alveoli of LPS-injured mice with and without DBCO-IgG liposome treatment ( $n = 3$  animals for all groups). Dashed line indicates CXCL2 levels in alveoli of naive mice.  $**P = 0.024$ , 0.079 and 0.034 for comparison of sham treatment with 2.5, 5 and 10 mg kg<sup>-1</sup> DBCO-IgG liposome treatment, respectively. **f**, Concentration of neutrophils in BALF of naive mice ( $n = 5$  animals), mice with model ARDS ( $n = 9$  animals) and mice with model ARDS dosed with 30 mg kg<sup>-1</sup> DBCO-IgG liposomes ( $n = 9$  animals). For comparison of DBCO-IgG liposome treatment to sham treatment,  $*P = 0.009$ . **g**, Biodistributions of anti-Ly6G antibody in naive mice ( $n = 3$  animals), LPS-injured mice ( $n = 3$  animals) and mice treated with 10 mg kg<sup>-1</sup> DBCO-IgG liposomes, with organs sampled at 1 h after treatment ( $n = 3$  animals) or 22 h after treatment ( $n = 3$  animals). Naive and untreated LPS-affected data are identical to data in Supplementary Fig. 1a.  $*P < 1 \times 10^{-10}$  for all comparisons of anti-Ly6G uptake in lungs or spleen of liposome-treated mice versus sham-treated mice. **h**, Complete blood count analysis of circulating leukocyte concentrations in naive mice ( $n = 3$  animals), LPS-injured mice ( $n = 3$  animals) and mice treated with 10 mg kg<sup>-1</sup> DBCO-IgG liposomes, with blood sampled 22 h after treatment ( $n = 3$  animals).  $*P = 0.019$ , 0.025 and 0.047 for comparison of DBCO-IgG liposome-treated to sham-treated values for total white blood cell (WBC), lymphocyte (Lym.) and neutrophil (Neu.) counts, respectively. **i**, Schematic for the fate of neutrophils in mice with model ARDS, with and without DBCO-IgG liposome treatment, based on data in **f-h** (created with [BioRender.com](#)). Statistical significance in **a**, **b**, **e** and **f** is derived from one-way ANOVA with Tukey's multiple-comparisons test. Statistical significance in **g** and **h** is derived from two-way ANOVA with Tukey's multiple-comparisons test. All error bars indicate mean  $\pm$  s.e.m.