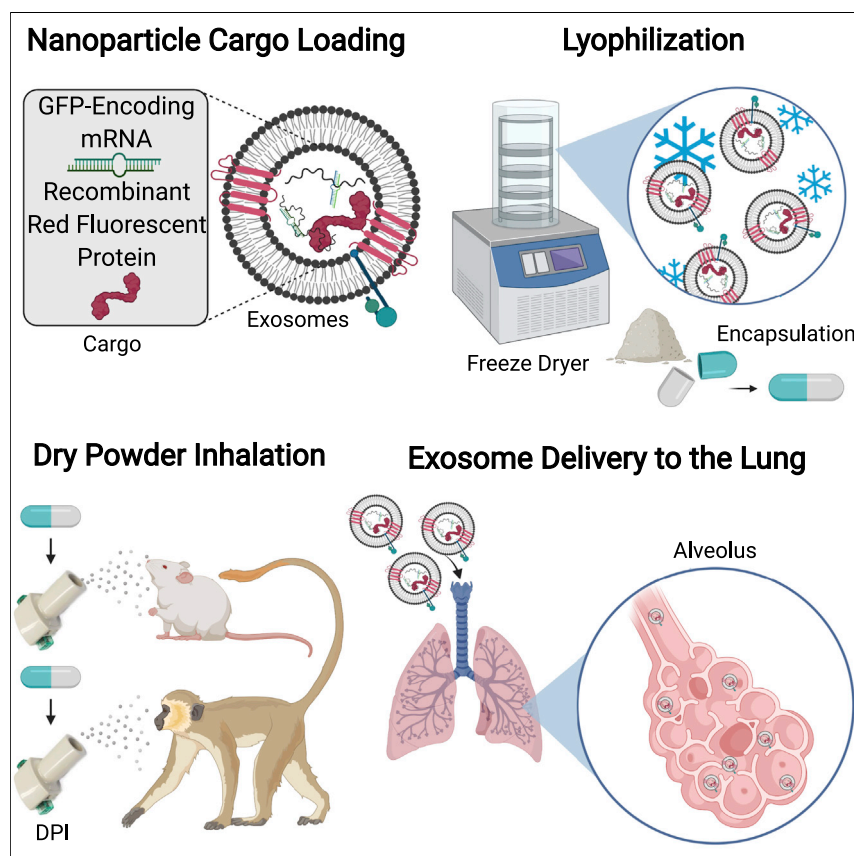


## Article

## Inhalable dry powder mRNA vaccines based on extracellular vesicles



Lipid nanoparticles have limitations in inhaled drug delivery, including low pulmonary bioavailability and unoptimized formulation. Lung-derived extracellular vesicles (Lung-Exos) may be naturally equipped for drug delivery to the lung. We determined the biodistribution of Lung-Exos following nebulization and dry powder inhalation, where Lung-Exos outperformed their biological and synthetic nanoparticle counterparts in drug distribution and retention. As an inhalable vaccine, Lung-Exos elicited greater protective antibody responses and pseudoviral clearance than their synthetic counterpart.

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**Highlights**

Lung extracellular vesicles (Lung-Exos) can package mRNA and protein drugs

Lung-Exos are deliverable through nebulization and dry powder inhalation

Dry powder Lung-Exos are room-temperature stable up to 28 days

Drug-loaded Lung-Exos can serve as an inhalable vaccine to illicit immune responses

**Development**

Practical, real world, technological considerations and constraints

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## Article

# Inhalable dry powder mRNA vaccines based on extracellular vesicles

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## SUMMARY

Respiratory diseases are a global burden, with millions of deaths attributed to pulmonary illnesses and dysfunctions. Therapeutics have been developed, but they present major limitations regarding pulmonary bioavailability and product stability. To circumvent such limitations, we developed room-temperature-stable inhalable lung-derived extracellular vesicles or exosomes (Lung-Exos) as mRNA and protein drug carriers. Compared with standard synthetic nanoparticle liposomes (Lipos), Lung-Exos exhibited superior distribution to the bronchioles and parenchyma and are deliverable to the lungs of rodents and nonhuman primates (NHPs) by dry powder inhalation. In a vaccine application, severe acute respiratory coronavirus 2 (SARS-CoV-2) spike (S) protein encoding mRNA-loaded Lung-Exos (S-Exos) elicited greater immunoglobulin G (IgG) and secretory IgA (SIgA) responses than its loaded liposome (S-Lipo) counterpart. Importantly, S-Exos remained functional at room-temperature storage for one month. Our results suggest that extracellular vesicles can serve as an inhaled mRNA drug-delivery system that is superior to synthetic liposomes.

## INTRODUCTION

Respiratory diseases are among the leading causes of morbidity and mortality worldwide,<sup>1</sup> with coronavirus disease 2019 (COVID-19)<sup>2</sup> remaining prevalent in the ongoing pandemic. A wide range of therapeutics have been developed and repurposed to treat respiratory diseases, including small-molecule drugs,<sup>3,4</sup> stem cells<sup>5–9</sup> and their derivatives,<sup>10–13</sup> and, notably, messenger RNA (mRNA)<sup>14–17</sup> and protein vaccines.<sup>18,19</sup> Currently, three intramuscular (IM) vaccines (Pfizer-BioNTech, Moderna, and Janssen) have been authorized for COVID-19 by the US Food and Drug Administration. However, formulating drugs and vaccines for inhaled delivery may improve their efficacy and bioavailability in the lung as well as improve patient compliance.

Although inhalation delivery is attractive due to its local, noninvasive, and highly absorptive properties, drug formulation with optimized physicochemical parameters remains the key obstacle.<sup>20</sup> A possible solution is colloidal drug-delivery systems, which help overcome poor drug solubility and hydrophobicity by providing protection through nanoparticle encapsulation.<sup>21</sup> Plus, nanoparticles retained in the lung prolong drug release in the highly vascularized pulmonary and bronchial circulations.<sup>22</sup> Lipid nanoparticles (LNPs) have been successful as drug-delivery vesicles

## PROGRESS AND POTENTIAL STATEMENT

Research in extracellular vesicles (EVs) is important to the field of translational medicine to develop therapeutics that are limited by poor cellular targeting and efficacy. The biological composition of EVs can be exploited as drug-delivery vehicles that may be engineered for cellular targeting or eliciting specific immune responses through their functions in membrane trafficking and cellular signaling. With the molecular composition of EVs varying depending on their parent-cell origin, the derivation of EVs can further refine nanomedicine by utilizing nanoparticles that are recognized by specific cellular microenvironments. EVs are found in almost all biological fluids, opening the application of EVs as tailored drug-delivery vesicles to a wide range of diseases.

for vaccines, most notably in the liposome-encapsulated COVID-19 mRNA IM vaccines.<sup>23</sup> However, major limitations such as deep-freezing storage and healthcare professional administration have led to the expansion of the LNP platform to alternative delivery routes such as self-administered inhalation. In addition, the lung has sophisticated pulmonary defense mechanisms and surfactants that protect it against inhaled particulates and microbes.<sup>22,24</sup> Therefore, the formulation of nanoparticle drug-delivery systems must be optimized to overcome these inherent obstacles.

Exosomes, biologically derived nanoparticles, may be naturally equipped to withstand pulmonary conditions. They are nanosized extracellular vesicles secreted by numerous cell types and found in almost all biological fluids.<sup>25</sup> Initially regarded as cellular debris, exosomes are now understood to have potent roles in autocrine and paracrine signaling.<sup>26–28</sup> Originating from the endosomal system and shedding from the plasma membrane, exosomes contain various cocktails of RNA, protein, and lipid cargo with unique parent-cell signatures.<sup>29,30</sup> Lung-derived exosomes can be utilized as sophisticated drug-delivery systems that offer cargo components and membrane features tailored to the lung microenvironment.<sup>20,31</sup> Our group has demonstrated the lung regenerative abilities of human lung spheroid cells (LSCs)<sup>8,9</sup> in rodent models of idiopathic pulmonary fibrosis (IPF), and their safety and efficacy are being tested in a human clinical trial (HALT-IPF, ClinicalTrials.gov: NCT04262167). LSCs and their secreted exosomes (lung-derived extracellular vesicles [Lung-Exos]) have regenerative abilities in IPF models<sup>13</sup> and protective abilities against COVID-19 as decoys.<sup>32</sup> In both disease models, Lung-Exos maintained therapeutic efficacy through jet-nebulization administration, demonstrating the ability of Lung-Exos to function as an inhalable drug-delivery and vaccine vehicle. Additionally, exosomes can be synthetically supplemented to enhance cellular targeting and therapeutic efficacy.<sup>33–35</sup> For example, anti-inflammatory peptides<sup>36</sup> can be linked onto the exosomal membrane, engineering an immune cell targeting nanomedicine. The combination of vesicle derivation and supplementation allows for a customizable nanoparticle delivery platform that can be utilized across many lung diseases.

Although exosomes as inhaled therapeutics and RNA delivery vehicles have been demonstrated,<sup>13,37</sup> the distribution and retention of exosome particles in the lung have yet to be determined. Drug effectiveness depends on proper deposition of particles within the respirable fraction, requiring optimized nanoparticle formulation. In this study, we sought to elucidate the biodistribution of lung-derived exosomes upon nebulization, baselined to liposomes (Lipos) as a commercial standard. Furthermore, to provide a room-temperature-stable product, we formulated exosomes as a lyophilized dry powder to investigate their stability and inhaled biodistribution in the lung of both the mouse and African green monkey (AGM). The parent-cell signature of Lung-Exos may suggest that they are naturally optimized for the distribution and retention within the lung, which may allow them to bypass pulmonary clearance more efficiently than Lipos or exosomes derived from other cell types. Through this enhanced pulmonary bioavailability, we hypothesize that lung-derived exosomes elicit greater therapeutic responses for pulmonary diseases and serve as a customizable drug-delivery vehicle for room-temperature-stable inhaled mRNA therapeutics.

## RESULTS

### Exosome distribution in the bronchioles and parenchyma are superior to that of synthetic nanoparticles

Red fluorescent protein (RFP)-labeled lung-derived exosomes (RFP-Exos) and Lipos (RFP-Lipos) were fabricated to generate trackable nanoparticles for biodistribution

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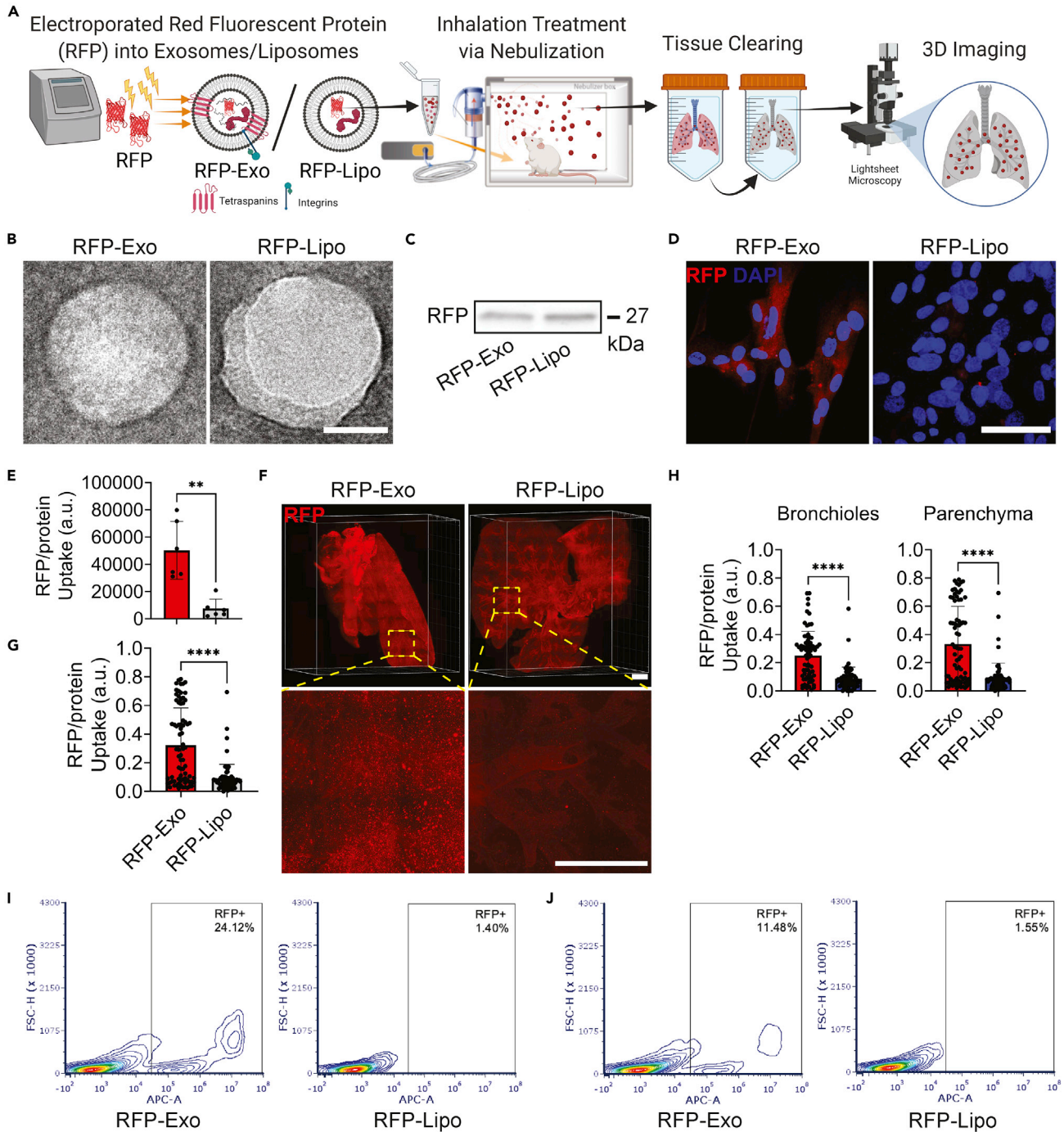
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**Figure 1. Fabrication and distribution of exosomes and liposomes**

(A) Schematic showing protein loading into lung-derived exosomes (RFP-Exos) and liposomes (RFP-Lipos), nebulization administration, ex vivo lung-tissue clearing, and 3D imaging by LSFM. Created with [BioRender.com](https://www.biorender.com).

(B) TEM images of RFP-Exos and RFP-Lipos; scale bar: 50 nm.

(C) Immunoblot of RFP in exosome and liposome lysate.

(D) Representative immunostaining images of lung parenchymal cells for RFP (red) and DAPI (blue); scale bar: 50  $\mu$ m.

(E) Quantification of RFP-Exo and RFP-Lipo pixel intensity normalized to nuclei in lung parenchymal cell images; n = 6 per group; data are represented as mean  $\pm$  standard deviation.

(F) LSFM images of cleared mouse lungs after RFP-Exo and RFP-Lipo nebulization; scale bar: 1,000  $\mu$ m.

**Figure 1. Continued**

(G) Quantification of the integrated density of RFP normalized to the whole-lung area; n = 74 total slices from two biological replicates per group; data are represented as mean  $\pm$  standard deviation.

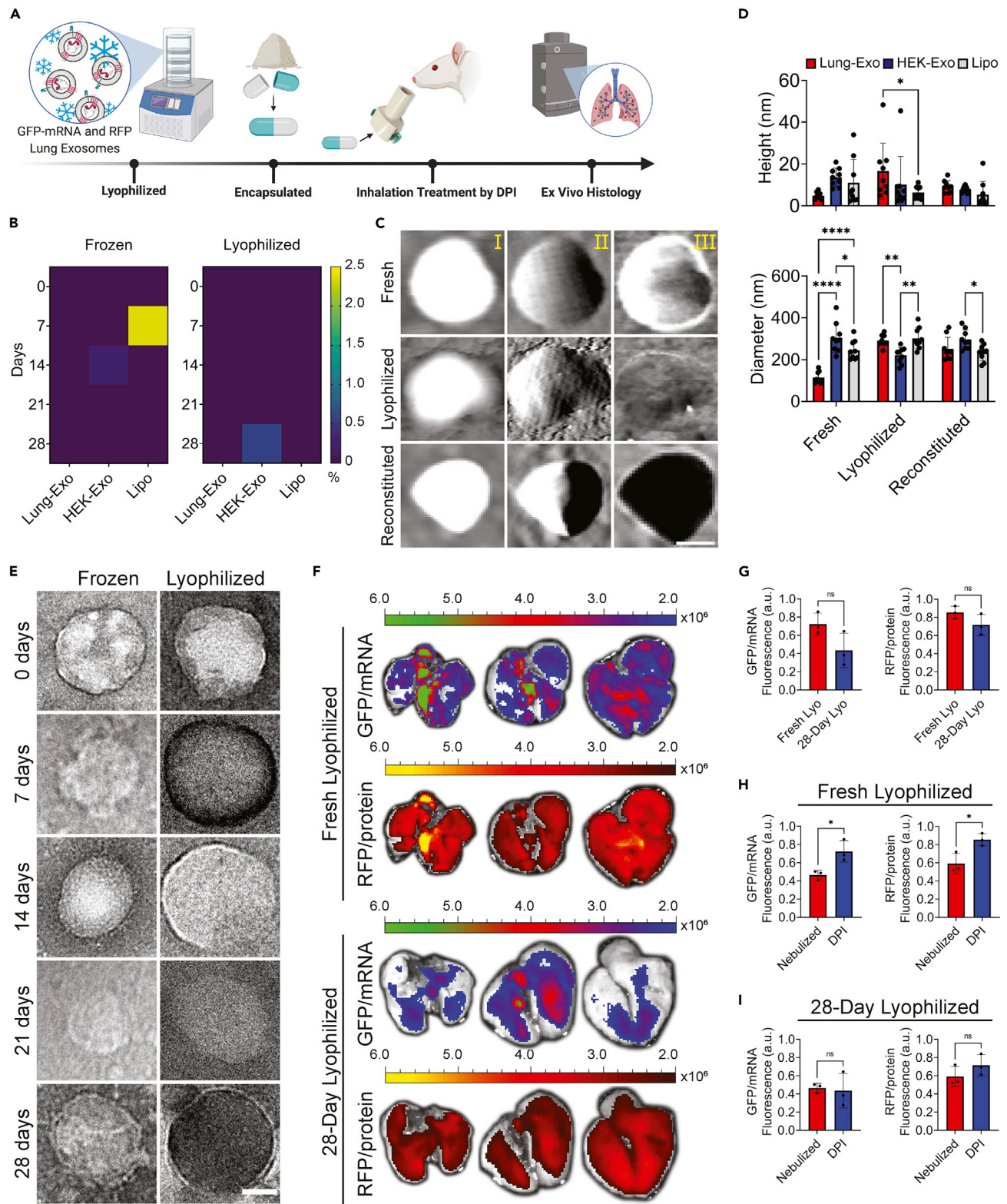
(H) Quantification of the integrated density of RFP normalized to segmented bronchiole and parenchymal regions from whole-lung images; n = 74 total slices from two biological replicates per group; data are represented as mean  $\pm$  standard deviation.

(I and J) Flow cytometry analysis of lung parenchymal cells co-cultured with RFP-Exos or RFP-Lipos (I) and murine lung cells that received nebulized RFP-Exos or RFP-Lipos (J).

analysis in the murine lung after inhalation treatment through three-dimensional (3D) imaging (Figure 1A). The nanoparticles were characterized by transmission electron microscopy (TEM), confirming that the isolation of exosomes and Lipos did not disrupt vesicular membrane integrity (Figure 1B). RFP loading was verified by immunoblotting (Figure 1C). When co-cultured with lung parenchymal cells, RFP-Exo had a 6.7-fold increase in cellular uptake and RFP protein expression compared with cells cultured with RFP-Lipo (Figures 1D and 1E). Next, the biodistribution of nanoparticles *in vivo* were evaluated through light-sheet fluorescence microscopy (LSFM) (Figure 1F). Healthy mice received a single dose of RFP-Exos or RFP-Lipos via nebulization and were sacrificed after 24 h. LSFM imaging confirmed nanoparticle delivery to the conducting airways and the deep lung, with an accumulation of RFP-Exos in the upper pulmonary regions (Videos S1 and S2). Quantification of nanoparticle delivery to the whole lung demonstrated a 3.7-fold improvement in RFP-Exo retention and uptake compared with RFP-Lipo (Figure 1G). Segmentation of the lung into bronchial and parenchymal regions revealed 2.9- and 3.8-fold improvements in RFP-Exo retention and uptake, respectively, compared with RFP-Lipo (Figure 1H). Flow cytometry analysis in lung parenchymal cells (Figure 1I) and in the murine lung following nebulization (Figure 1J) confirmed greater cellular uptake of RFP-Exos than RFP-Lipos. The drug-loading capabilities of lung-derived exosomes (Lung-Exos and Lipos) were expanded by loading GFP-encoding mRNA to evaluate nanoparticle mRNA uptake. Lung parenchymal cells that received GFP-Exos demonstrated more rapid internalization of exosomal mRNA than liposomal mRNA (Figure S1). These data confirm that our nanoparticle labeling system maintains nanoparticle integrity while delivering functional and translatable cargo after jet nebulization. *In vitro* and *in vivo* analyses suggest superior retention and cellular uptake of exosomes over Lipos in the lung. The native lung signature of lung-derived exosomes may enhance pulmonary bioavailability, resulting in an optimized nanoparticle vesicle for drug delivery for respiratory diseases.

**Lung-derived exosomes efficiently penetrate mucus**

Delivery of inhaled therapeutics must penetrate the lung's protective mucus lining to provide pulmonary bioavailability. Lung-Exos were compared against human embryonic kidney (HEK)-derived exosomes (HEK-Exos) and Lipos to determine if nanoparticle derivation affected mucus penetrance. To test this, we used a model of the human airway at the air-liquid interface (Figure S2A), with human mucus-secreting bronchial epithelial cells lining the transwell membrane and human lung parenchymal cells lining the well (Figure S2B). Immunostaining confirmed the mucus lining in the transwell membrane and delivery of DiD-labeled nanoparticles (Figure S2C). Quantification of nanoparticle penetrance into the wells revealed the greatest uptake of Lung-Exos (Figure S2D), with the highest percentage of cellular uptake by lung parenchymal cells (Figure S2E) by 24 h. Likewise, Lung-Exos had the least entrapment by the mucus-lined membrane (Figure S2F) and the lowest percentage of cellular uptake by bronchial epithelial cells (Figure S2G). These data confirm mucus penetrance of the nanoparticles and suggest that Lung-Exos can most efficiently evade mucoadhesion, overcoming the lung's natural defense mechanism and allowing for greater parenchymal bioavailability.



**Figure 2. Stability and distribution of lung-derived exosomes in dry powder formulation in the murine lung**

(A) Schematic of mRNA and protein-loaded lung-derived exosome lyophilization, encapsulation, rodent DPI administration, and ex vivo histology. Created with [BioRender.com](https://www.biorender.com).

**Figure 2. Continued**

(B) Heatmaps of RFP leakage from Lung-Exos, HEK-Exos, and Lipos detected by ELISA;  $n = 2$  per group.

(C) Representative AFM height (I), amplitude (II), and phase (III) images of Lung-Exos; scale bar: 50 nm.

(D) Quantification of the height and diameter of Lung-Exos, HEK-Exos, and Lipos from AFM images;  $n = 9$  per group; data are represented as mean  $\pm$  standard deviation.

(E) TEM images of Lung-Exos at frozen (Frozen) or room (Lyophilized) temperatures; scale bar: 50 nm.

(F) *Ex vivo* images of mouse lungs that received fresh lyophilized (0 days) and 28-day-old lyophilized Lung-Exos via dry powder inhalation after 24 h.

(G) Quantification of the integrated density of GFP and RFP fluorescence in *ex vivo* mouse lungs 24 h after fresh (Fresh-Lyos) and 28-day-old (28-Day Lyos) dry powder inhalation;  $n = 3$  per group; data are represented as mean  $\pm$  standard deviation.

(H) Quantification of the integrated density of GFP and RFP fluorescence in *ex vivo* mouse lungs 24 h after nebulization and fresh (Fresh-Lyos) dry powder inhalation;  $n = 3$  per group; data are represented as mean  $\pm$  standard deviation.

(I) Quantification of the integrated density of GFP and RFP fluorescence in *ex vivo* mouse lungs 24 h after nebulization and 28-day-old (28-Day Lyos) dry powder inhalation;  $n = 3$  per group; data are represented as mean  $\pm$  standard deviation.

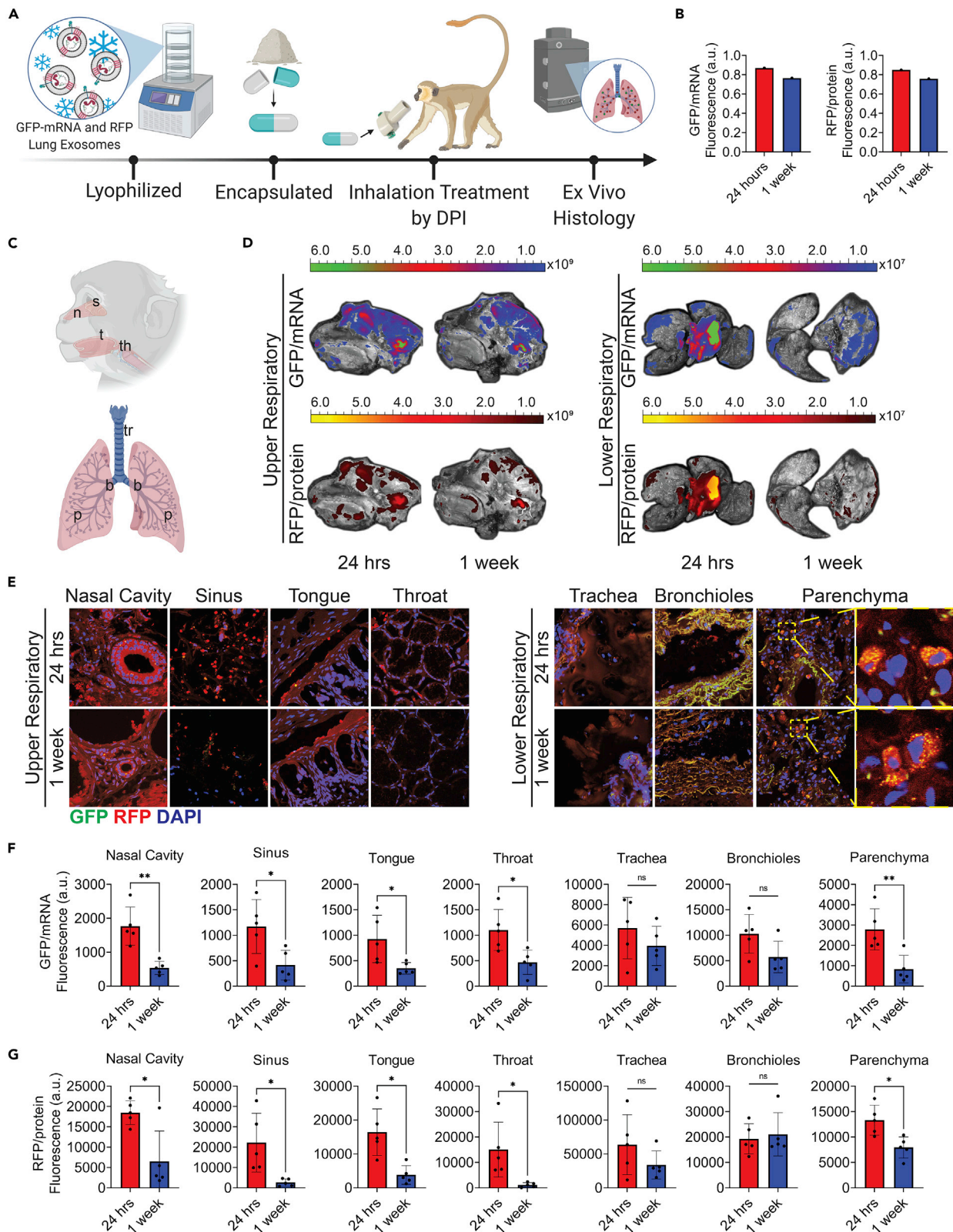
**Lung-derived exosomes are room-temperature stable and distributable in dry powder formulation in the murine lung**

Room-temperature formulation of therapeutics circumvents major limitations in traditional IM vaccine delivery: deep-freezing storage, healthcare professional administration, and reduced patient compliance. Therefore, we reformulated our liquid nanoparticle suspensions into dry, lyophilized powder for dry powder inhalation (DPI) administration. We verified the efficacy and stability of room-temperature lyophilized Lung-Exos up to 28 days in the murine lung (Figure 2A). To verify dry powder nanoparticle stability and shelf life, lyophilized nanoparticle cargo leakage was tested by an enzyme-linked immunosorbent assay (ELISA), where nanoparticles had less than 2.4% of total pg/mL cargo leakage at day 28 of room-temperature storage (Figures 2B and S3). Next, the morphology of nanoparticles was evaluated across their fresh and lyophilized formulations, as well as lyophilized powder reconstituted in water (reconstituted), to mimic rehydration of dry powder by saliva and mucus. TEM (Figures 2E and S4) and atomic force microscopy (AFM) verified that reformulation and rehydration did not affect nanoparticle membrane integrity (Figures 2C and S5) but did affect size distributions through clumping (Figure S6–S8). Lyophilization increased nanoparticle height and diameter (Figure 2D) but remained as small respiratory droplets upon reconstitution. Across all formulations, the nanoparticle diameters are approximately 10-fold larger, which may be explained by tip dilation<sup>38</sup> that reports larger lateral dimensions than 2D analysis such as through TEM.<sup>39</sup> Cross-section measurement curves demonstrate a restoration of membrane “smoothness” in reconstituted nanoparticles, mimicking fresh formulation (Figure S9). Next, we delivered the lyophilized Lung-Exos via DPI, where *ex vivo* images (Figure 2F) of mouse lungs who received fresh (fresh lyophilized) and 28-day-old (28-day lyophilized) dry powder Lung-Exos had no significant difference in exosomal mRNA and protein distribution (Figure 2G). mRNA activity showed greater variability at its 28-day-old state, but protein activity remained more stable.

Compared with nebulized exosome biodistribution, both fresh (Figure 2H) and 28-day-old (Figure 2I) lyophilized exosomes have trends of greater pulmonary distribution 24 h after administration. Lyophilized Lung-Exo is a room-temperature-stable exosome formulation that can deliver functional and translatable cargo via DPI.

**Distribution of exosomes via DPI in AGMs**

Dry powder inhalers offer an at-home or on-the-go electronic-free administration of therapeutics through a user-friendly device designed specifically for pulmonary disease treatment, providing local drug delivery and reducing side effects associated with IM and oral drugs. Lung-Exos, HEK-Exos, and Lipos were lyophilized and encapsulated into commercially available hydroxypropyl methylcellulose (HPMC) capsules, and particle-distribution analysis was performed on DPIs with a





**Figure 3. Distribution of lung-derived exosomes via dry powder inhalation in African green monkeys**

(A) Schematic of mRNA and protein-loaded lung-derived exosome lyophilization, encapsulation, non-human primate DPI administration, and *ex vivo* histology. Created with [BioRender.com](https://www.biorender.com).

(B) Quantification of the integrated density of GFP and RFP fluorescence in *ex vivo* primate lungs 24 h and 1 week after dry powder inhalation;  $n = 1$  per group.

(C) Schematic showing upper respiratory tissue sectioning for nasal (n), sinus (s), tongue (t), and throat (th) sections and lower respiratory tissue sectioning for tracheal (tr), bronchial (b), and parenchymal (p) sections. Created with [BioRender.com](https://www.biorender.com).

(D) *Ex vivo* images of primate head cross-sections and lungs 24 h and 1 week after lyophilized Lung-Exos via dry powder inhalation.

(E) Representative immunostaining images of nasal cavity, sinus, tongue, throat, trachea, bronchioles, and parenchyma sections for GFP (green), RFP (red), and DAPI (blue); scale bar: 100  $\mu\text{m}$  in representative images; scale bar: 1  $\mu\text{m}$  in parenchyma sections.

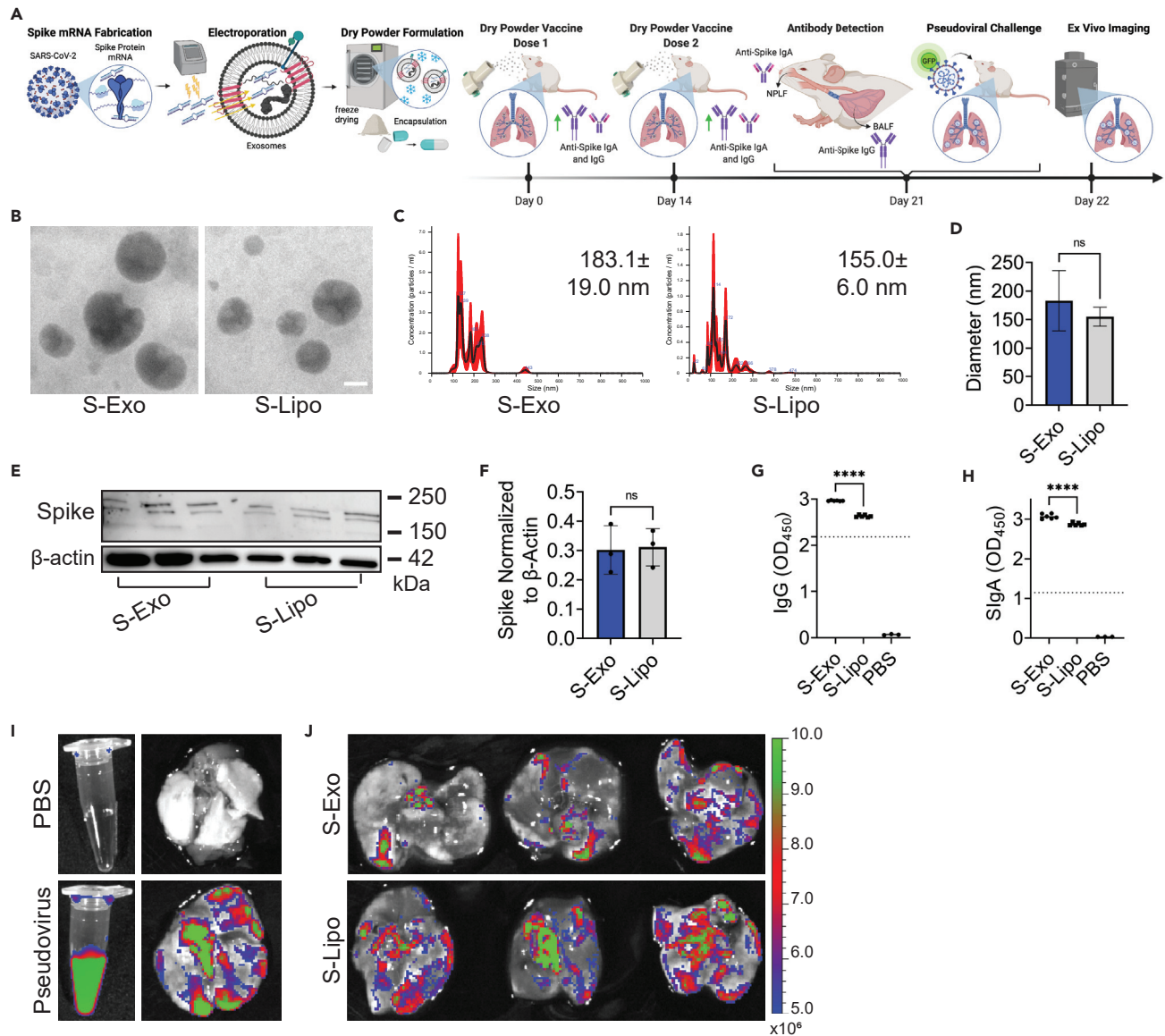
(F) Quantification of Lung-Exo GFP pixel intensity normalized to nuclei in nasal cavity, sinus, tongue, throat, trachea, bronchioles, and parenchyma sections;  $n = 5$  per group; data are represented as mean  $\pm$  standard deviation.

(G) Quantification of Lung-Exo RFP pixel intensity normalized to nuclei in nasal cavity, sinus, tongue, throat, trachea, bronchioles, and parenchyma sections;  $n = 5$  per group; data are represented as mean  $\pm$  standard deviation.

range of resistances from low to ultra high. A commercially available RS01 high-resistance DPI was selected for its overall greater output of aerosols within the respirable fraction (Figure S10)<sup>40</sup> and consistent particle distributions across the nanoparticles (Figure S11). Using this DPI, we tracked exosome biodistribution in the AGM by delivering a single dose of lyophilized Lung-Exos and sacrificing after 24 h and 1 week for further analysis (Figure 3A). AGMs receiving Lung-Exos were compared against a negative control AGM to evaluate autofluorescence. *Ex vivo* imaging revealed a similar biodistribution of exosomal mRNA and protein cargo throughout the lung (Figures 3B and 3D). Lung-Exos were maintained in the lung 1 week after administration (Figures 3B and 3D). Further immunostaining analysis of the upper and lower respiratory tracts confirmed exosome delivery (Figures 3C, 3E, and S12) and the greatest nanoparticle deposition into the lung (Figure S13). Exosomal mRNA (Figure 3F) and protein (Figure 3G) were significantly cleared from the upper respiratory and parenchymal regions after 1 week. Tracheal and bronchial regions maintained similar GFP and RFP fluorescence, which may be attributed to by the autofluorescent nature of airway tissue due to collagen, particularly in the fluorescein isothiocyanate (FITC) range (Figure S13).<sup>41,42</sup> Collagen fibers in the trachea and bronchioles are highly fluorescent in both the exosome-treated (Figure S12) and control (Figure S14) tissue sections and do not reflect exosomal mRNA and protein deposition. Overall, Lung-Exos are distributable in the simian upper and lower respiratory tracts through DPI.

**SARS-CoV-2 spike-loaded exosomes elicit antibody protection through DPI**

In our final study, the therapeutic potential of lyophilized Lung-Exos was tested as an inhaled vaccine against COVID-19. mRNA encoding the severe acute respiratory coronavirus 2 (SARS-CoV-2) spike (S) protein was loaded into Lung-Exos and Lipos and formulated for DPI, generating S-Exos and S-Lipos, respectively (Figure 4A). Although not naturally immunogenic, exosomes can be supplemented with agents such as mRNA to activate specific immune responses.<sup>43</sup> To confirm long-term storage efficiency, S-Exos and S-Lipos were stored for 1 month at room temperature prior to characterization and mouse vaccination. Nanoparticle vaccine integrity was verified by TEM (Figure 4B) and nanoparticle-tracking analysis (NTA) (Figures 4C and 4D). S-protein-encoding mRNA loading and *in vitro* cellular translation were verified by immunoblotting (Figure 4E), where S-Exos and S-Lipos had similar results (Figure 3F). Our inhaled vaccine was then tested in healthy mice, which received two doses of S-Exos or S-Lipos via DPI. The mice were sacrificed 1 week after the second dose, and bronchoalveolar lavage fluid (BALF) and nasopharyngeal lavage fluid (NPLF) were collected to assess anti-S immunoglobulin G (IgG) and secretory IgA (SIgA) antibody production, respectively (Figure 4A). ELISAs revealed that both S-Exos and S-Lipos produced



**Figure 4. Dry powder inhalation of S-protein-loaded lung-derived exosomes elicit greater immune responses than their synthetic counterpart** (A) Schematic of S protein mRNA loading into lung-derived exosomes, dry powder formulation, inhaled vaccine delivery doses, antibody production against SARS-CoV-2 spike protein, and pseudoviral challenge. Created with [BioRender.com](#).

(B) TEM images of S-Exos and S-Lipos at room temperature; scale bar: 50 nm.  
 (C) NTA size-distribution analysis.  
 (D) Quantification of NTA size-distribution analysis of the average mean  $\pm$  standard error of five replicates;  $n = 1$  per group.  
 (E) Immunoblots of S protein in mouse lung lysate.  
 (F) Quantification of immunoblots normalized to  $\beta$ -actin;  $n = 3$  per group; data are represented as mean  $\pm$  standard deviation.  
 (G) Anti-spike IgG antibody titer from murine BALF detected by ELISA;  $n = 6$  per group; data are represented as mean  $\pm$  standard deviation.  
 (H) Anti-spike SIgA antibody titer from murine NPLF detected by ELISA;  $n = 6$  per group; data are represented as mean  $\pm$  standard deviation.  
 (I) Ex vivo images of PBS or pseudovirus in solution (left) and in lungs 24 h after dry powder inhalation (right).  
 (J) Ex vivo images of S-Exo- or S-Lipo-vaccinated lungs 24 h after pseudoviral challenge.

sufficient IgG ( $>2.18325$ ; [Figure 4G](#)) and SIgA ( $>1.14595$ ; [Figure 4H](#)) antibodies to induce neutralizing antibody responses. However, DPI of S-Exos produced significantly higher amounts of antibodies than S-Lipos ([Figures 4G and 4H](#)). This suggests that S-Exos would have superior protection against SARS-CoV-2 infection over S-Lipos as an inhaled vaccine. Coupled with enhanced mucus

penetrance, S-Exos may further facilitate passive IgG diffusion into the pulmonary epithelial lining fluid.<sup>44</sup> To assess the ability to clear inhaled pseudovirus, mice vaccinated with two doses of S-Exos or S-Lipos were challenged with wild-type SARS-CoV-2 pseudoviral particles with a GFP reporter (Figure 4I). Twenty-four h after pseudoviral challenge, mice vaccinated with S-Exos showed significantly greater viral clearance than their S-Lipo counterpart (Figure 4J), suggesting that S-Exos would provide more rapid protection against viral infection. The features of Lung-Exos have significant advantages over their synthetic nanoparticle counterpart, suggesting that exosomes contain additional therapeutic benefits regarding pulmonary targeting, retention, and immune responses. Biological nanoparticles such as exosomes can be exploited as inhaled drug-delivery vehicles to maximize drug targeting, delivery, and therapeutic efficacy.

## DISCUSSION

Altogether, we have demonstrated the formulation and biodistribution of inhaled exosomes and Lipos in mouse and nonhuman primate (NHP) lungs. Exosomal and liposomal mRNA and protein cargo are stable and maintain biological function upon dry powder formulation and DPI, with a shelf life extending 28 days. We showed that lung-derived exosomes most efficiently evade mucoadhesion while maintaining higher exosomal mRNA and protein cargo deposition, retention, and distribution in lung than their Lipo counterpart. Lung-derived exosomes had the highest nanoparticle delivery to the bronchioles and parenchyma, suggesting that nanoparticle phenotypes that are native to the lung microenvironment have enhanced cellular targeting and bioavailability within the lung.

For the first time, we showed that dry powder formulation of lung-derived exosomes is room-temperature stable and is compatible with a clinically used DPI device for at-home administration. The *chlorocebus sabaesus* NHP model most closely replicates the human airway and respiratory physiology, and exosome delivery was verified in the upper and lower respiratory tracts through DPI administration. Lung-derived exosomes were retained in the primate lung 1 week after a single DPI dose, delivering functional mRNA and protein exosomal cargo to the nose and to the deep lung. Lung-derived exosomes are functional as an inhaled therapeutic and drug-delivery vesicle for both upper and lower respiratory diseases.

Additionally, lung-derived exosomes have enhanced efficacy for pulmonary disease applications. S-protein-encoding mRNA remains effective in exosomes after 1 month of room temperature storage when lyophilized. As an inhaled vaccine, S-protein-loaded lung-derived exosomes elicited stronger immune responses than their synthetic counterpart, emphasizing the immunological advantages of biological nanoparticles for inhaled vaccines. Mice vaccinated with S-protein-loaded exosomes were able to more rapidly clear pseudoviral infection than Lipos, further emphasizing the enhanced efficacy of exosomes. Lung-derived exosomes offer a unique room-temperature-stable nanoparticle drug-delivery system, with enhanced bioavailability, that can serve as an mRNA and protein drug-delivery vesicle tailored for lung diseases.

## EXPERIMENTAL PROCEDURES

### Resource availability

#### Lead contact

Further information and requests for resources and reagents should be directed to and will be fulfilled by the lead contact, Ke Cheng ([ke\\_cheng@ncsu.edu](mailto:ke_cheng@ncsu.edu)).

### Materials availability

All unique/stable reagents generated in this study are available from the lead contact with a completed materials transfer agreement.

### Data and code availability

All data reported in this paper will be shared by the [lead contact](#) upon request. This paper does not report original code. Any additional information required to reanalyze the data reported in this paper is available from the [lead contact](#) upon request.

### Cell culture

Human LSCs were generated from healthy whole-lung samples from the Cystic Fibrosis and Pulmonary Diseases Research and Treatment Center at the University of North Carolina at Chapel Hill and expanded as previously described.<sup>7,8,45</sup> LSCs were plated on a fibronectin-coated (Corning, Corning, NY, USA) flask and maintained in Iscove's modified Dulbecco's media (IMDM; Thermo Fisher Scientific, Waltham, MA, USA) containing 20% fetal bovine serum (FBS; Corning), 1% L-glutamine (Thermo Fisher Scientific), 0.5% Gentamicin (Thermo Fisher Scientific), and 0.18% 2-mercaptoethanol (Thermo Fisher Scientific). HEK 293T cells were purchased from American Type Culture Collection (ATCC; American Type Culture Collection, Manassas, VA, USA). HEK cells were plated on a flask and maintained in minimum essential media (MEM; Thermo Fisher Scientific) containing 10% FBS, 1% L-glutamine, 0.5% Gentamicin, and 0.18% 2-mercaptoethanol. Human bronchial epithelial cells were purchased from Lonza (CC-2540B; Lonza, Basel, Switzerland) and maintained according to manufacturer's instructions. Media changes on all cultures were performed every other day. LSCs and HEK cells were allowed to reach 70%–80% confluence before generating serum-free secretome (Lung-Secretome, HEK-Secretome), as previously described.<sup>13</sup> Lung- and HEK-Secretome were collected and filtered through a 0.22- $\mu$ m filter to remove cellular debris. All procedures performed in this study involving human samples were in accordance with the ethical standards of the institutional research committee and with the guidelines set by the Declaration of Helsinki.

### Exosome isolation and characterization

Lung-Exos and HEK-Exos were collected and isolated from Lung-Secretome and HEK-Secretome using an ultrafiltration method.<sup>46</sup> Filtered secretome was pipetted into a 100-kDa Amicon centrifugal filter unit (MilliporeSigma, Burlington, MA, USA) and centrifuged at 400 RCF and 10°C. After all media passed through the centrifugal filter unit, remaining exosomes were detached from the filter and resuspended using 1X Dulbecco's phosphate-buffered saline (DPBS; Thermo Fisher Scientific) with 25 mM trehalose (MilliporeSigma) for further analysis.<sup>47</sup> Pegylated Remote Loadable Lipos were purchased from Avanti Polar Lipids (Avanti Polar Lipids, Alabaster, AL, USA). LSC-Exos, HEK-Exos, and Lipos were fixed with 4% paraformaldehyde (PFA; Electron Microscopy Sciences, Hatfield, PA, USA) and 1% glutaraldehyde (Sigma-Aldrich, St. Louis, MO, USA) onto 100 mesh copper grids (Electron Microscopy Sciences, Hatfield, PA, USA) for TEM imaging (JEOL JEM-2000FX, Peabody, MA, USA). Samples were stained with Vanadium Negative Stain (ab172780; Abcam, Cambridge, UK). Sample concentrations and mean diameters were quantified by NTA before and after fluorescent label loading (NanoSight NS3000, Malvern Panalytical, Malvern, UK).

### Nanoparticle fluorescent label loading

RFP (ab268535; Abcam) was loaded into Lung-Exo and Lipo particles via electroporation, yielding RFP-Exos and RFP-Lipos.<sup>48</sup> One billion nanoparticles from each

sample were diluted in Gene Pulser Electroporation Buffer (Bio-Rad, Hercules, CA, USA) at a 1:9 ratio of nanoparticles to buffer. Ten  $\mu\text{g}$  of RFP were added to the nanoparticle-buffer solution and transferred to an ice-cold 0.4 cm Gene Pulser/MicroPulser Electroporation Cuvette (Bio-Rad). The electroporation cuvette was inserted into the Gene Pulser Xcell Total System (Bio-Rad) and electroporated under the following conditions: pulse type: square waveforms; voltage: 200 V; pulse length: 10 ms; number of pulses: 5; pulse interval: 1 s. Electroporation buffer was filtered out of the fluorescently labeled nanoparticles by the ultrafiltration method described above. GFP-encoding mRNA DasherGFP (Aldevron, Fargo, ND, USA) and RFP were loaded into Lung-Exos and Lipos, as described above. Lung-Exos, HEK-Exos, and Lipos were incubated with DiD labeling solution (V22889; Thermo Fisher Scientific) according to the manufacturer's instructions.

### Dry powder inhaler formulation and room-temperature stability

Lung-Exos, HEK-Exos, and Lipos were formulated into dry powder through lyophilization (Labconco, Kansas City, MO, USA). The nanoparticles were diluted in DPBS with 25 mM trehalose at a 1:8 ratio of nanoparticle solution volume to DPBS with 25 mM trehalose solution volume. Trehalose solution serves as a cryoprotectant and bulking agent for sufficient powder production.<sup>49</sup> Diluted nanoparticle solutions were stored at  $-80^{\circ}\text{C}$  overnight and lyophilized for 24 h. To test the room-temperature stability of lyophilized nanoparticles, RFP-loaded nanoparticles were stored in ambient room conditions for 0, 7, 14, 21, and 28 days. RFP cargo leakage was measured by ELISA (AKR-122; Cell Biolabs, San Diego, CA, USA) per the manufacturer's instructions in non-denatured RFP-loaded nanoparticle solutions. An RFP standard curve was generated using a second-order polynomial model, and cargo-leakage concentrations were interpolated from the standard curve. Percentage of leakage is reported as  $\left(\frac{\text{Interpolated Concentration}}{\text{Total Loaded RFP Concentration}}\right) \times 100$ .

### Nanoparticle S protein loading

Full-length DNA sequence of the S surface glycoprotein (SARS-CoV 2 isolate Wuhan-Hu-1, gene ID: 43740568, NC\_045512.2:21563-25384) was used to design a plasmid for *in vitro* transcription (IVT). Briefly, a stretch of DNA sequence with two restriction sites, I-CeuI and I-SceI, was cloned into pCR4Blunt-TOPO vector (45-0031; Thermo Fisher Scientific). The full length of the S protein sequence was split to design two gBlocks (Integrated DNA Technologies, Coralville, Iowa, USA), each of which was approximately 1.5- and 2.3-kb long. The T7 promoter sequence (5'-taatacagactcactataggg-3') was added to the 5' end of the 1.5-kb gBlock. The two gBlocks were ligated into the backbone vector using I-CeuI, BstEII, and I-SceI. Following sequence confirmation by Sanger sequencing (Genewiz, South Plainfield, NJ, USA), the full-length S protein DNA sequence, cut by I-CeuI and I-SceI, was extracted from TAE agarose gel. IVT was done following the manufacturer's manual of the MEGAscript Kit (AM1333; Thermo Fisher Scientific). High-yield RNA was treated with DNase and cleaned up using the Monarch RNA cleanup kit (T2040L; New England Biolab, Ipswich, MA, USA). Size-confirmed pure IVT S protein mRNA was aliquoted and stored in  $-80^{\circ}\text{C}$  until nanoparticle loading. S protein mRNA was loaded into LSC-Exo and Lipo nanoparticles via the electroporation method described above.

### Animal procedures

Seven-week-old male CD1 mice (022) were obtained from Charles River Laboratory (Wilmington, MA, USA). RFP-Exos, RFP-Lipos, Lung-Exos, S-Exos, and S-Lipos were administered via jet nebulization (Pari Trek S Portable 459 Compressor Nebulizer

Aerosol System, PARI, Starnberg, Germany) or dry powder inhalation as described above. Fluorescently labeled nanoparticles were given in a single dose of  $10^9$  particles per kg of body weight. Immediately after sacrifice, the lungs, heart, liver, kidneys, spleen, cecum, and brain were excised and imaged using a Xenogen Live Imager (PerkinElmer, Waltham, MA, USA). Blood was collected in Vacuette ethylenediaminetetraacetic acid (EDTA) tubes (Greiner Bio-One, Kremsmünster, Austria) and centrifuged at maximum speed for 5 min to separate out serum. All animal studies complied with the requirements of the Institutional Animal Care and Use Committee (IACUC) of North Carolina State University.

### NHP studies

Three *chlorocebus sabaues* monkeys were housed at Bioqual. The primates received a single dose of  $10^9$  lyophilized fluorescently labeled Lung-Exo particles per kg of body weight via DPI. The primates were necropsied 24 h and 1 week after DPI of Lung-Exos. All animal studies complied with the requirements under local, state, and federal regulations and were approved by the Bioqual IACUC.

### IgG and SIgA antibody titers

To collect BALF, the trachea was exposed by thoracotomy, and a transverse incision was made at the top of the bronchial bifurcation. A needle was inserted into the trachea to wash the lungs with 200  $\mu$ L of DPBS. Washing was repeated three times for a total of 600- $\mu$ L wash fluid. To collect NPLF, the trachea was cut in the middle, and the nasopharynx was washed upwards from the incision with 200  $\mu$ L DBPS. Washing was repeated three times for a total of 600- $\mu$ L wash fluid. S-protein-specific IgG (20154; Cell Signaling Technology, Danvers, MA, USA) and SIgA (58873; Cell Signaling Technology) from BALF and NPLF were measured by ELISA per the manufacturer's instructions.

### Pseudoviral challenge

Wild-type SARS-CoV-2 pseudoviral particles with a GFP reporter were purchased from Virongy (HaCoV2GFP; Virongy, Manassas, VA, USA). Pseudoviral particles were given in a single dose of  $10^9$  particles per kg of body weight via DPI, as described above.

### Statistical analysis

Statistical analysis was performed using GraphPad Prism analysis software (GraphPad Software, San Diego, CA, USA). Comparisons among two groups were performed using an unpaired t test, followed by Welch's correction test. Comparisons among more than two groups were performed using a parametric one-way ANOVA test followed by Bonferroni's multiple comparisons test.  $p \leq 0.05$  was considered statistically significant. The legend is as follows: \* $p \leq 0.05$ ; \*\* $p \leq 0.01$ ; \*\*\* $p \leq 0.001$ ; \*\*\*\* $p \leq 0.0001$ ; ns = not significant.

## SUPPLEMENTAL INFORMATION

Supplemental information can be found online at <https://doi.org/10.1016/j.matt.2022.06.012>.

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## AUTHOR CONTRIBUTIONS

Conceptualization, P.-U.C.D. and K.C.; methodology, K.D.P., A.M., G.S., D.Z., X.M., X.C., Y.C.; formal analysis, K.D.P.; investigation, K.D.P., A.M., G.S., D.S., H.L., B.L.d.J.A., A. George, E.B., M.C., A. Ghodsi., K.H., and J.L.; resources, A.C.B., A. Greebaum, and K.C.; writing, K.D.P., B.L.d.J.A., and A. George; supervision, P.-U.C.D. and K.C.; funding acquisition, K.C.

## DECLARATION OF INTERESTS

North Carolina State University has filed a patent on the technologies related to this study. K.C. is an equity holder and consultant of Xsome Biotech, Inc. Xsome has entered an exclusive license agreement with North Carolina State University.

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**Matter, Volume 5**

## **Supplemental information**

### **Inhalable dry powder mRNA vaccines**

#### **based on extracellular vesicles**

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## Supplementary Materials

### SUPPLEMENTARY METHODS

#### SDS-PAGE and western blot

Samples were lysed, denatured, and reduced by Laemmli sample buffer (Bio-Rad, Hercules, CA, USA) and  $\beta$ -mercaptoethanol (Bio-Rad, Hercules, CA, USA) at 90°C for 5 minutes. Protein samples and molecular ladder (Precision Plus Protein Unstained Standards; Bio-Rad, Hercules, CA, USA) were loaded into a 10% acrylamide precast Tris-Glycine gel (Bio-Rad, Hercules, CA, USA) for sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) separation. Gels were run at a stacking voltage of 100V until samples ran out of the wells, followed by a constant voltage of 200V. Gels were visualized and imaged in a Bio-Rad Imager (Bio-Rad, Hercules, CA, USA). Gels were transferred onto polyvinylidene fluoride membranes (PVDF; Bio-Rad, Hercules, CA, USA) using the Bio-Rad wet electroblotting transfer system (Bio-Rad, Hercules, CA, USA). Following transfer, membranes were washed three times in 1X phosphate-buffered saline with 0.1% Tween detergent (PBS-T; MilliporeSigma, Burlington, MA, USA) for 5 minutes each and blocked using 5% milk in PBS-T for one hour at room temperature. Membranes were blotted against anti- $\beta$ -Actin (ab6276; Abcam, Cambridge, United Kingdom), anti-RFP (ab62341; Abcam, Cambridge, United Kingdom), and anti-Spike (40592-T62; Sino Biological, Beijing, China) primary antibodies in 5% milk in PBS-T and incubated at 4°C for one week. After incubation, membranes were incubated with the corresponding goat anti-rabbit (ab6721; Abcam, Cambridge, United Kingdom) and goat anti-mouse (ab6789; Abcam, Cambridge, United Kingdom) HRP-conjugated secondary antibodies for 1 hour at room temperature. Membranes were then visualized using Clarity Western ECL Substrate (Bio-Rad,

23 Hercules, CA, USA) and imaged in a Bio-Rad Imager (Bio-Rad, Hercules, CA, USA). Band  
24 intensities were analyzed using ImageJ analysis software (NIH; <https://imagej.nih.gov/ij/>).

## 25 **Tissue clearing and imaging**

26 Mouse lungs were cleared using the BoneClear protocol.<sup>S1,S2</sup> Anesthetized mice were perfused  
27 with DPBS and 50 µg/mL heparin (Sigma-Aldrich, St. Louis, MO, USA). Lung tissues were  
28 dissected and fixed in a DPBS, 0.5% PFA (Electron Microscopy Sciences, Hatfield, PA, USA),  
29 and 10% sucrose (Sigma-Aldrich, St. Louis, MO, USA) solution at room temperature, then  
30 further fixed overnight in a DPBS and 0.5% PFA solution at 4°C. Fixed lung samples were  
31 incubated in a methanol (VWR, Randor, PA, USA) gradient of 20%, 40%, 60%, 80%, and 100%  
32 for two hours per percentage at room temperature. Then, the samples were decolorized overnight  
33 in a 30% hydrogen peroxide (Sigma-Aldrich, St. Louis, MO, USA) and 100% methanol solution  
34 at a 1:10 ratio respectively at 4°C. Lung tissues were then incubated in a reverse methanol  
35 gradient at room temperature and permeabilized with a DPBS, 0.02% Triton X-100 (Sigma-  
36 Aldrich, St. Louis, MO, USA), 0.01% sodium deoxycholate (Sigma-Aldrich, St. Louis, MO,  
37 USA), 10% dimethyl sulfoxide (DMSO; Sigma-Aldrich, St. Louis, MO, USA), 25 mM EDTA  
38 (Sigma-Aldrich, St. Louis, MO, USA) solution at 37°C overnight. Lung samples were then  
39 blocked with a DPBS, 0.02% Triton X-100, 10% DMSO, 5% normal donkey serum (Sigma-  
40 Aldrich, St. Louis, MO, USA), and 25 mM EDTA solution overnight at 37°C. Lung tissues were  
41 immunolabeled with anti-RFP (ab62341; Abcam, Cambridge, United Kingdom) primary  
42 antibody diluted in a DPBS, 0.02% Tween-20, 1 g/mL heparin, 5% normal donkey serum, and  
43 25 mM EDTA solution at a 1:200 ratio respectively for one week at 37°C. Tissues were washed  
44 and further immunolabeled with Cy3 (711-165-152; Jackson ImmunoResearch Laboratories,  
45 West Grove, PA, USA) secondary antibody diluted in a DBPS, 0.02% Tween-20, 1 g/mL

46 heparin, 5% normal donkey serum, and 25 mM EDTA solution at a 1:500 ratio respectively for 5  
47 days at 37°C. Lung tissues were washed, incubated in a methanol gradient, then incubated twice  
48 in a dichloromethane (Sigma-Aldrich, St. Louis, MO, USA) and methanol solution at a 1:2 ratio  
49 respectively at room temperature, and followed by four incubations in 100% dichloromethane.  
50 Finally, lung samples were cleared three times with 100% dibenzyl-ether (Sigma-Aldrich, St.  
51 Louis, MO, USA) at room temperature and imaged using a custom-built light sheet  
52 microscope.<sup>S1</sup> Whole lung images and movies were captured using Imaris image analysis  
53 software (Imaris, Oxford Instruments, Abingdon, United Kingdom).

#### 54 **Particle segmentation**

55 Cleared mouse lungs were analyzed using ImageJ analysis software. In each image, pixels that  
56 belonged to exosomes or liposomes were segmented via thresholding, during which the intensity  
57 threshold was decided manually. The selection brush tool was used to refine the masks generated  
58 by thresholding and to segment the airway regions. Quantification of areas or pixels was then  
59 performed based on the extracted masks.

#### 60 **Flow cytometry**

61 Exosome and liposome uptake were evaluated in lung parenchymal cells and single-cell  
62 suspensions of murine lung cells by flow cytometry. Lung parenchymal cells were co-cultured  
63 with RFP-Exo and RFP-Lipo for 24 hours to evaluate nanoparticle protein cargo uptake. Lung  
64 parenchymal cells were co-cultured with RFP-Exo and RFP-Lipo for 1 minute to evaluate  
65 nanoparticle mRNA cargo uptake. Single-cell suspensions of murine lung cells were generated  
66 from mice that received nebulized RFP-Exo or RFP-Lipo. Lung parenchymal cells and murine  
67 lung cells were permeabilized (BD, Franklin Lakes, NJ, USA), GFP (ab290; Abcam, Cambridge,  
68 United Kingdom) and RFP primary antibodies were incubated for 1 hour at 4°C, and their

69 corresponding goat anti-rabbit AF488-conjugated (ab150077; Abcam, Cambridge, United  
70 Kingdom) and AF647-conjugated (ab150079; Abcam, Cambridge, United Kingdom) secondary  
71 antibodies were incubated for 1 hour at 4°C. The internalization of nanoparticle mRNA and  
72 protein cargo by cells was examined by flow cytometry (CytoFlex, Beckman Coulter, Brea, CA,  
73 USA).

#### 74 **Air-liquid interface system**

75 A model of the human airway at the air-liquid interface was created by seeding human bronchial  
76 epithelial cells onto a 0.4 µm pore polycarbonate membrane and lung parenchymal cells onto a  
77 6.5 mm well in a transwell system (Corning Incorporated, Corning, NY, USA). Cells were  
78 maintained for one week before administering DiD-labeled exosomes and liposomes to the  
79 human bronchial epithelial cells. Nuclei in the transwell wells were visualized by adding  
80 NucBlue™ Live ReadyProbes™ Reagent (R37605; ThermoFisher Scientific, Waltham, MA,  
81 USA) to the media and analyzed using ImageJ analysis software.

#### 82 **DPI fabrication**

83 A DPI for nanoparticle inhalation to mice was fabricated as previously described, with  
84 modifications.<sup>S3</sup> The inhalation apparatus was adapted by using a plastic microcentrifuge tube as  
85 the powder receptacle. A plastic 250 mL centrifuge tube was attached to the powder receptacle to  
86 serve as a containment chamber for the un-anesthetized mouse; this optimized mouse muzzle  
87 orientation. A DPI for nanoparticle inhalation to primates was assembled using the RS01 high-  
88 resistance DPI (239700002AA; Berry Global, Evansville, IN, USA) connected to an aerosol  
89 chamber inhaler spacer (Canack Technology Ltd., Vancouver, Canada).

#### 90 **Histology**

91 Immunostaining was performed on transwell membranes and tissue slides fixed in 4% PFA  
92 (Electron Microscopy Sciences, Hatfield, PA, USA) in DPBS for 30 minutes, followed by  
93 permeabilization and blocking with Dako Protein blocking solution (Aglient Technologies, Santa  
94 Clara, CA, USA) with 0.1% saponin (Sigma-Aldrich, St. Louis, MO, USA) at room temperature  
95 for 1 hour. Membranes were immunolabeled with anti-MUC5b (ab77995; Abcam, Cambridge,  
96 United Kingdom) primary antibody diluted in Dako Protein blocking solution and its  
97 corresponding goat anti-mouse (A10667; Invitrogen, Waltham, MA, USA) AF488-conjugated  
98 secondary antibody diluted in Dako Protein blocking solution. Membranes and slides were  
99 mounted with ProLong Gold Antifade Mountant (Invitrogen, Waltham, MA, USA) and ProLong  
100 Gold Antifade Mountant with DAPI (Invitrogen, Waltham, MA, USA). Membrane and slides  
101 were imaged on the Olympus FLUOVIEW CLSM (Olympus; FV3000, Shinjuku, Tokyo, Japan)  
102 with an Olympus UPlanSAPO 10x objective (Olympus; 1-U2B824, Shinjuku, Tokyo, Japan) and  
103 Olympus UPlanSAPO 60x objective (Olympus; 1-U2B832, Shinjuku, Tokyo, Japan). H&E  
104 staining was performed on tissue slides (Hematoxylin HHS16; Eosin 318906; Sigma-Aldrich, St.  
105 Louis, MO, USA) and imaged on the Leica DMI8 (Leica Microsystems, Wetzlar, Germany).  
106 Tissue slides were analyzed using ImageJ analysis software.

### 107 **AFM imaging**

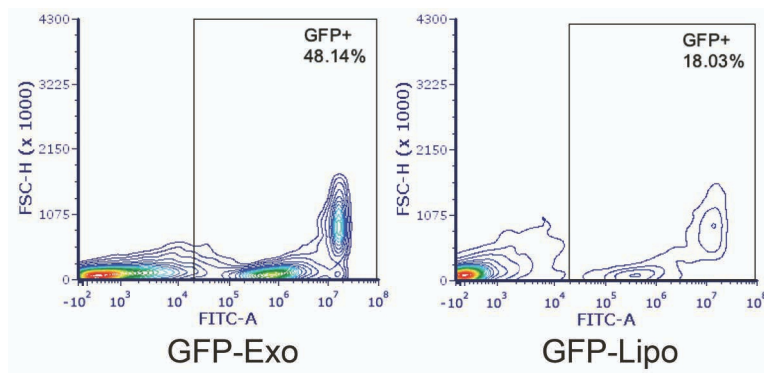
108 Lung-Exo, HEK-Exo, and Lipo were prepared as freshly isolated particles in PBS and trehalose  
109 solution (Fresh), lyophilized powder (Lyophilized), and lyophilized powder reconstituted in  
110 DNase/RNase-free distilled water (Invitrogen, Waltham, MA, USA) (Reconstituted). Samples  
111 were added onto coverslips functionalized by (3-Aminopropyl)triethoxysilane (APTMS; Sigma-  
112 Aldrich, St. Louis, MO, USA) as previously described.<sup>S4,S5</sup> Samples were imaged on the MFP-  
113 3D AFM (Asylum Research, Oxford Instruments, Abingdon, United Kingdom) with a silicon

114 scanning probe microscopy-sensor probe (ARROW-NCR-50; Neuchâtel, Switzerland). Height  
115 and diameter measurements were analyzed using Asylum Research Software Version 16  
116 (Asylum Research, Oxford Instruments, Abingdon, United Kingdom) ran on Igor Pro 6  
117 (WaveMetrics, Tigard, OR, USA).

### 118 **Particle counting**

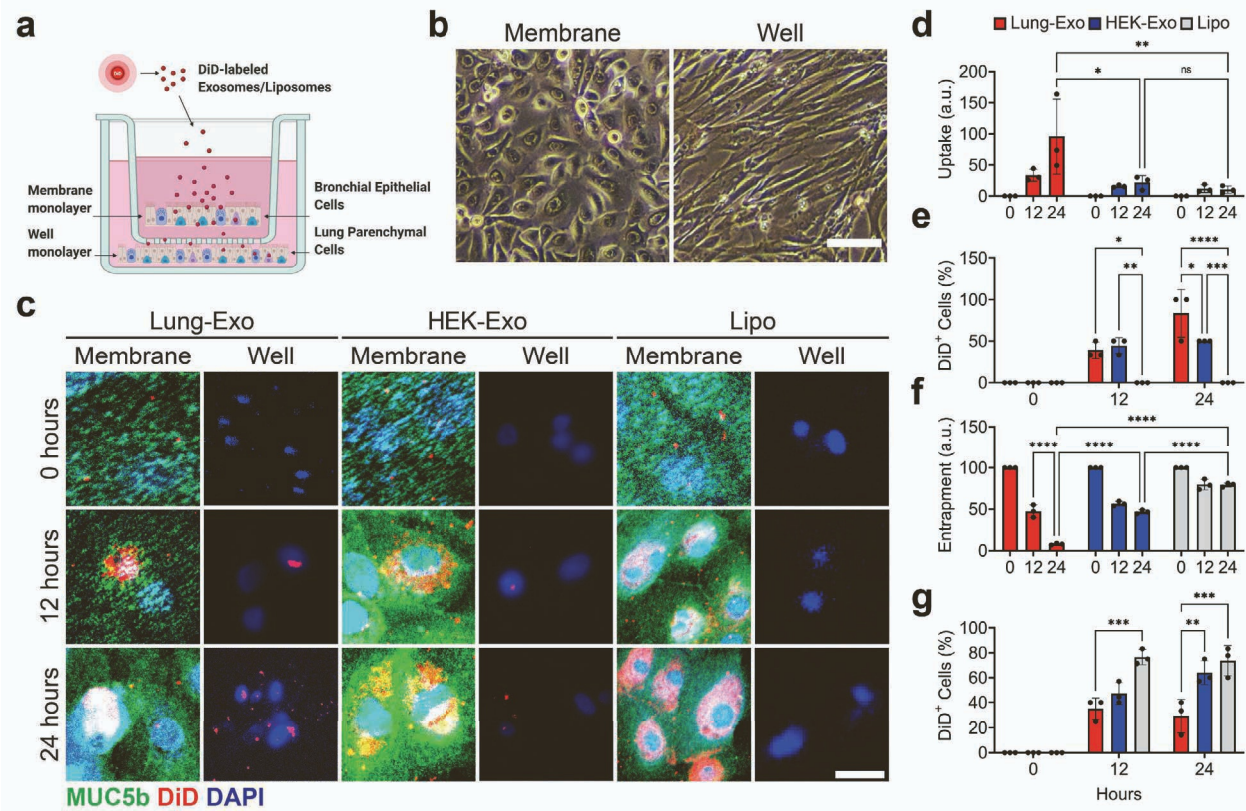
119 Lyophilized Lung-Exo, HEK-Exo, and Lipo were encapsulated in Quali V Inhalation capsules  
120 (Qualicaps, Inc., Whitsett, NC, USA) and inserted into low-, medium-, high-, and ultra-high-  
121 resistance DPIs (Berry Global, Evansville, IN, USA). The DPIs were inserted into a negative-  
122 pressure vacuum chamber to aerosolize the samples. Samples were aerosolized for 30 seconds,  
123 followed by particle size distribution measurement using the TSI AeroTrak particle counter  
124 (9306-V2, TSI, Shoreview, MN, USA) for 1-minute time intervals at ambient room temperature.  
125 Differential particle counts measure values ranging from one bin size (i.e. 3  $\mu\text{m}$ ) up to, but not  
126 including, its successive bin size (i.e. 5  $\mu\text{m}$ ). Cumulative particle counts measure values from  
127 one bin size (i.e. 3  $\mu\text{m}$ ) and greater. Bin sizes within the respirable fraction and nanoparticle  
128 biodistribution<sup>S6</sup> in the lung determined the optimal resistance DPI.

### 129 **SUPPLEMENTARY FIGURES**



130

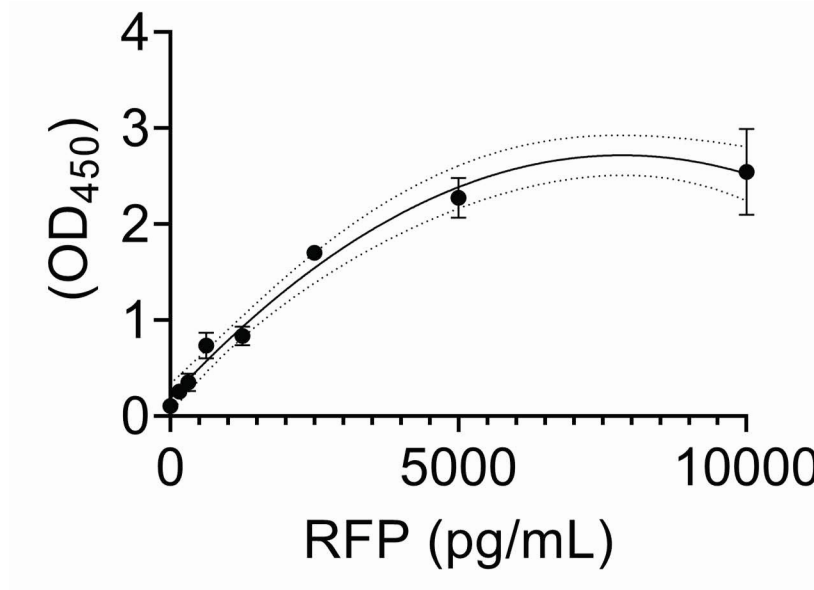
131 **Fig. S1. mRNA uptake by lung parenchymal cells.** Flow cytometry analysis of lung  
 132 parenchymal cells co-cultured with GFP-Exo or GFP-Lipo.



133  
 134 **Fig. S2. Mucus penetrance of nanoparticles.** (A) Schematic of DiD labeling and administration  
 135 to an air-liquid interface transwell system. Created with BioRender.com. (B) Brightfield images  
 136 of human bronchial epithelial cells on the transwell membrane and lung parenchymal cells on the  
 137 transwell well; scale bar=10 μm. (C) Representative immunostaining images of transwell  
 138 membranes and transwell wells for MUC5b (green), DiD (red), and DAPI (blue); scale bar=10  
 139 μm. (D) Quantification of Lung-Exo, HEK-Exo, and Lipo pixel intensity normalized to nuclei in  
 140 lung parenchymal cells; n=3 per group. (E) Percentage of DiD+ lung parenchymal cells; n=3 per  
 141 group. (F) Quantification of Lung-Exo, HEK-Exo, and Lipo pixel intensity normalized to nuclei

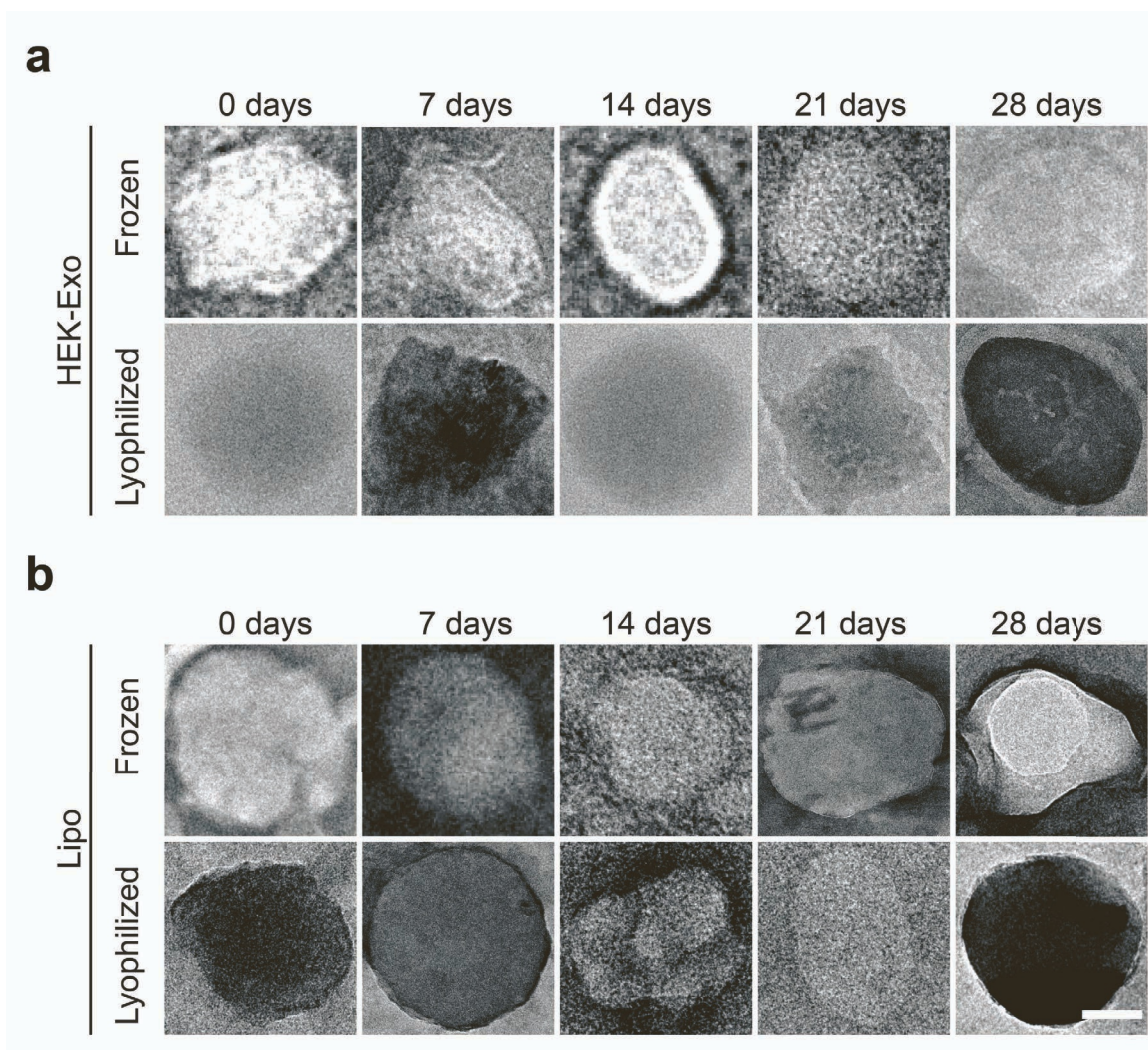


142 in human bronchial epithelial cells; n=3 per group. (G) Percentage of DiD<sup>+</sup> human bronchial  
143 epithelial cells; n=3 per group.



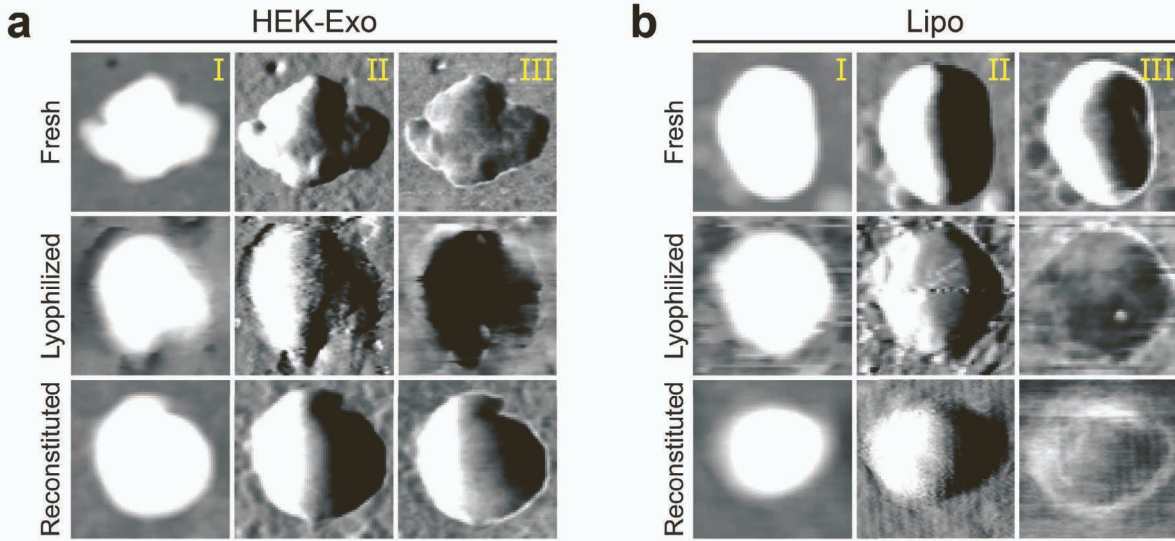
144

145 **Fig. S3. Standard curve for cargo leakage.** Standard curve of RFP concentrations in  
146 duplicates. Interpolation of the standard curve is represented by the solid line. The 95%  
147 confidence interval is represented by the dashed line.



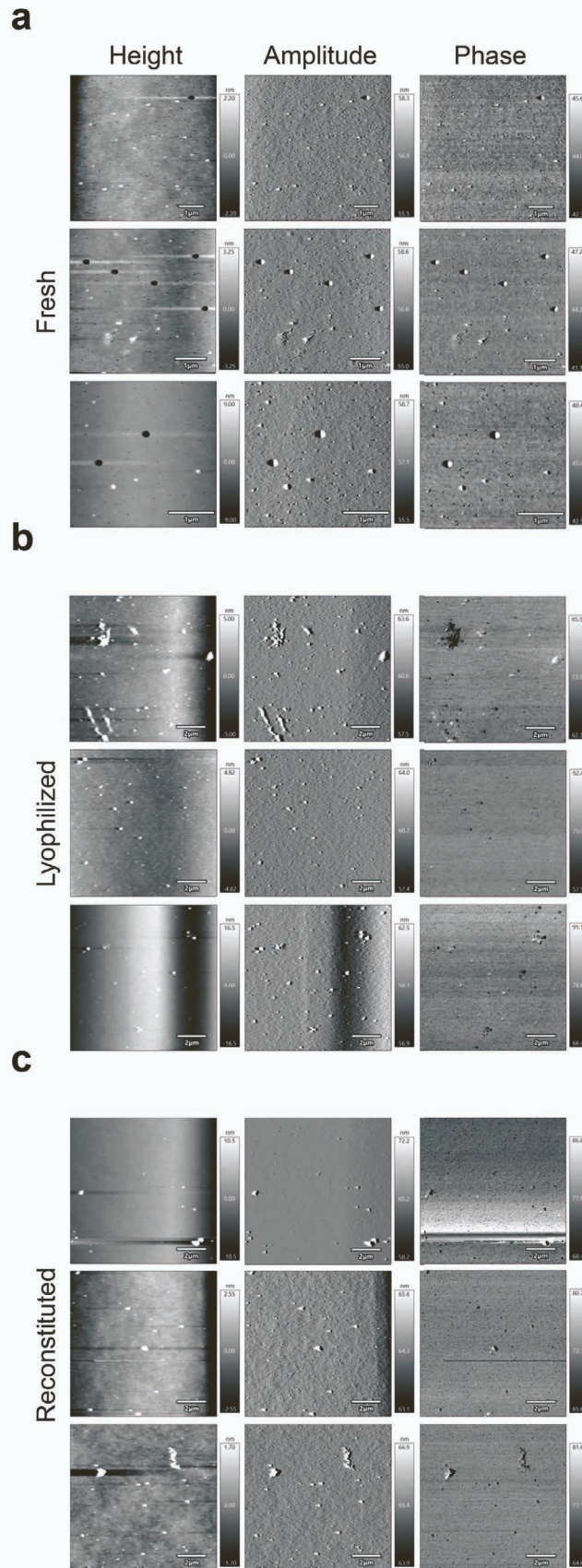
148

149 **Fig. S4. Morphology of nanoparticles at fresh and lyophilized formulations.** TEM images of  
 150 HEK-Exo (A) and Lipo (B) at frozen (Frozen) or room (Lyophilized) temperatures; scale bar=50  
 151 nm.

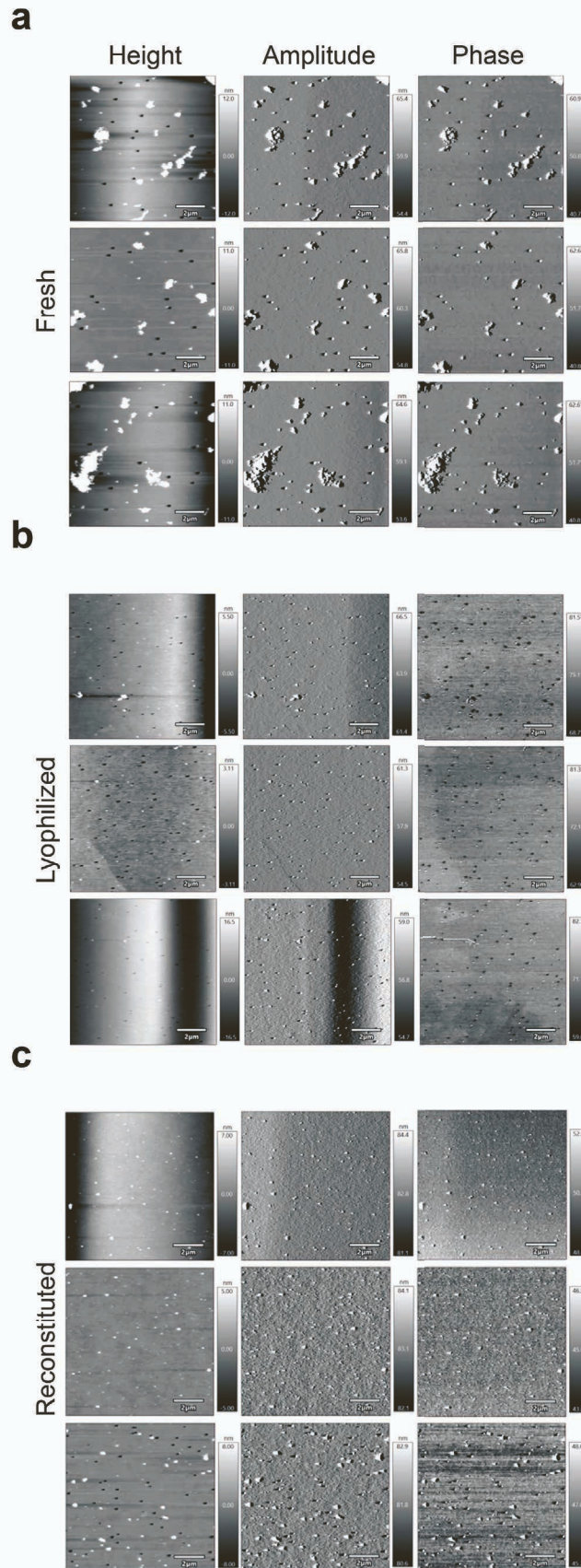


152

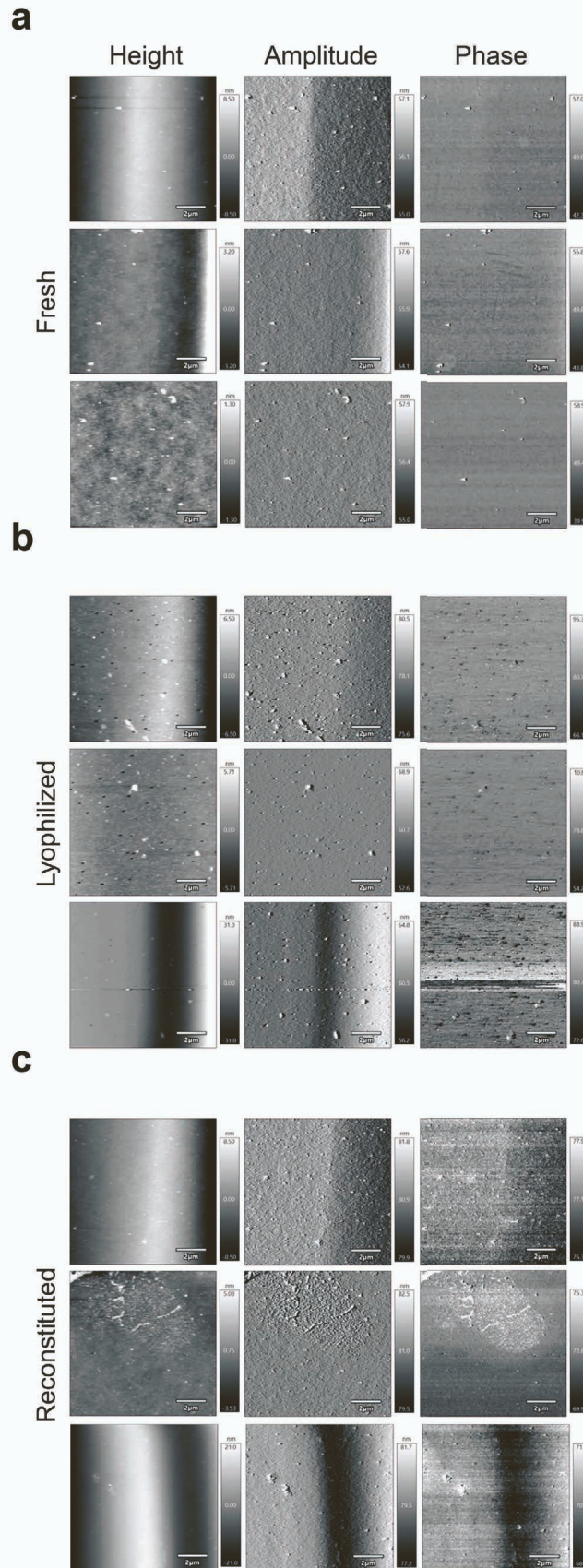
153 **Fig. S5. Membrane integrity of nanoparticles across formulations.** Representative AFM  
 154 height (i), amplitude (ii), and phase (iii) images of HEK-Exo (A) and Lipo (B) across fresh,  
 155 lyophilized, and reconstituted formulations.



157 **Fig. S6. Clumping properties of Lung-Exo.** AFM images of fresh (A), lyophilized (B) and  
158 reconstituted (C) Lung-Exo across fresh, lyophilized, and reconstituted formulations.

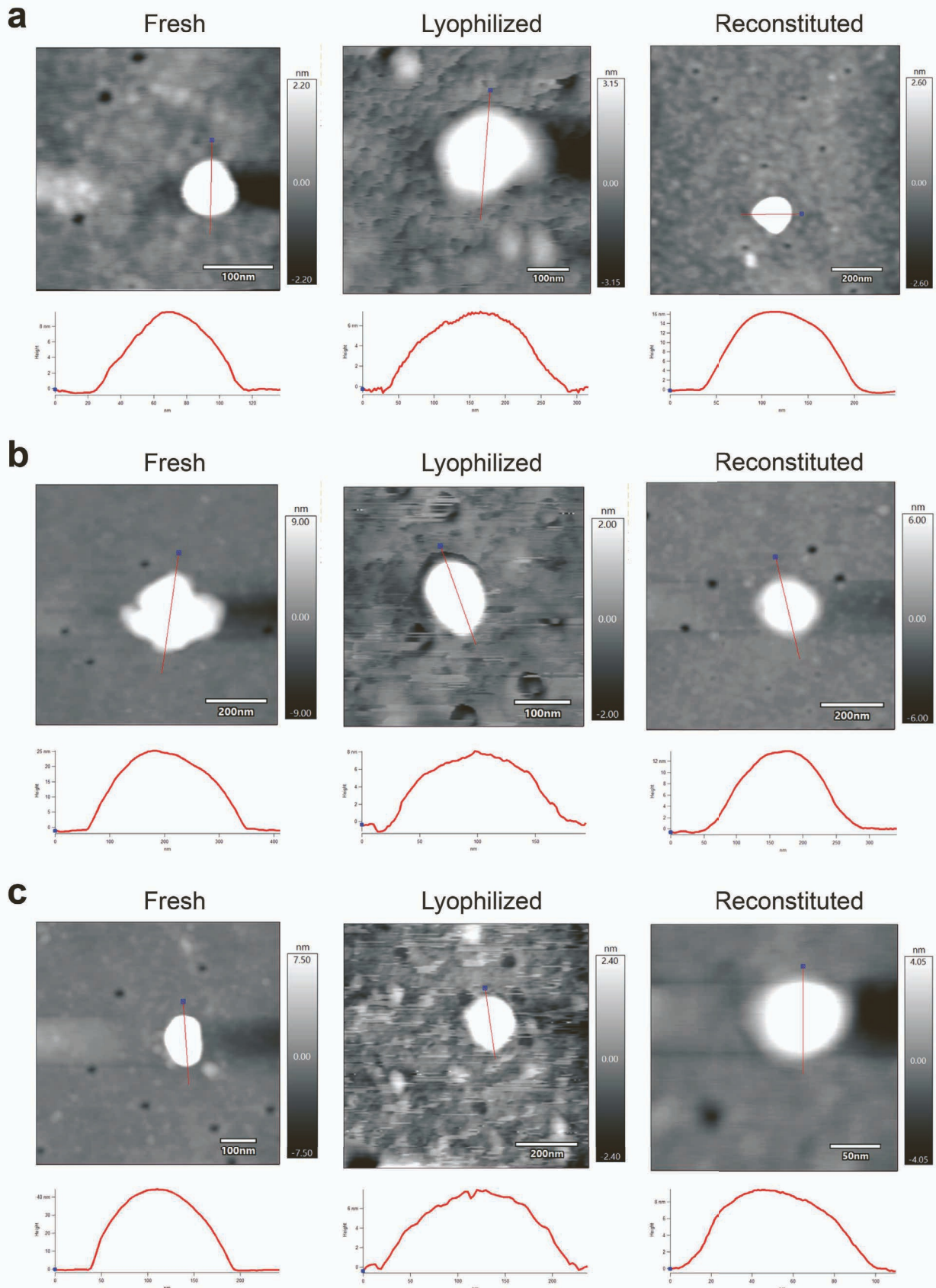


160 **Fig. S7. Clumping properties of HEK-Exo.** AFM images of fresh (A), lyophilized (B) and  
161 reconstituted (C) HEK-Exo across fresh, lyophilized, and reconstituted formulations.

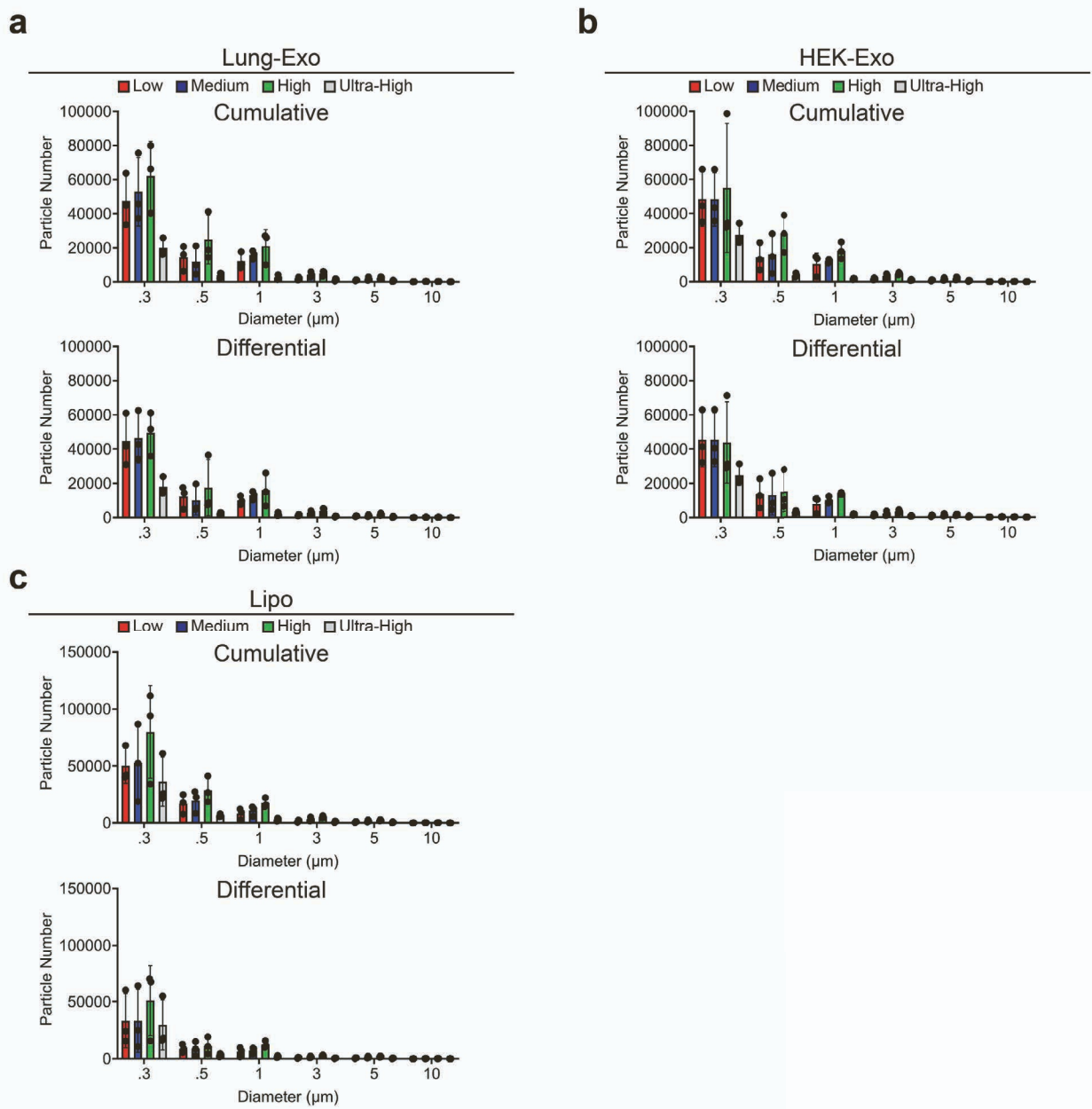




163 **Fig. S8. Clumping properties of Lipo.** AFM images of fresh (A), lyophilized (B) and  
164 reconstituted (C) Lipo across fresh, lyophilized, and reconstituted formulations.

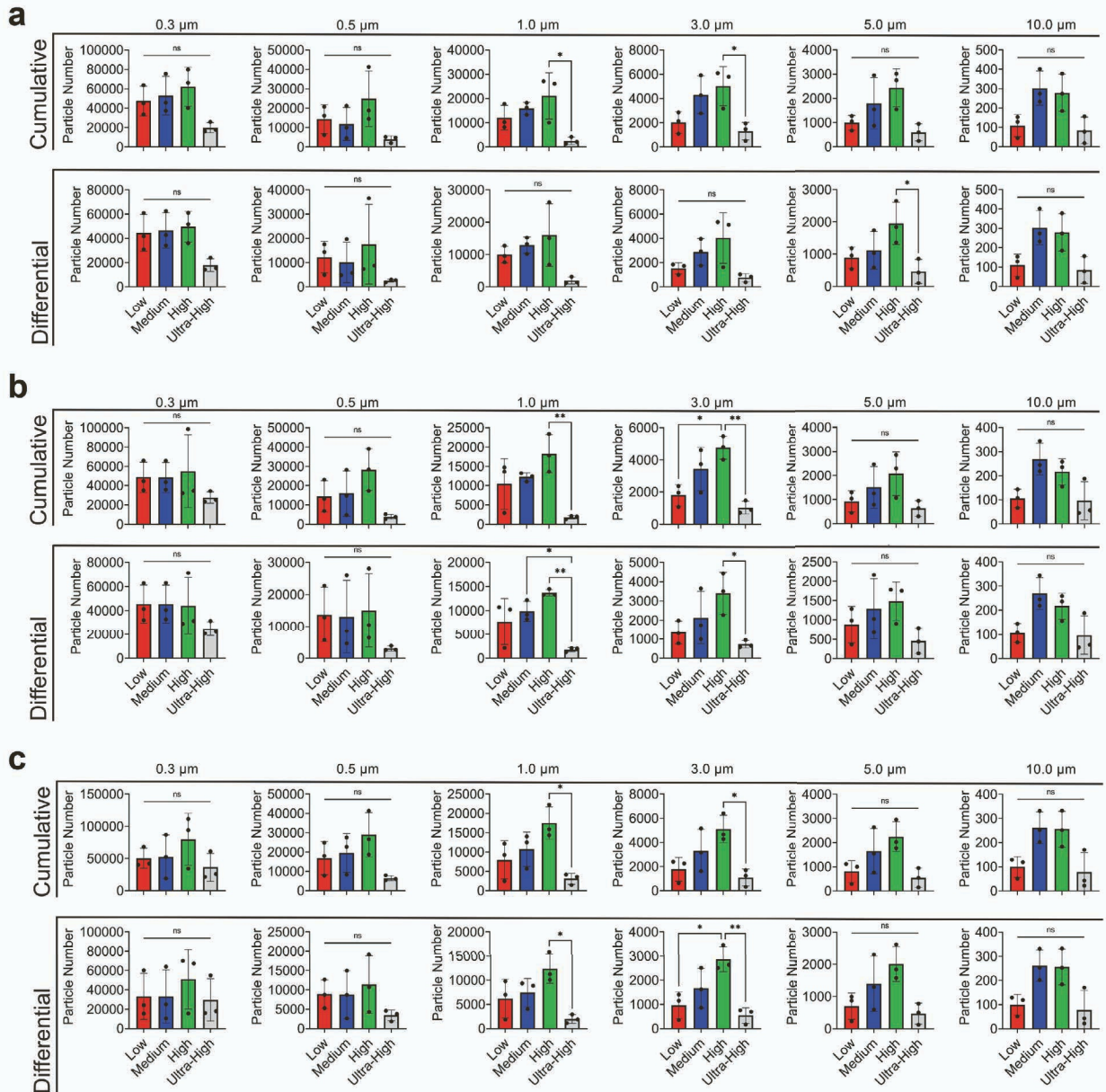


166 **Fig. S9. Cross-section measurements of nanoparticles.** Representative AFM height images of  
 167 Lung-Exo (A), HEK-Exo (B), and Lipo (C) across fresh, lyophilized, and reconstituted  
 168 formulations. Cross-section measurements were repeated on nine singular Lung-Exo (see Fig.  
 169 2C), HEK-Exo (see Fig. S4), and Lipo (see Fig. S4) height images to obtain height and diameter  
 170 measurements (see Fig. 2D).



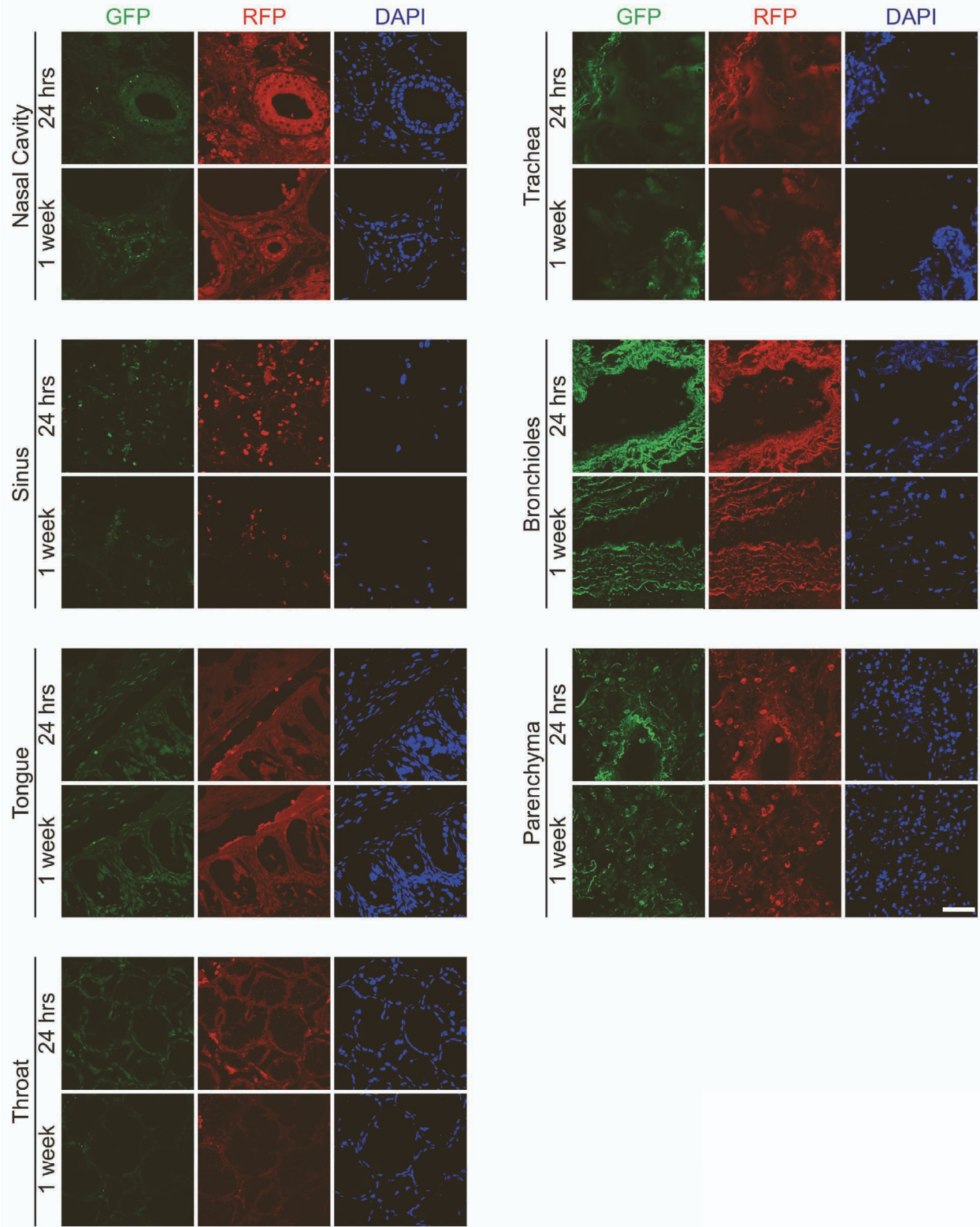
171

172 **Fig. S10. Total particle counts from various resistance DPIs.** Total cumulative and  
 173 differential particle counts of Lung-Exo (A), HEK-Exo (B), and Lipo (C) distributed by low-,  
 174 medium-, high-, and ultra-high-resistance DPIs.

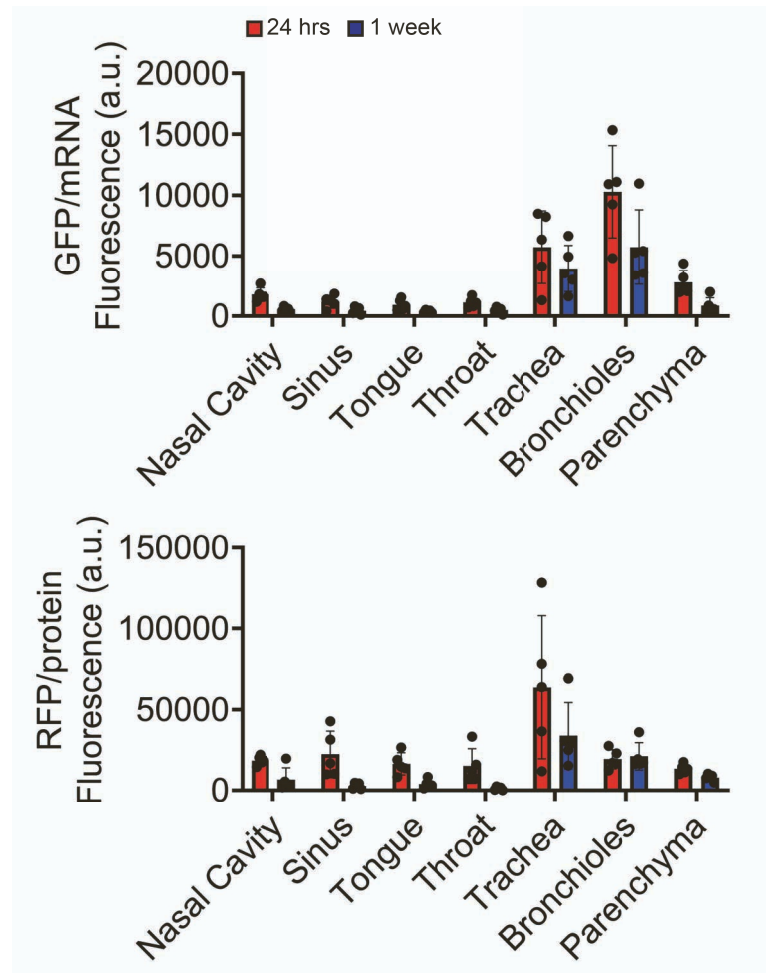


175

176 **Fig. S11. Particle counts at six bin sizes from various resistance DPIs.** Cumulative and  
177 differential particle count evaluation of LSC-Exo (A), HEK-Exo (B) and Lipo (C) distributed by  
178 low-, medium-, high-, and ultra-high-resistance DPI.



180 **Fig. S12. Immunostaining of simian upper and lower respiratory tracts.** Representative split-  
181 channel immunostaining images of the following sections from AGMs that received Lung-Exo  
182 via DPI: nasal cavity, sinus, tongue, throat, trachea, bronchioles, and parenchyma sections for  
183 GFP (green), RFP (red) and DAPI (blue); scale bar=100  $\mu\text{m}$ .

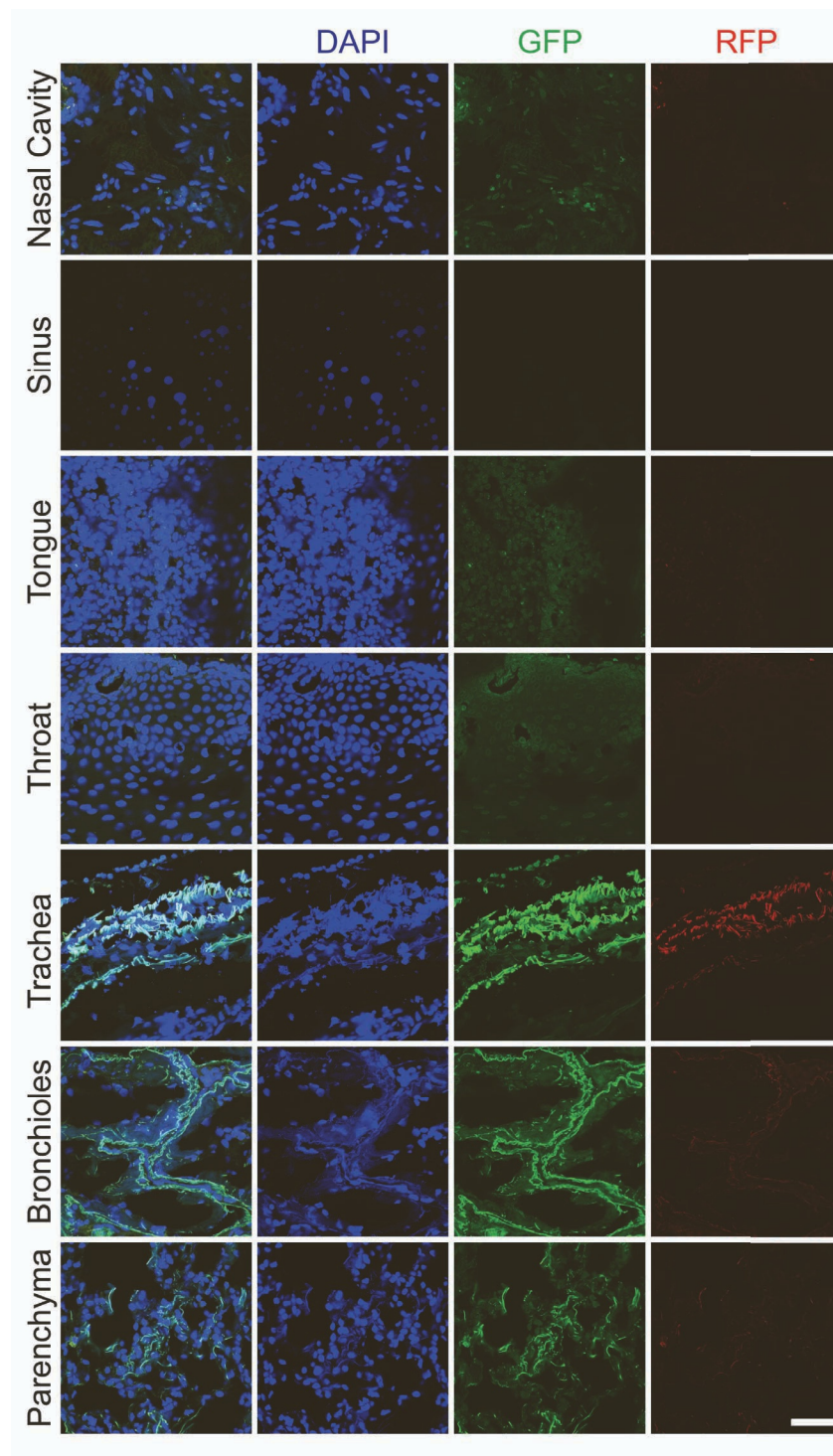


184

185 **Fig. S13. Total exosome deposition in the simian upper and lower respiratory tracts.**

186 Quantification of mRNA and protein fluorescence in simian upper and lower respiratory tissues

187 24 hours and 1 week after lyophilized Lung-Exo via dry-powder inhalation.



188

189 **Fig. S14. Autofluorescence of simian tissues.** Representative autofluorescence of the following  
 190 sections from AGMs that received no treatment: nasal cavity, sinus, tongue, throat, trachea,



191 bronchioles, and parenchyma sections for GFP (green) channel, RFP (red) channel, and DAPI  
192 (blue) staining; scale bar=100  $\mu$ m.

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