

Testing thousands of nanoparticles *in vivo*

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I co-founded GuideTx, which was acquired by Beam.

I am an advisor to GV.

What have we learned?

Many lipids are unexplored.

Keep LNP structure as simple as possible.

You find what you look for, so run predictive assays.

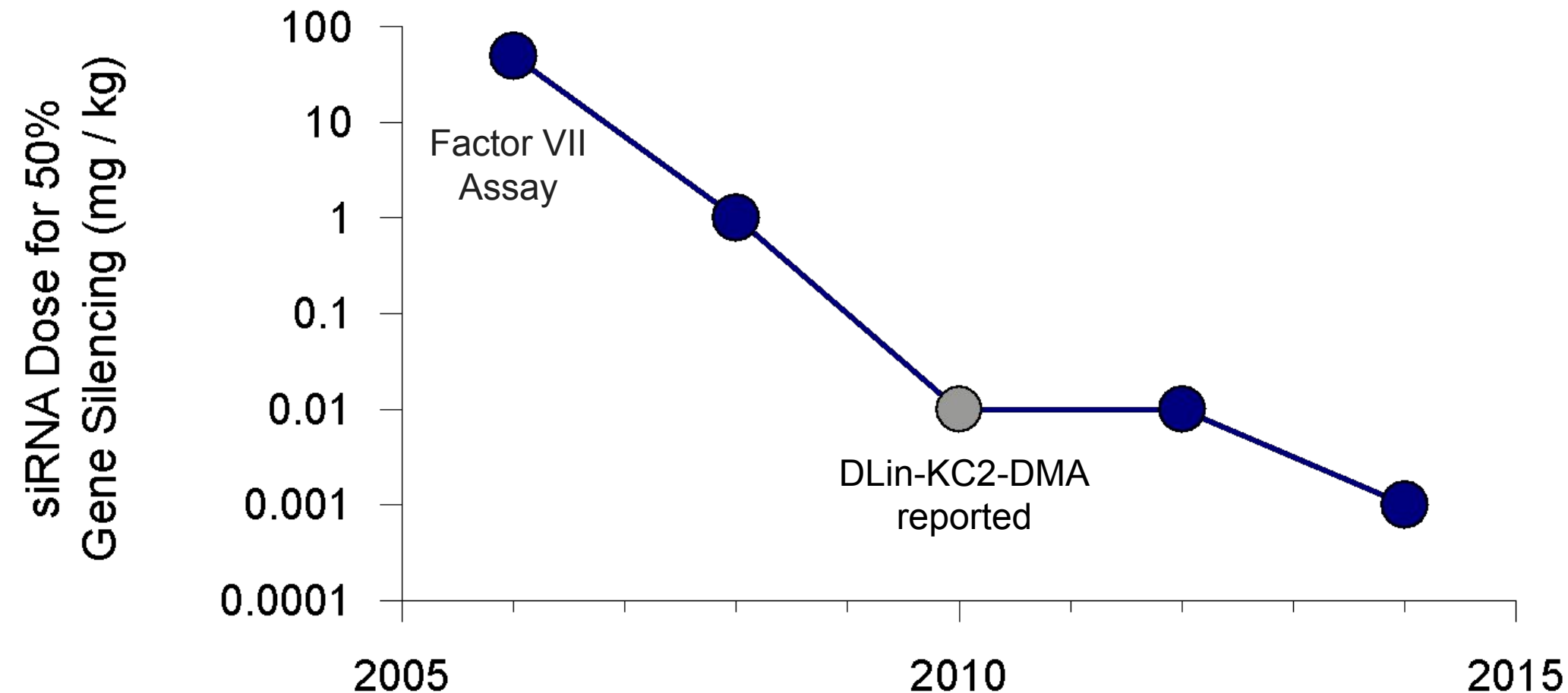
Non-hepatocyte delivery will be important



The Onpattro story and the clinical translation of nanomedicines containing nucleic acid-based drugs

The regulatory approval of Onpattro, a lipid nanoparticle-based short interfering RNA drug for the treatment of polyneuropathies induced by hereditary transthyretin amyloidosis, paves the way for clinical development of many nucleic acid-based therapies enabled by nanoparticle delivery.

The field looked for hepatocyte delivery for a decade



History lesson from hepatocyte RNA drugs:

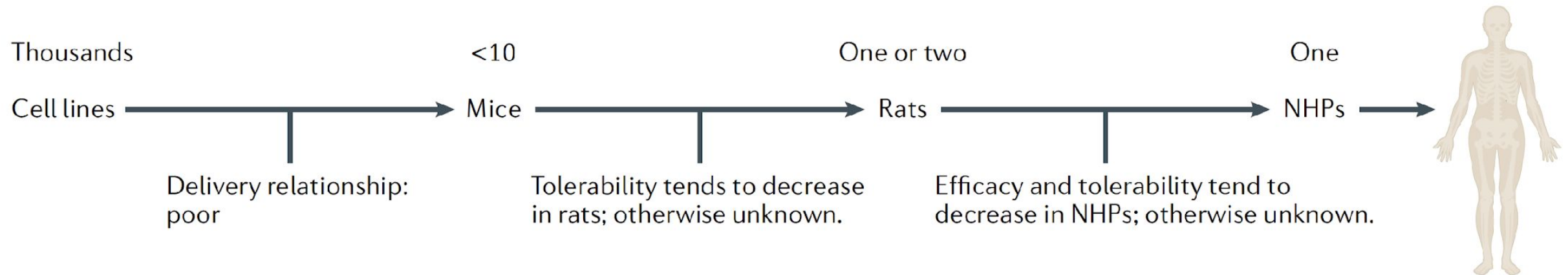
By iteratively quantifying how hundreds of nanoparticles delivered RNA to hepatocytes *in vivo*, the field found clinically relevant delivery to hepatocytes.

Our hypothesis:

By iteratively quantifying how thousands of nanoparticles deliver RNA to any combination of desired cell types *in vivo*, you may find clinically relevant delivery to other cell types.

Is this an assay problem?

We study / optimize the nanoparticle discovery pipeline

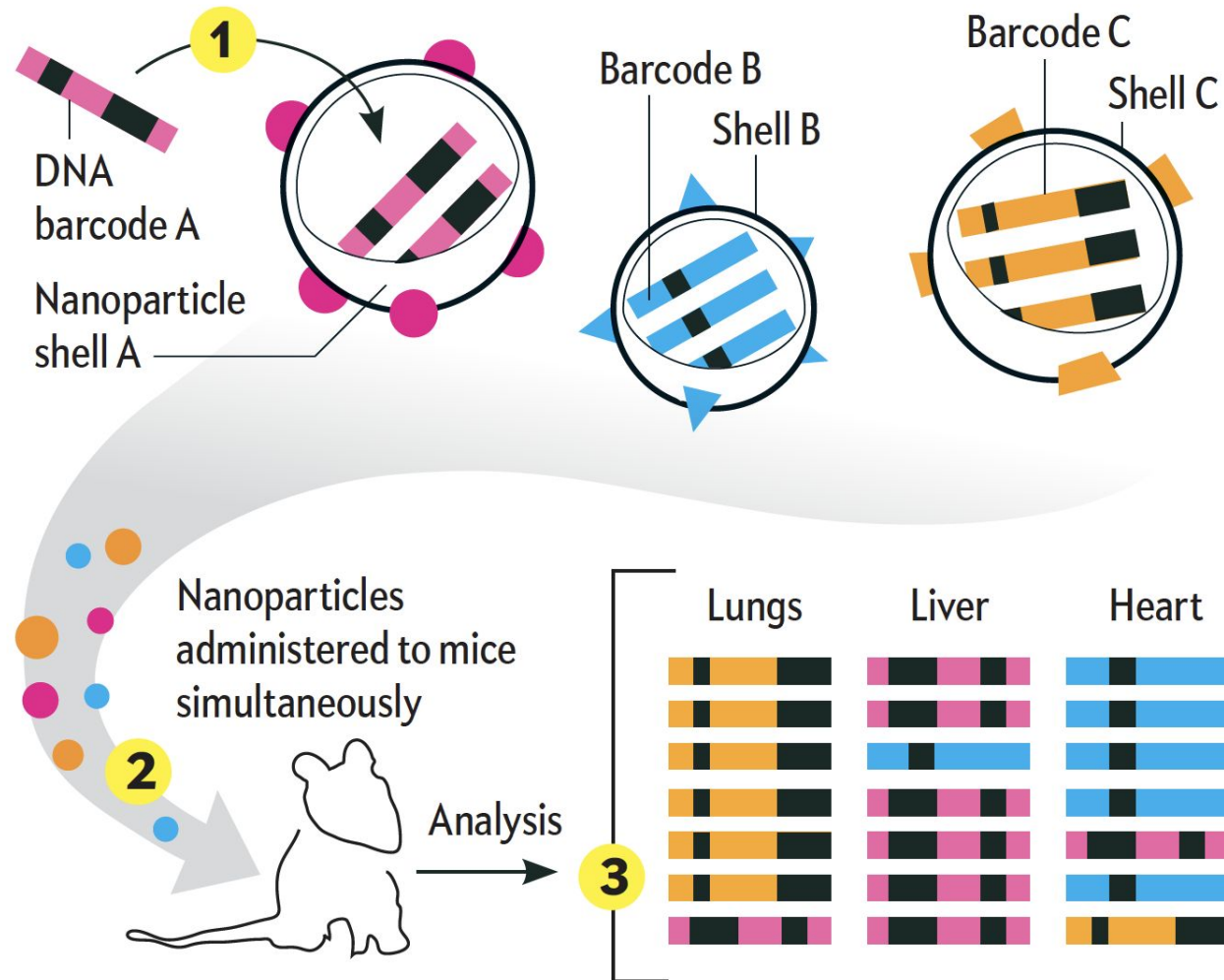


We optimize the pipeline with DNA barcoded LNPs

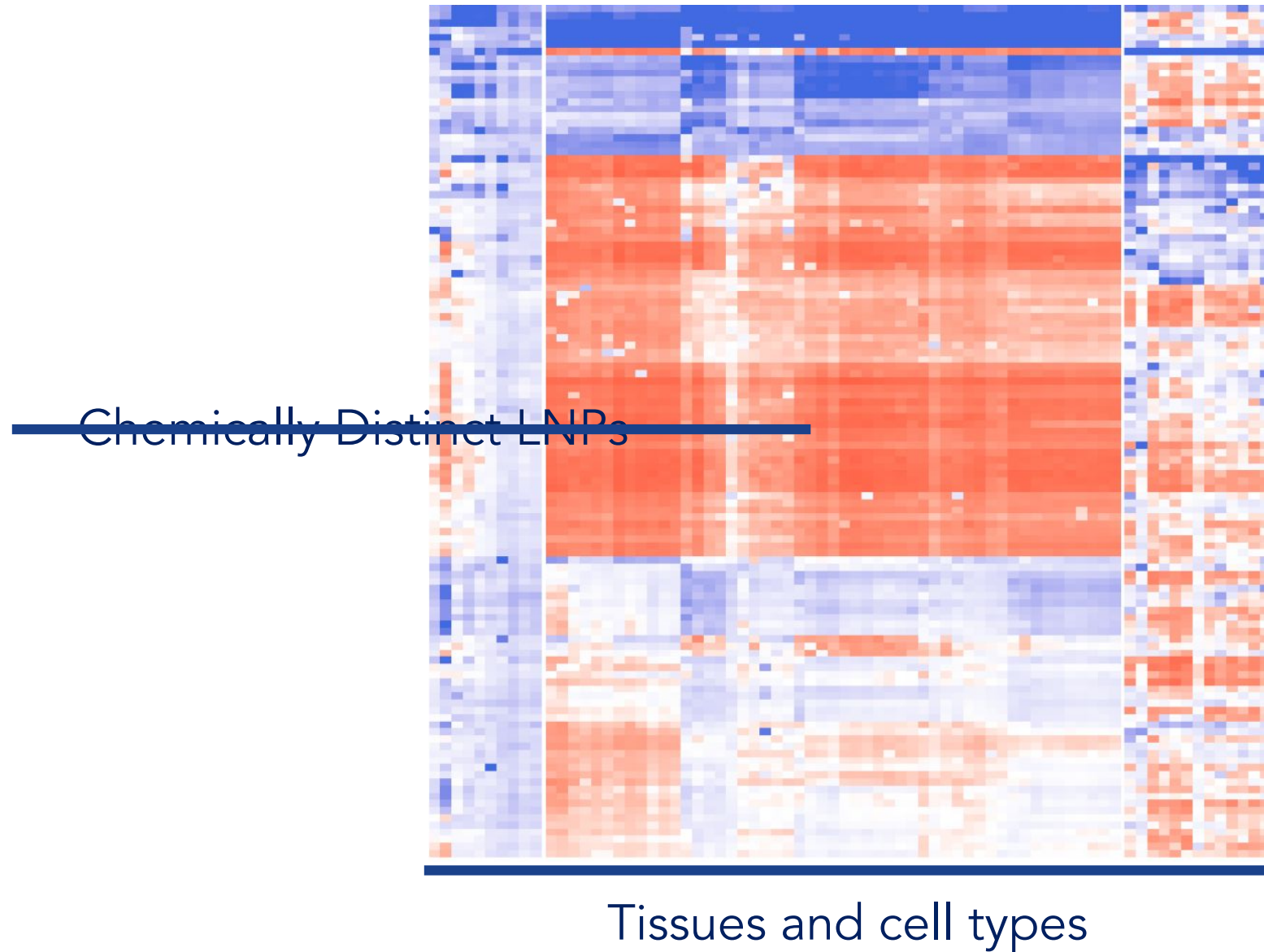
These assays are:

- (1) High throughput (up to ~300 LNPs / mouse)
- (2) *In vivo* (mice)
- (3) Functional (mRNA ☐ protein, siRNA ☐ silencing protein)
- (4) Iterative (data from LNPs 1-200 inform LNPs 201-400)

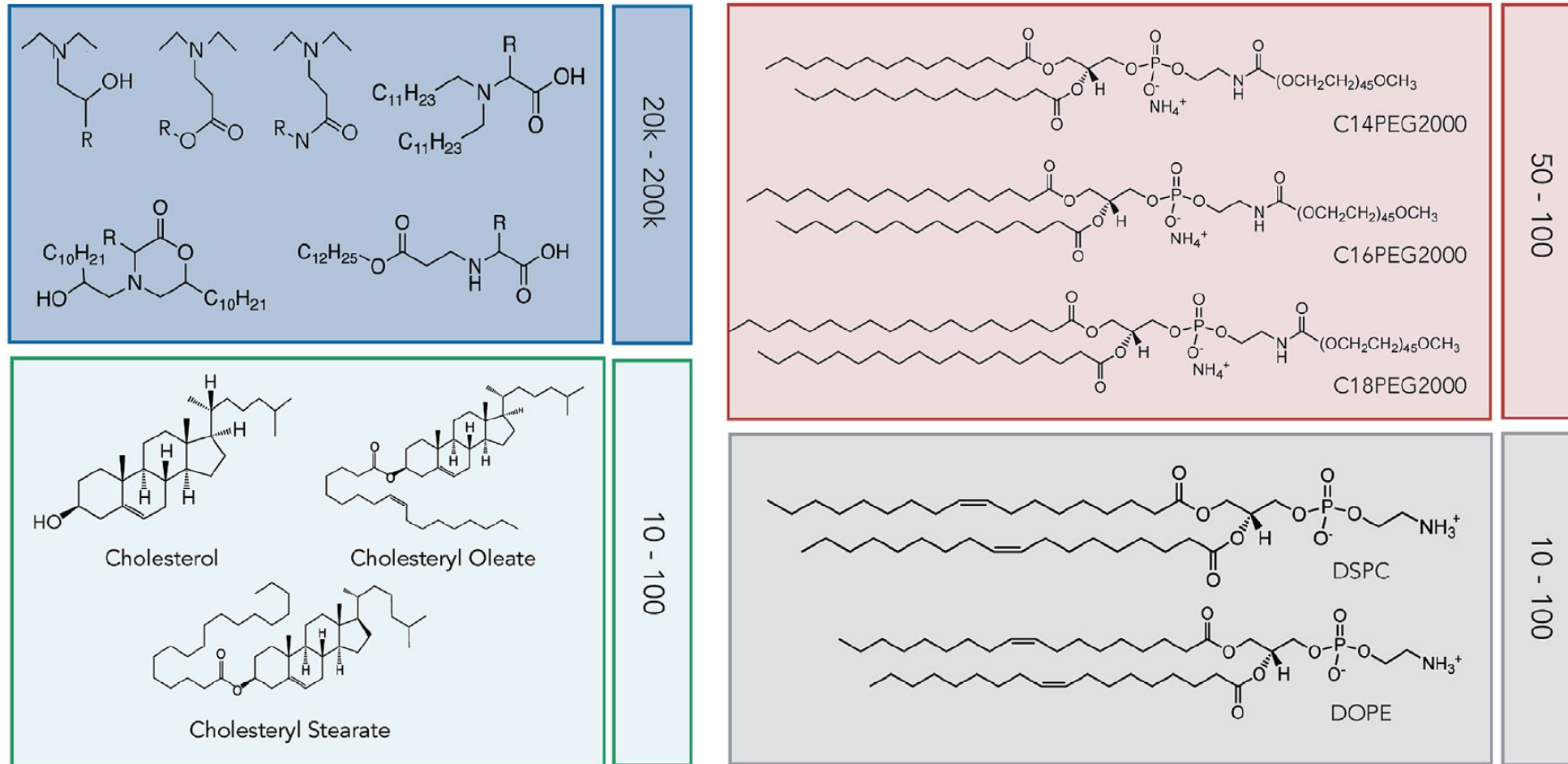
We label individual LNPs with barcodes, then pool / administer



We study many LNPs / animal



Testing many LNPs matters: chemical space is very large

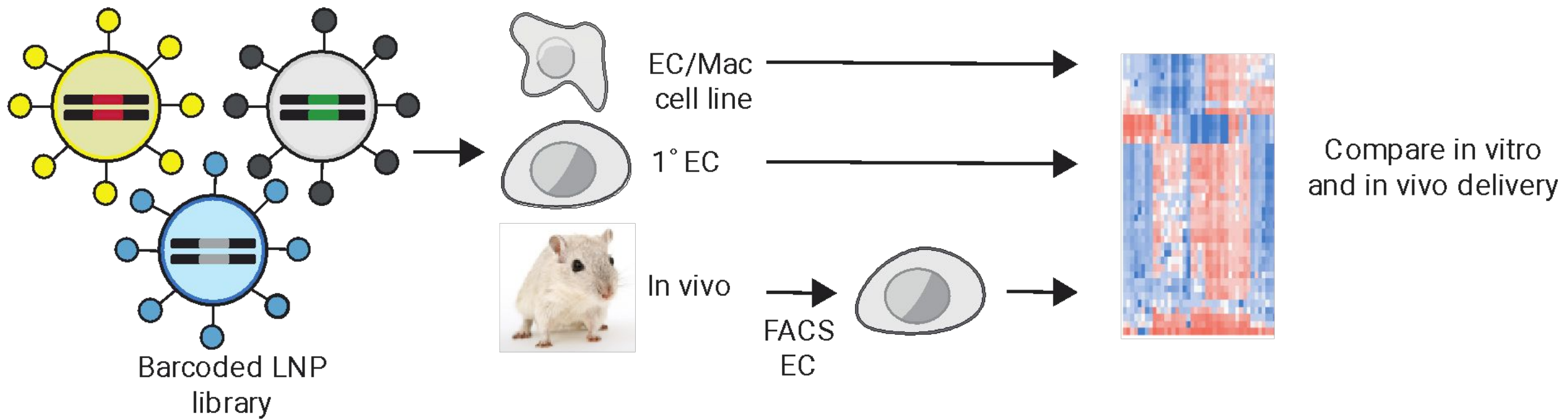


You find what you look for: *in vitro* / *in vivo*, mouse vs. NHP, and route of administration

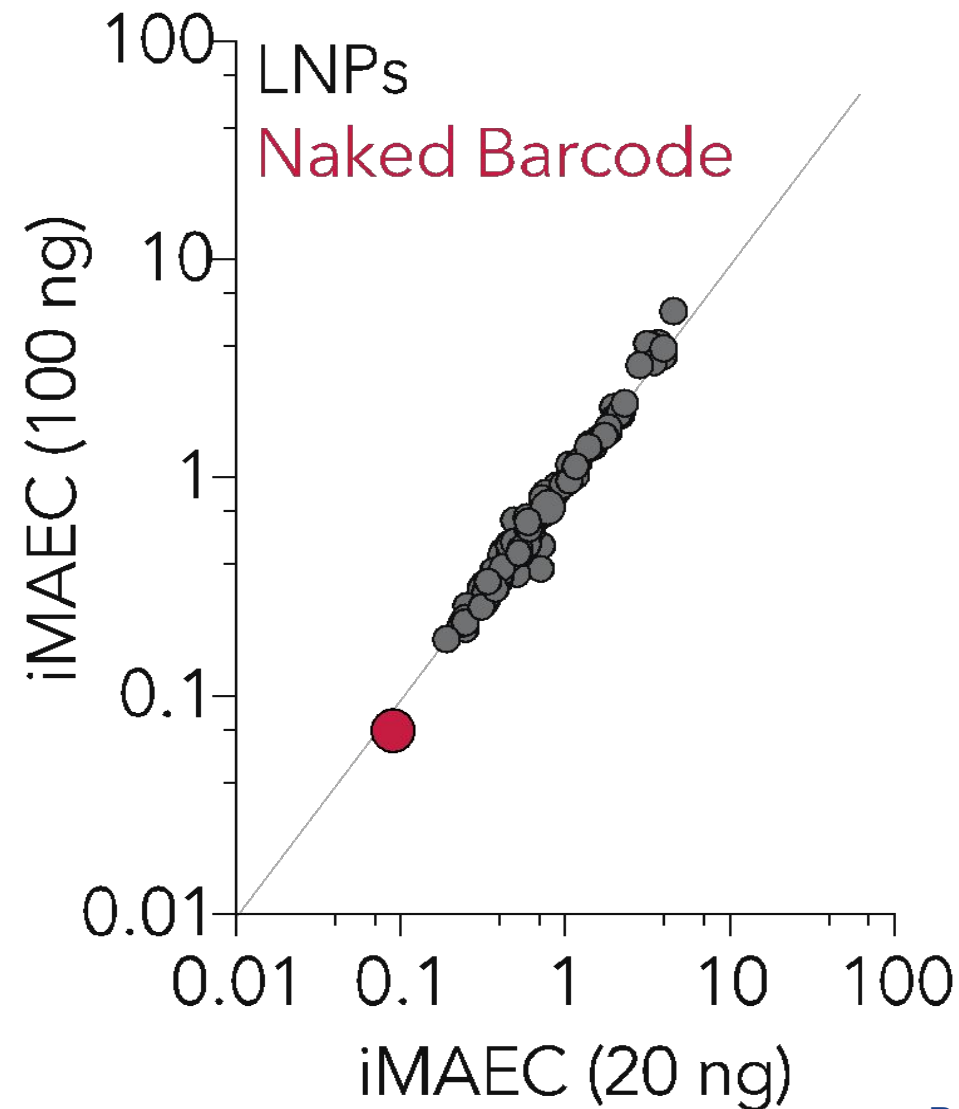
LNPs are typically screened *in vitro*



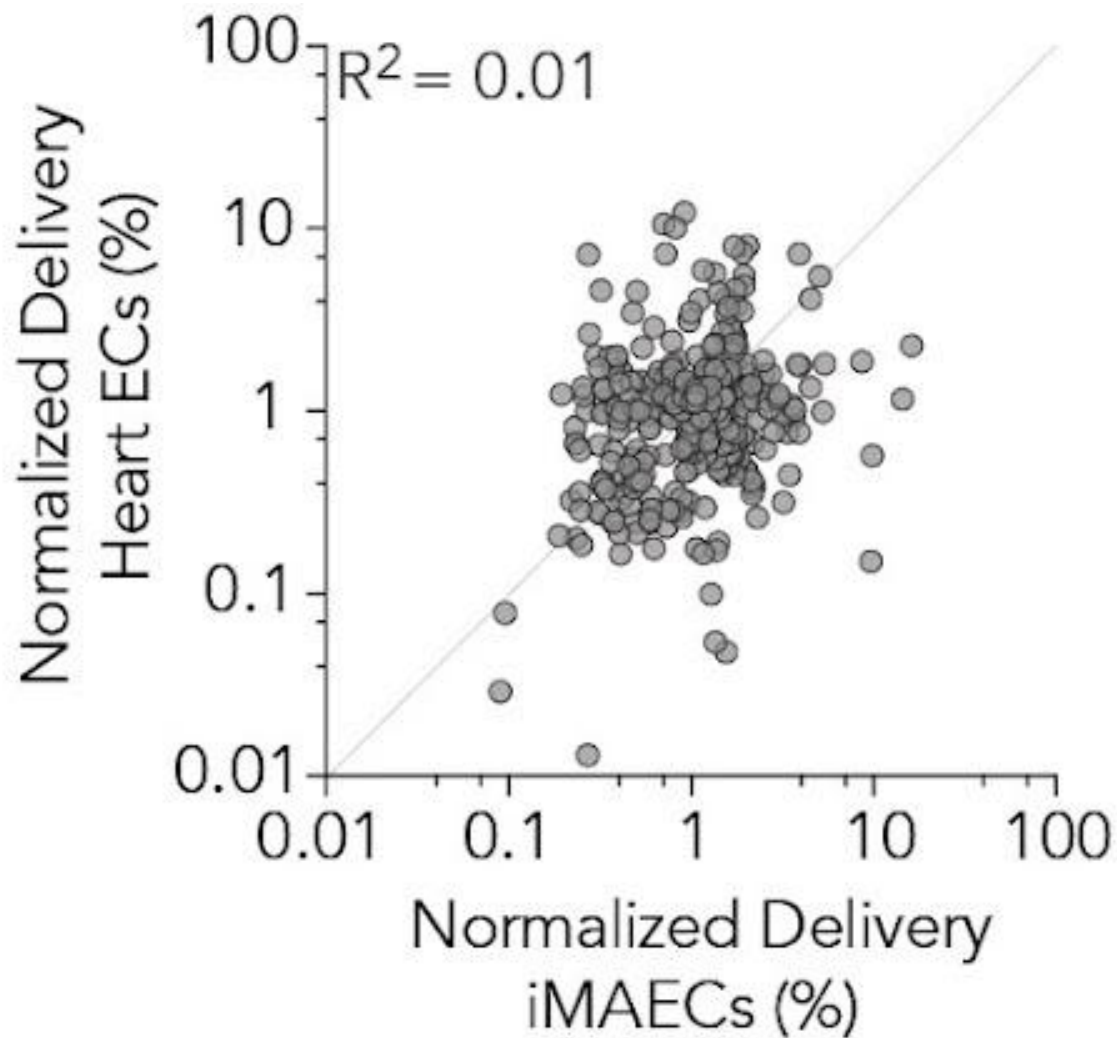
Does *in vitro* delivery predict *in vivo* delivery?



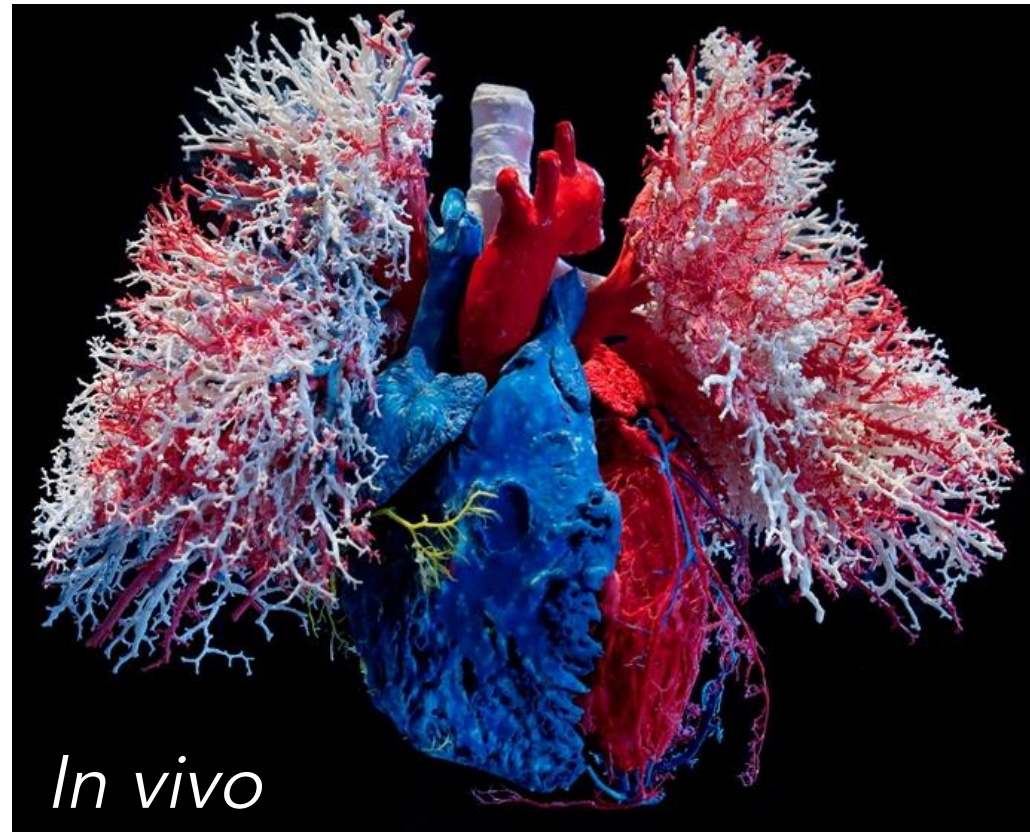
Control: *in vitro* delivery predicts *in vitro* delivery



In vitro delivery does not predict *in vivo* delivery

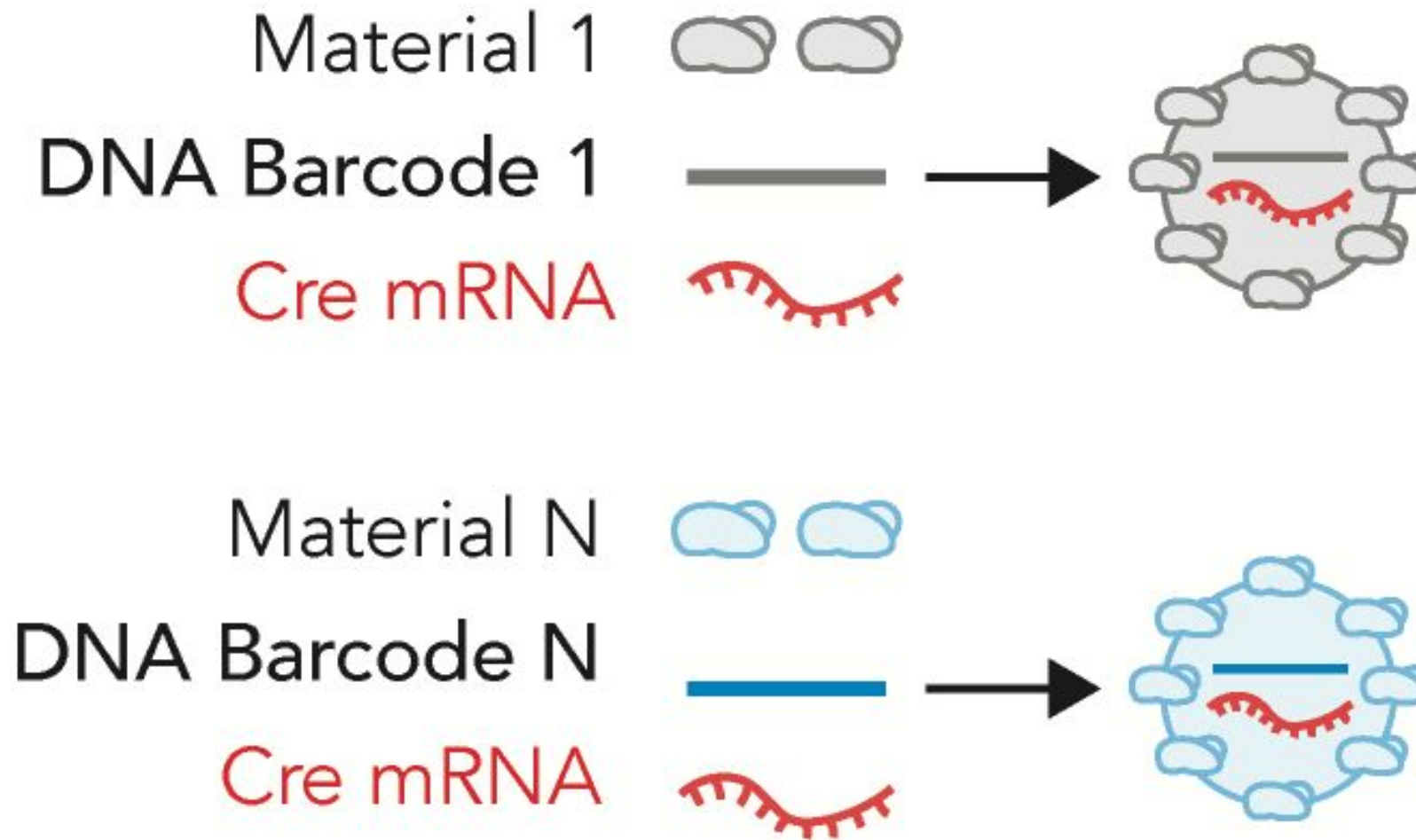


This is an opportunity to meaningfully improve LNP discovery



We designed assays to readout functional delivery

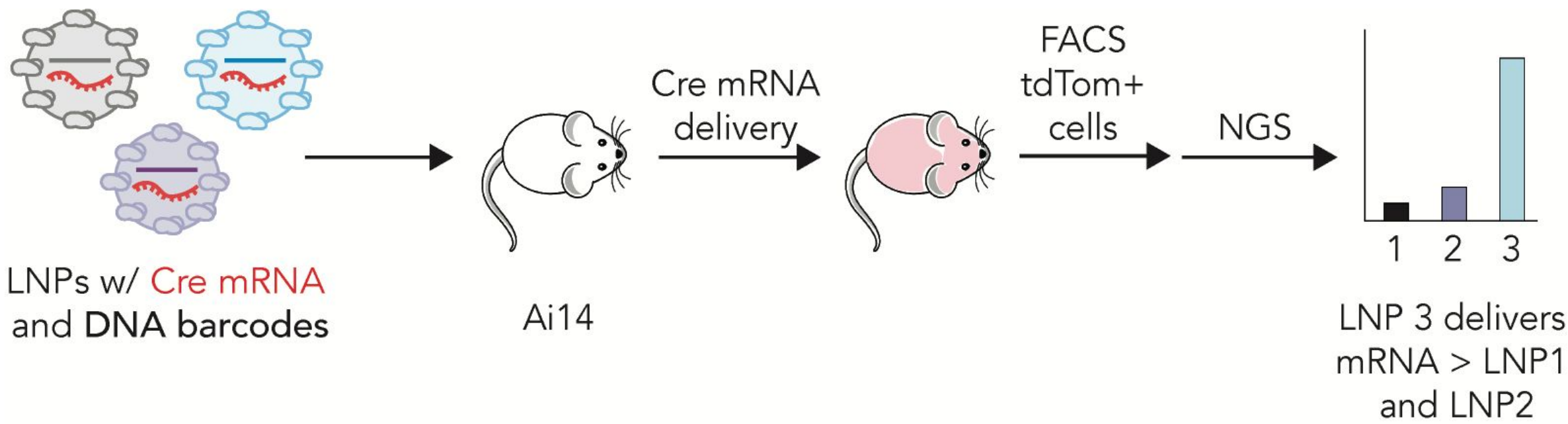
We can screen for Cre mRNA delivery *in vivo*



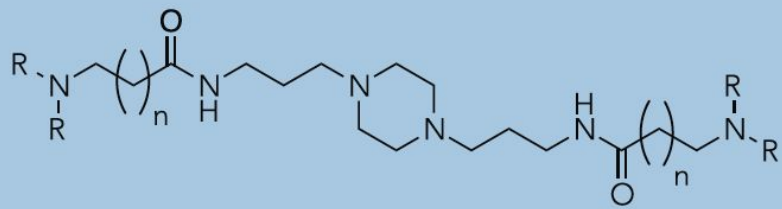
Fast Identification of Nanoparticle Delivery (FIND)

Control: Lox – Stop – Lox - tdTomato

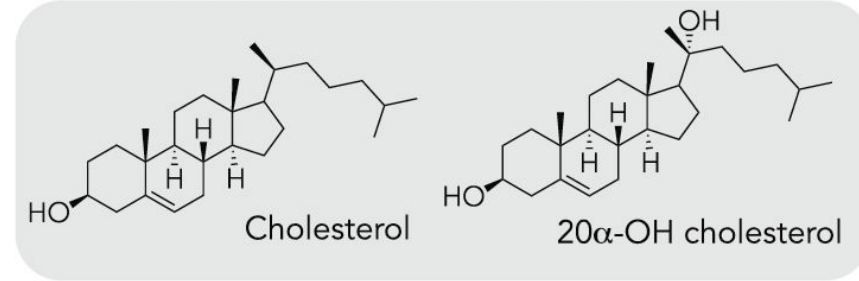
Control: **Lox - tdTomato**



FIND identified piperazine-based LNPs for immune delivery



Scaffold	Tail ID	R tails
PPZ-A (n = 1)	10	
	11	
	12	
	18-2Z	
PPZ-B (n = 2)	10	
	11	
	12	
	18-2Z	



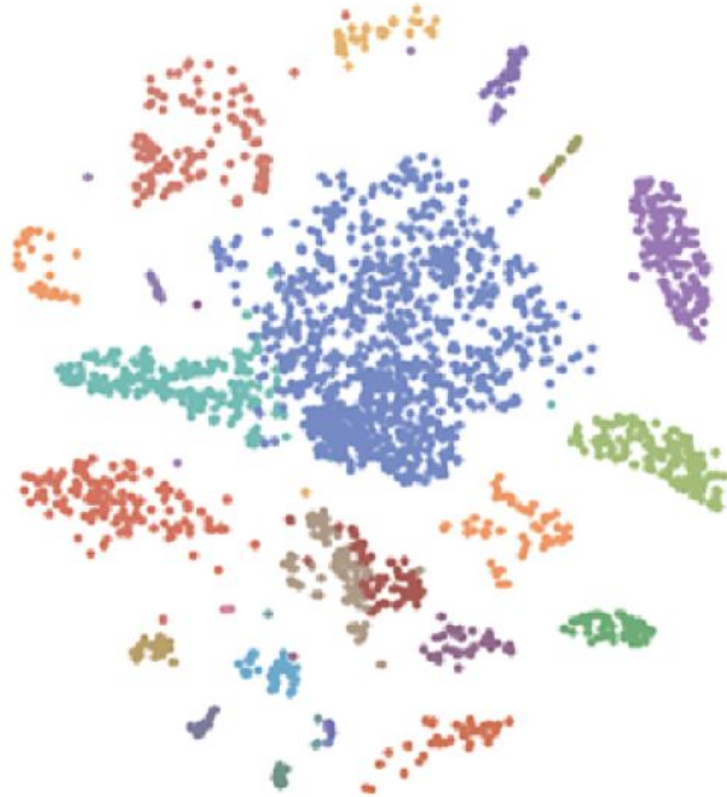
C₁₄PEG_{2K}

C₁₈PEG_{2K}

DOPE

Molar Ratio				
Ionizable Lipid	30	35	45	50
Cholesterol	30	46.5	39.5	35
PEG-Lipid	1	2.5	2.5	2.5
Helper Lipid	39	16	13	12.5

FIND also identified LNPs that deliver across lung










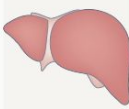

Cell type



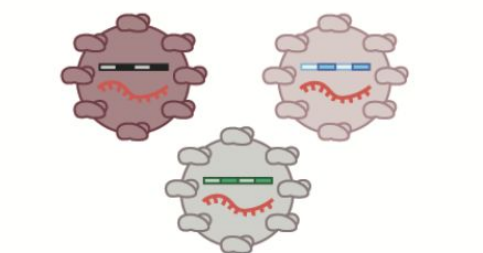
Delivered mRNA

You find what you look for: *in vitro* / *in vivo*, mouse vs. NHP, and route of administration

The species probably matters

	Mice	% of bw	Rats	% of bw	NHPs	% of bw	Humans	% of bw
Brain		2.1		0.9		1.4		1.9
Heart		0.6		0.4		0.4		0.4
Lung		0.7		0.5		0.6		1.0
Liver		4.5		3.0		2.2		2.0
Spleen		0.4		0.2		0.1		0.2
Kidney		1.4		0.8		0.4		0.3







We built a species-agnostic barcoding system



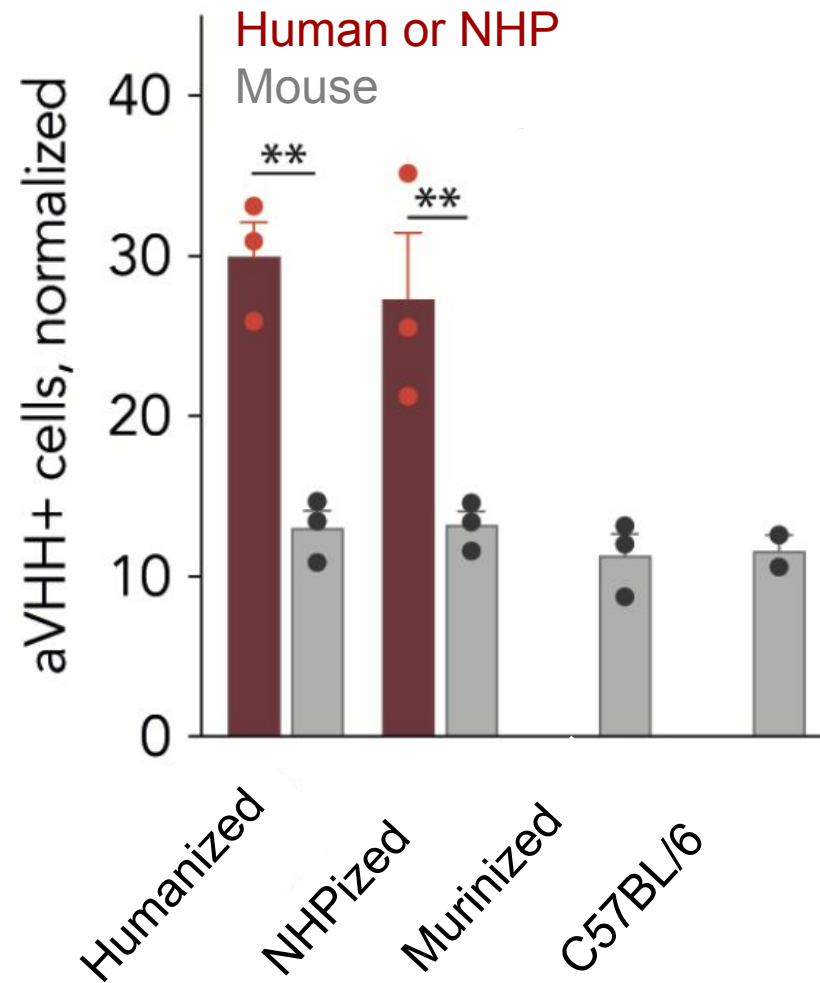
Nanoparticle N carries
Barcode N, aVHH mRNA



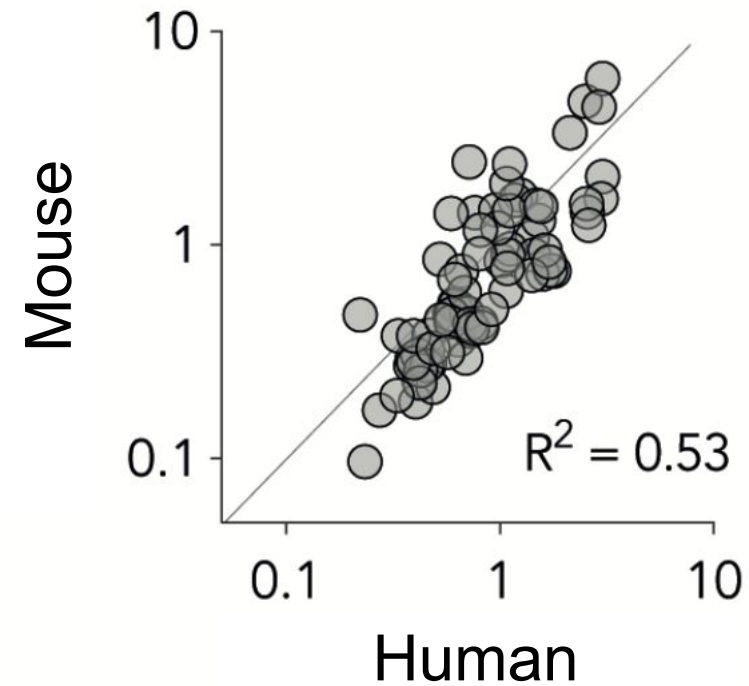
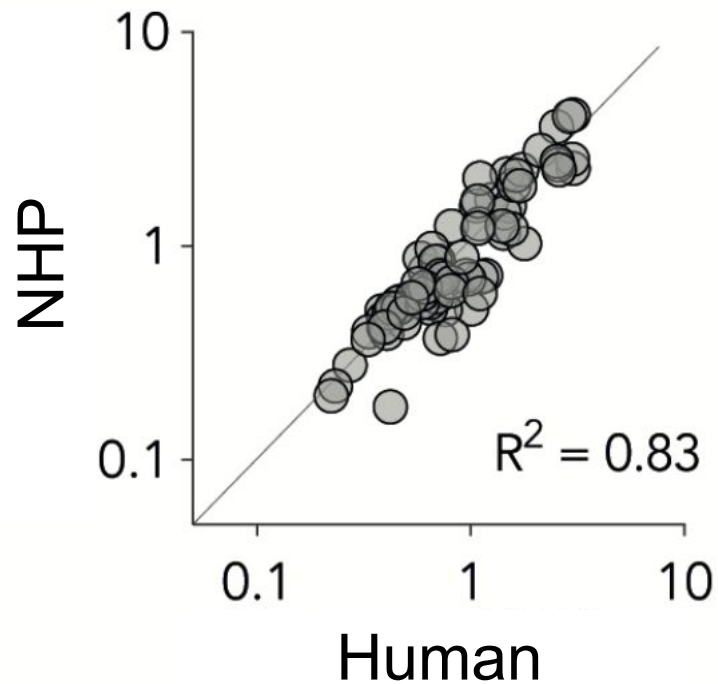
Hepatocyte
repopulated mice

<i>Yecuris FRG® KO</i> on C57Bl/6	 Human	 Primate	 Murine
Repopulated Hep. FACS anti-human CD47 ⁺	Humanized/H	Primatized/P	
Resident Hep. FACS CD31 ⁻ CD45 ⁻	Humanized/M	Primatized/M	Murinized/M
<i>Common mouse strains</i>	 C57BL/6J	 BALB/cJ	 NZB/BINJ
Resident Hep. FACS CD31 ⁻ CD45 ⁻	BL/6	Balb/C	NZB

We observed species-dependent delivery



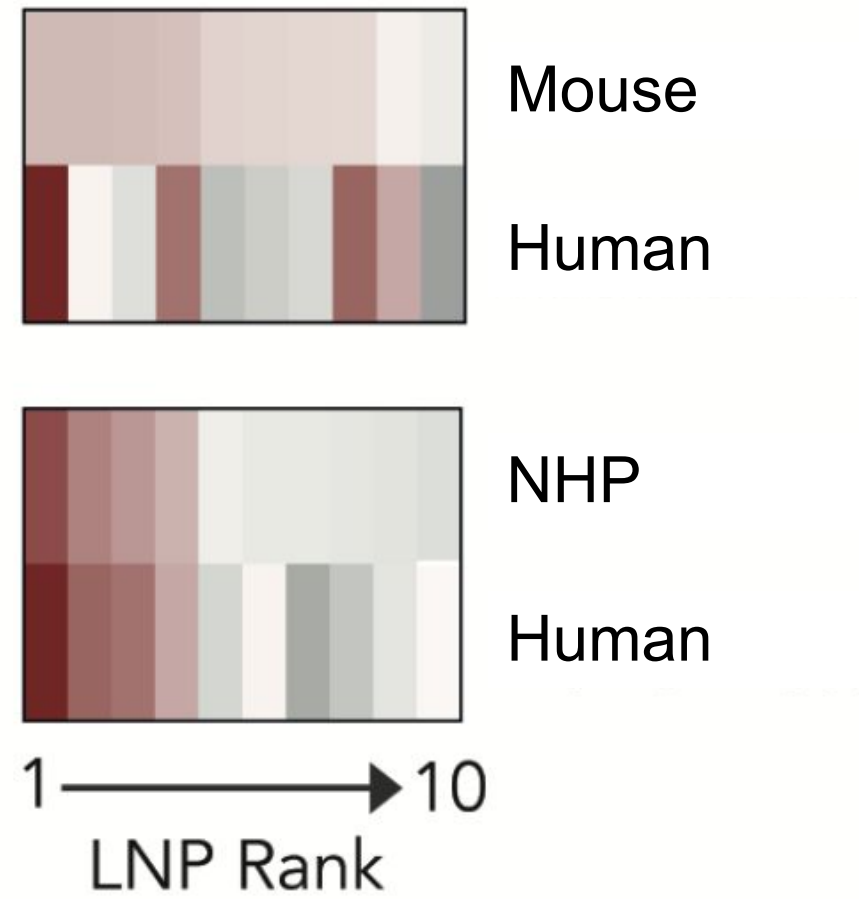
Delivery in primate cells predict human > mice



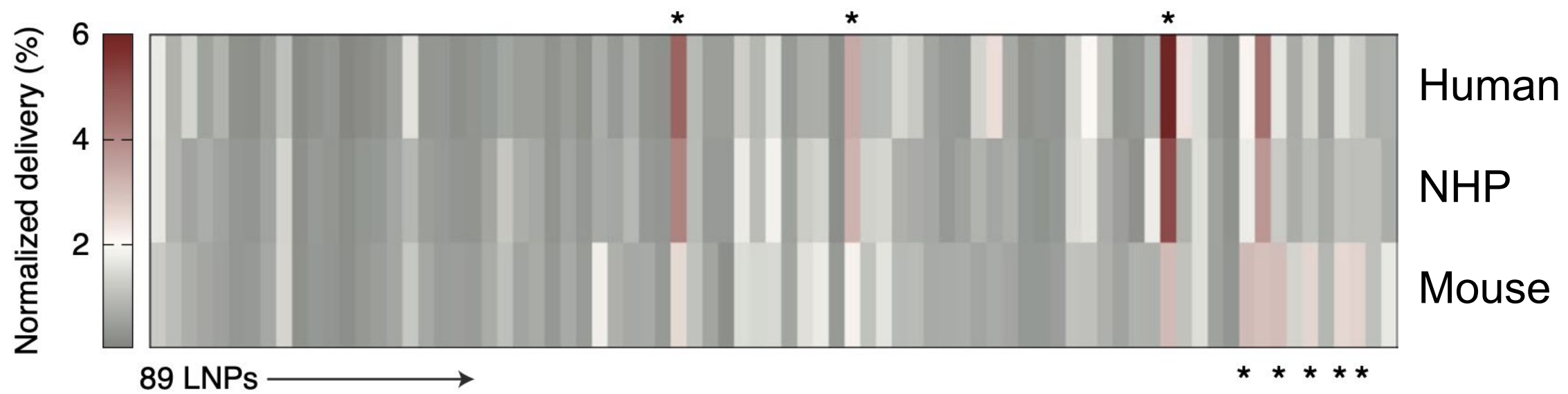
Mouse LNP readouts can give false positives / negatives



Mouse LNP readouts can give false positives / negatives



Mouse gives you false positives and false negatives



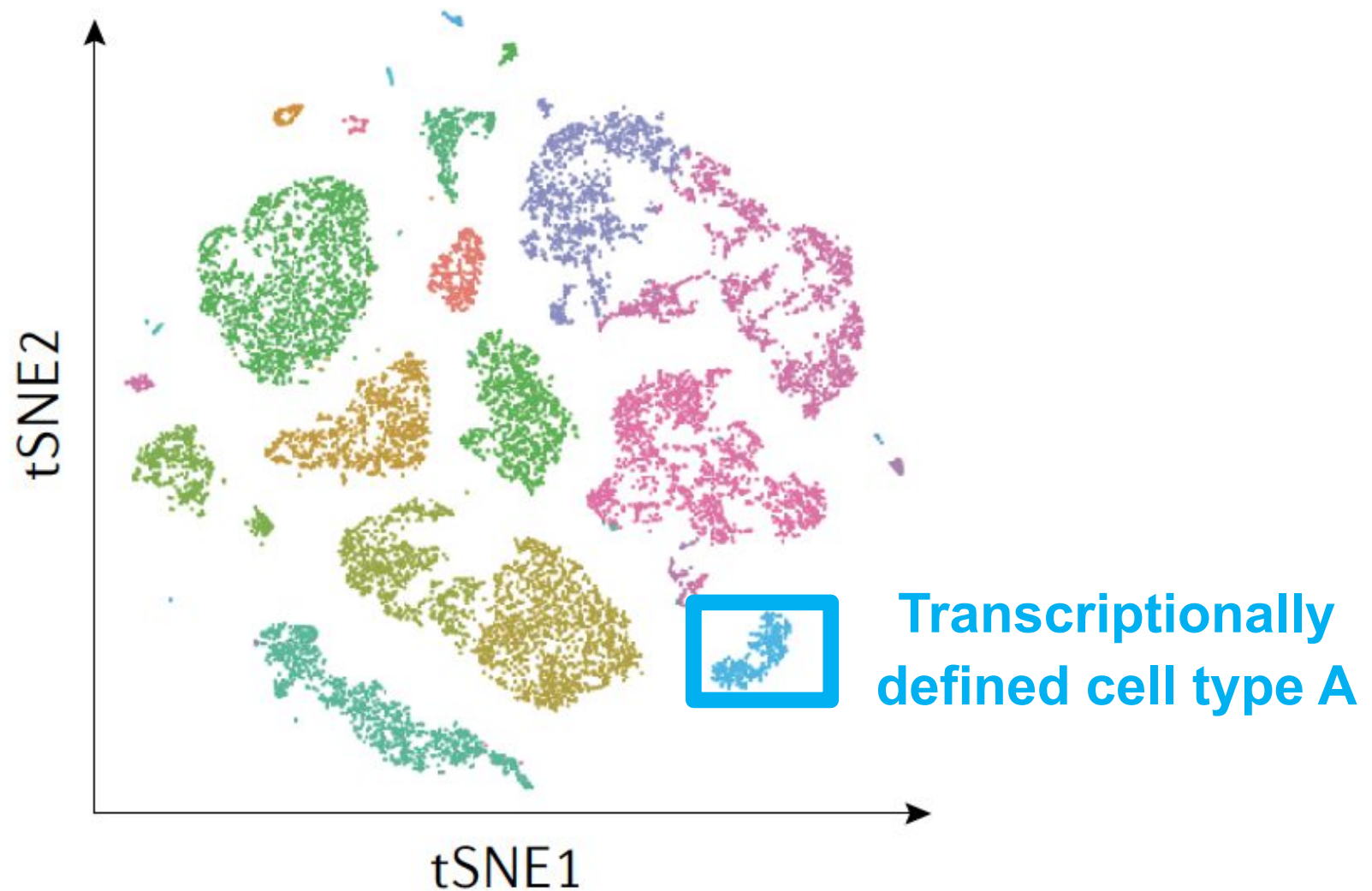
An ideal LNP screen would tell you...

1) Where did (many) LNPs go?

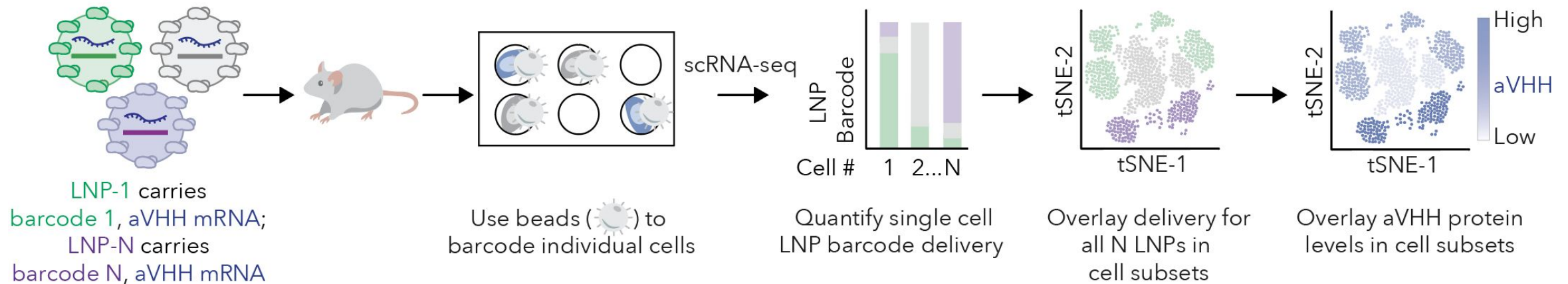
2) Where did (many) LNPs work?

... and it would do so with single cell resolution.

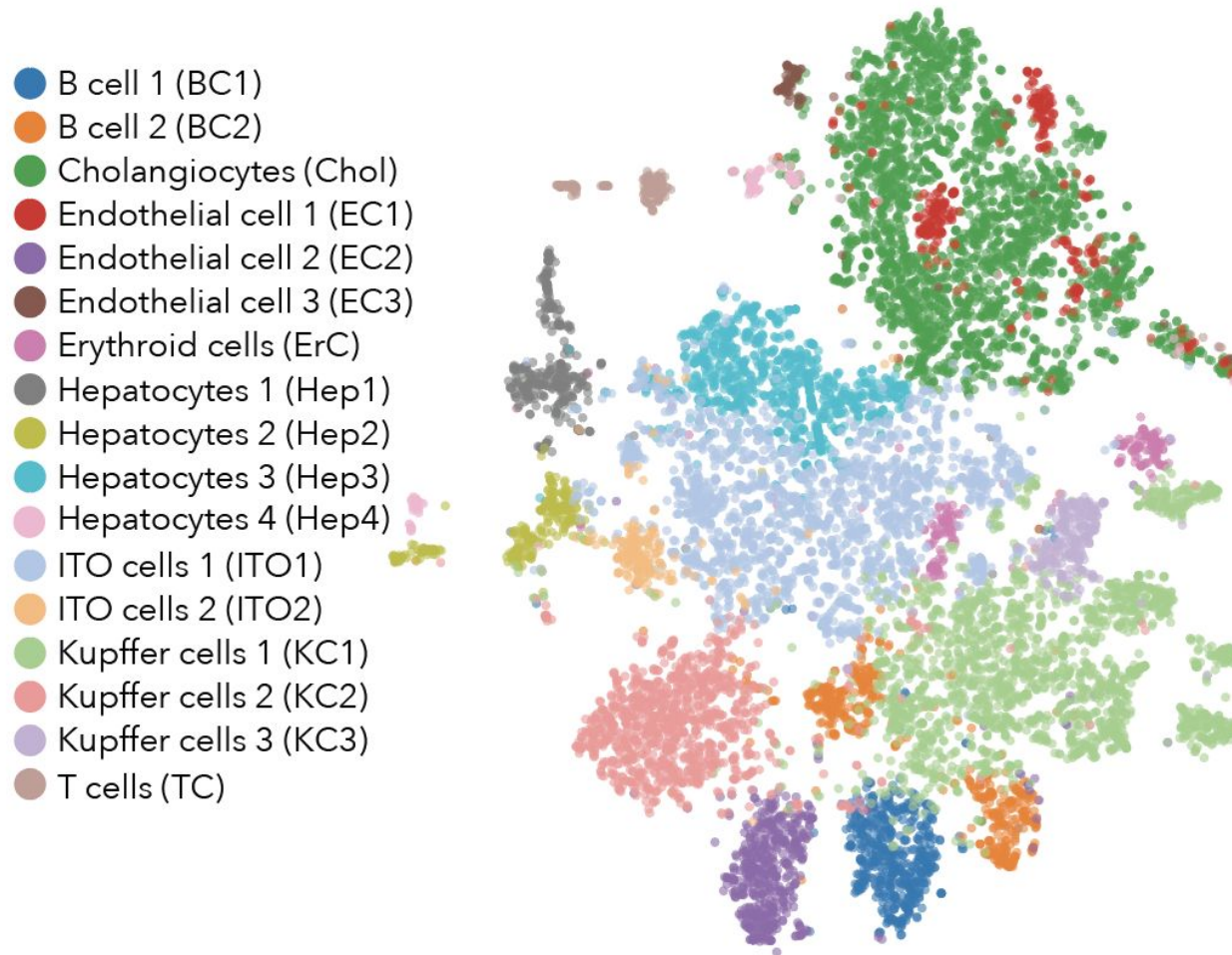
Single-cell RNA-sequencing gives you cell identity



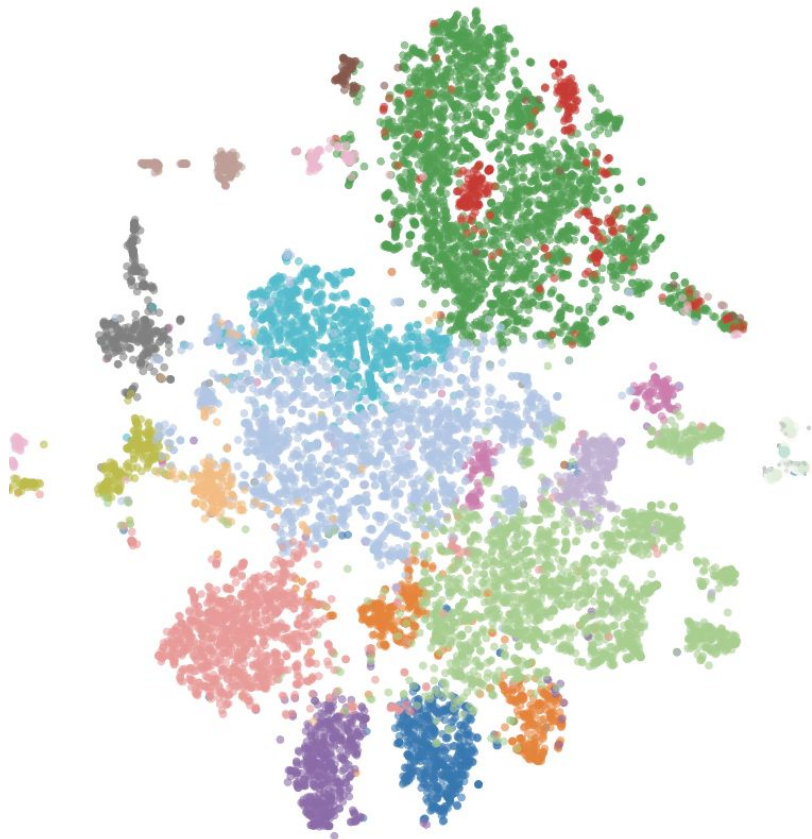
SENT-seq gives you barcode, protein, transcriptome



scRNA-seq = transcriptionally defined cell subtypes



Overlay protein delivery and barcodes onto single cells



Cell type



Protein delivery



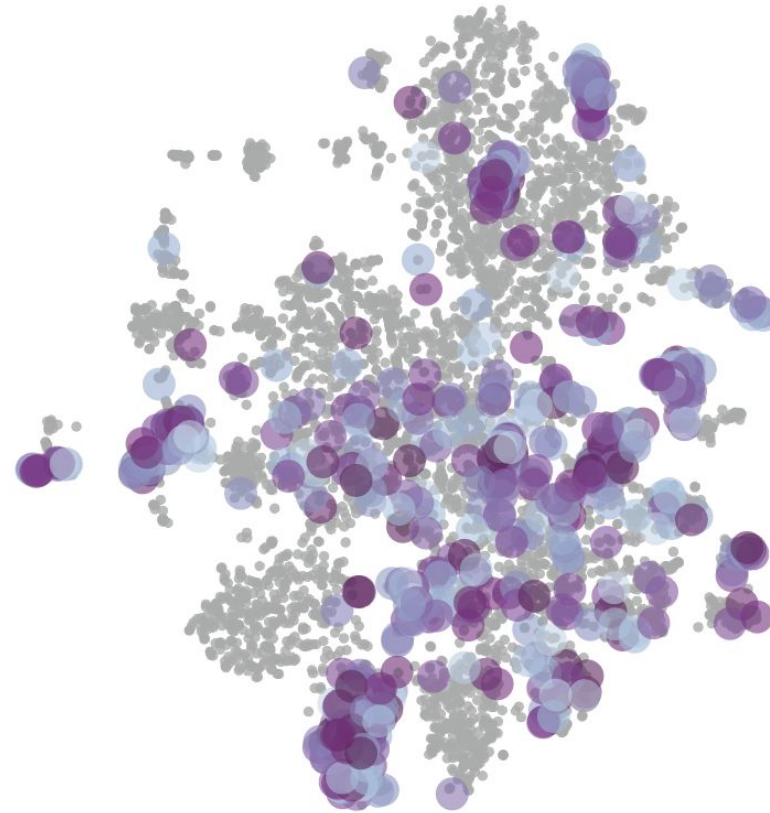
Barcodes

This lets us measure LNP tropism in any single cell(s) you want

.....

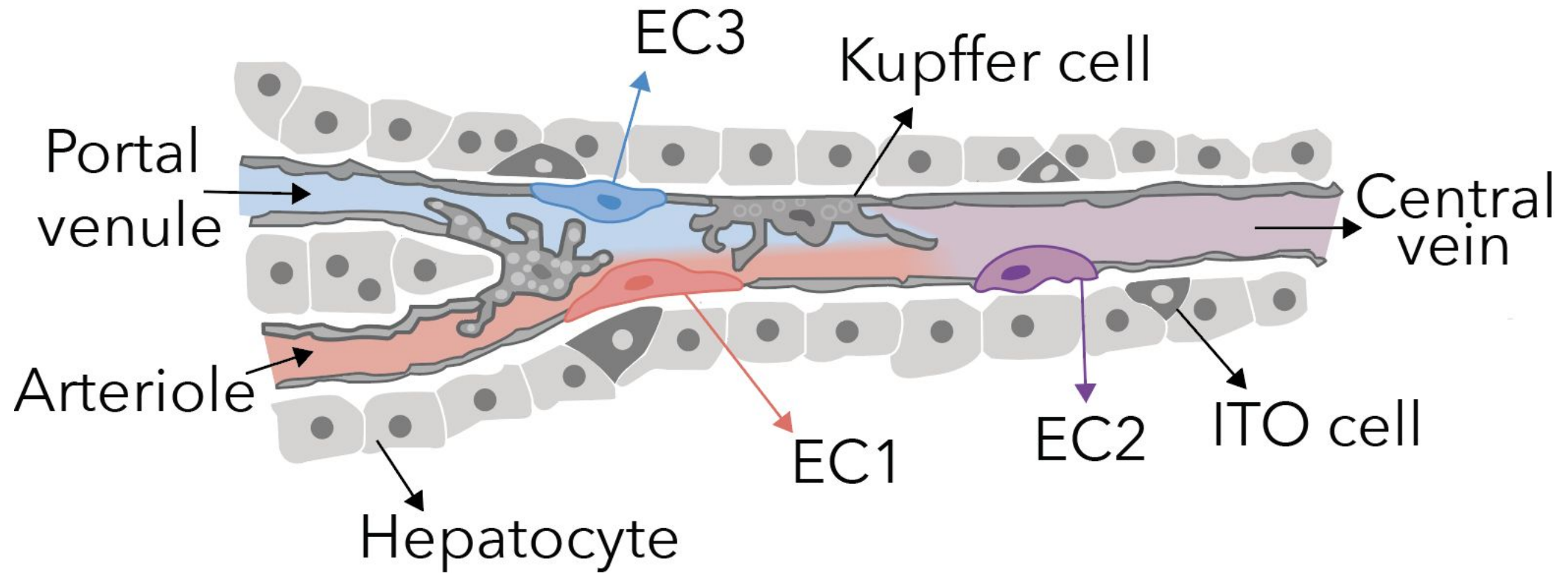


LNP-10

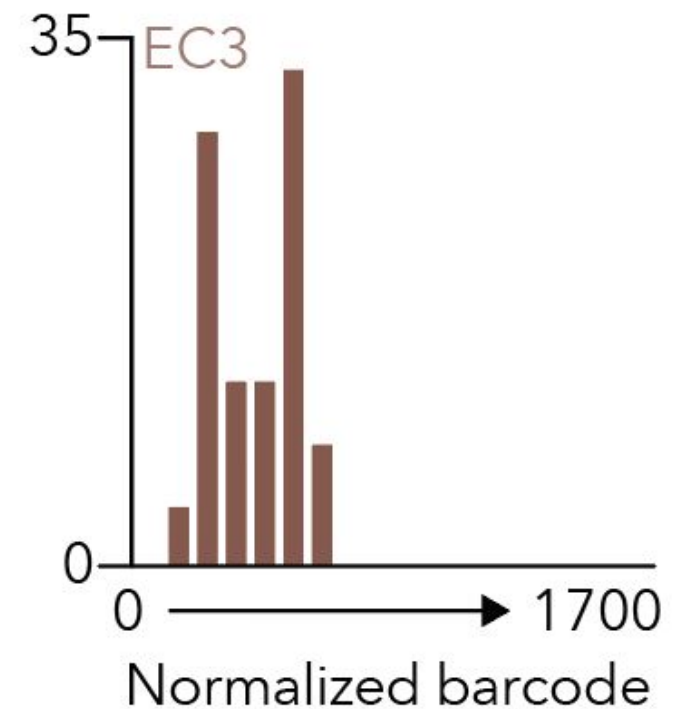
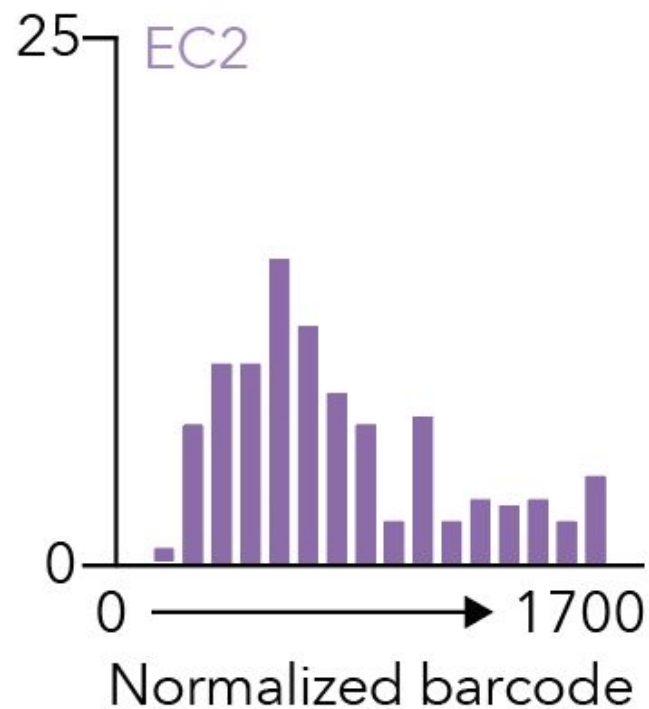
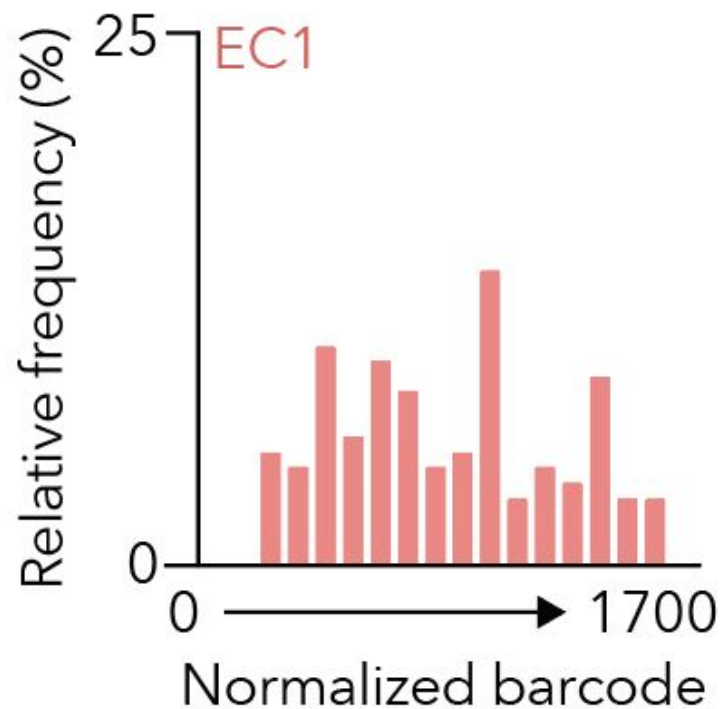


LNP-12

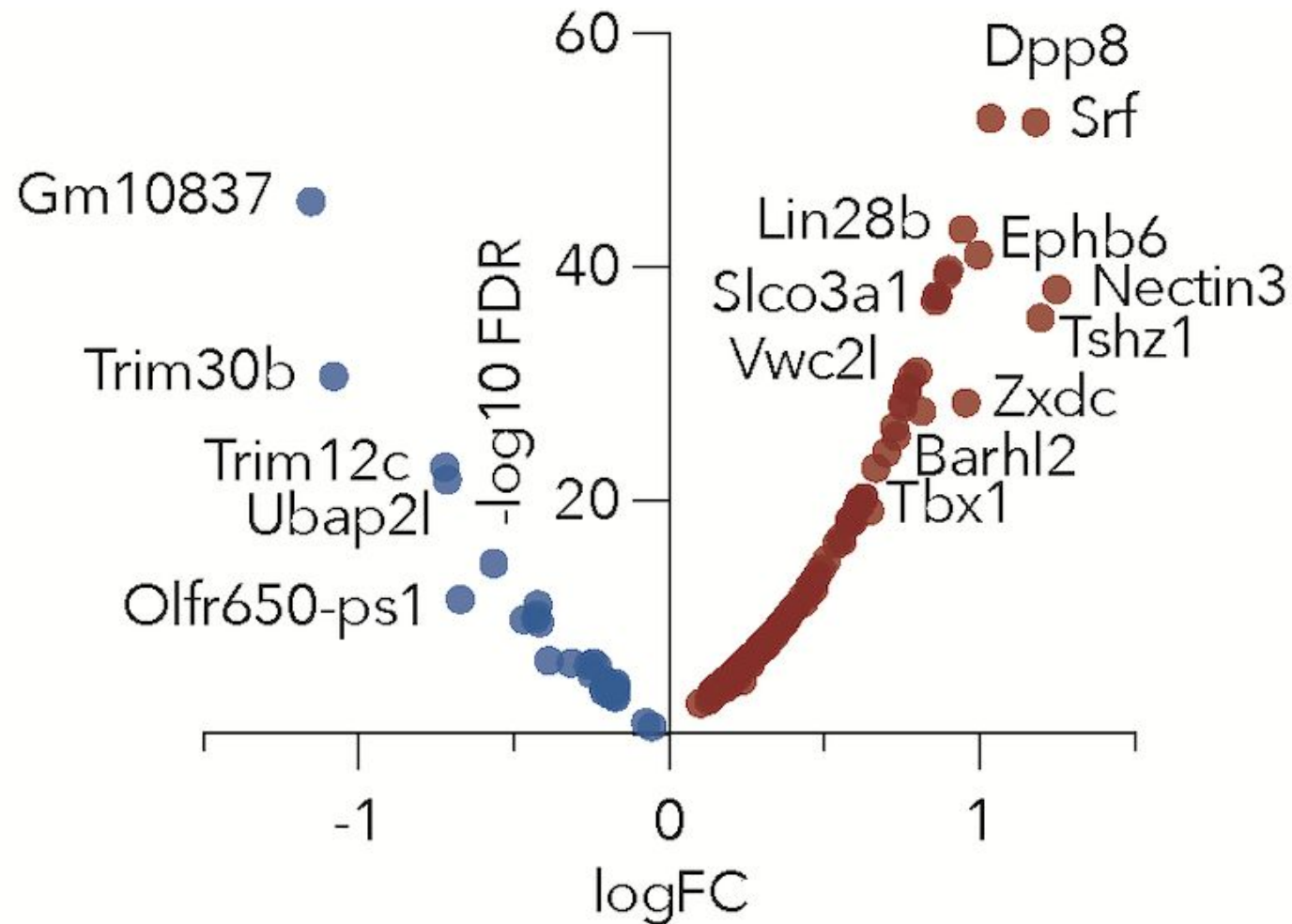
Looking at on / off-target in cell subtypes



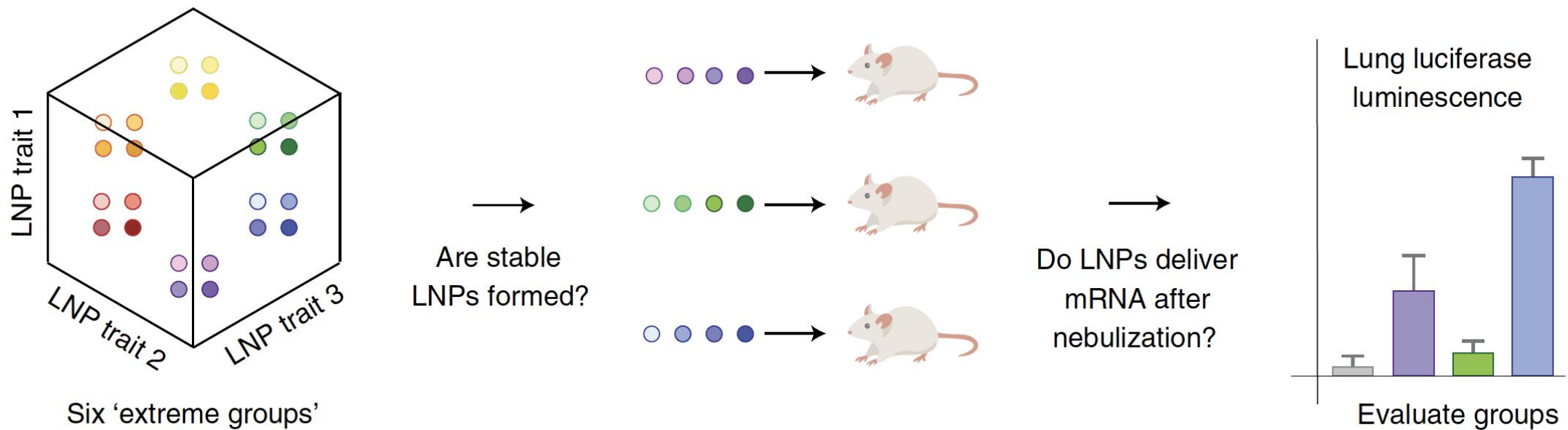
We can measure delivery distributions in single cells



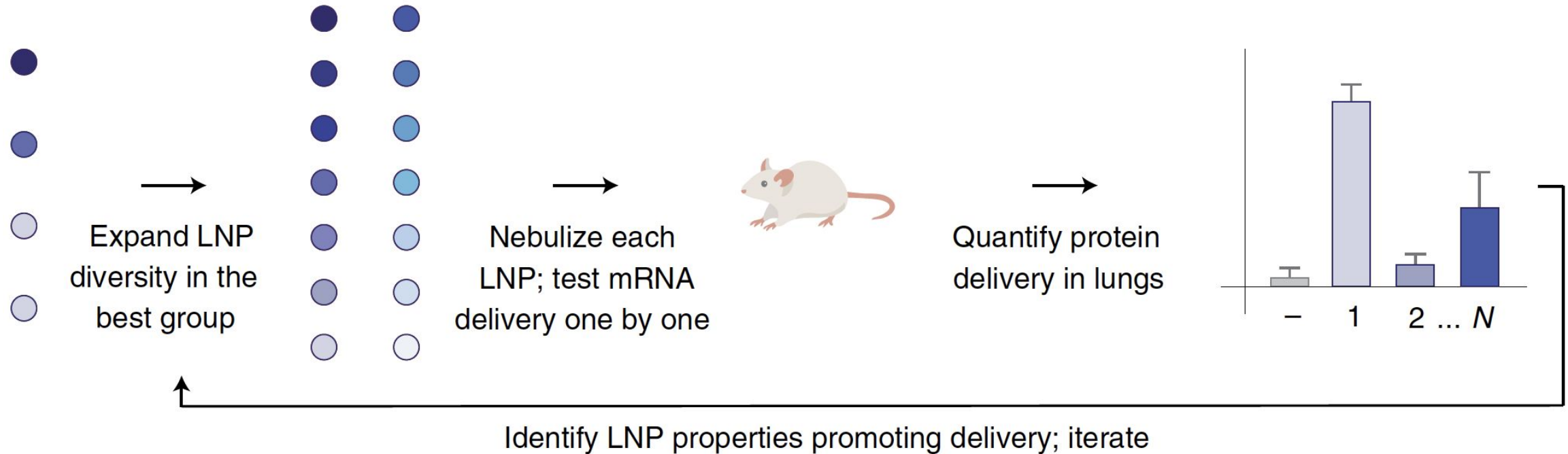
Transcriptional signatures in LNP^{High} vs LNP^{Low} cells



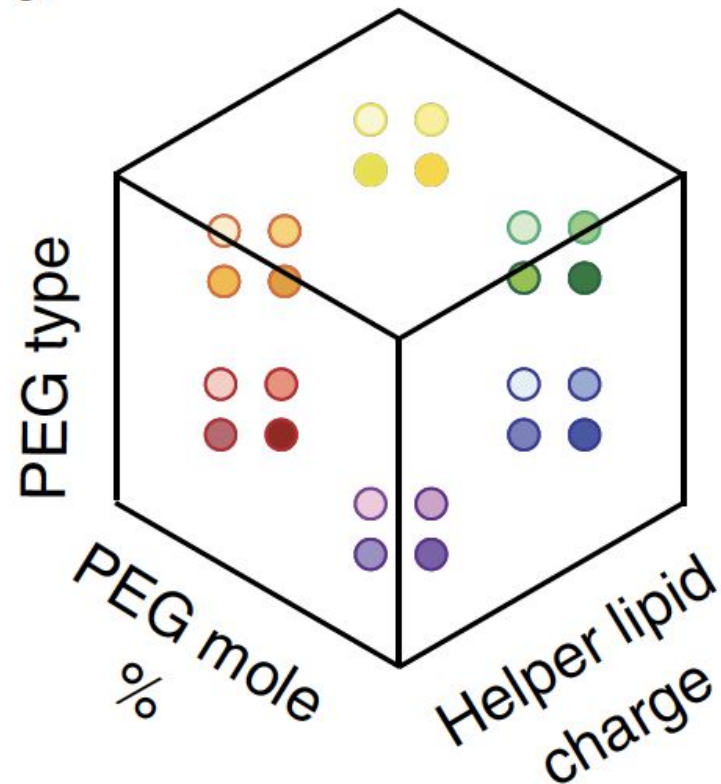
Finally, we screen across wide chemical spaces



After a wide LNP screen, we “zoom in”



This approach led us to nebulization LNP design rules



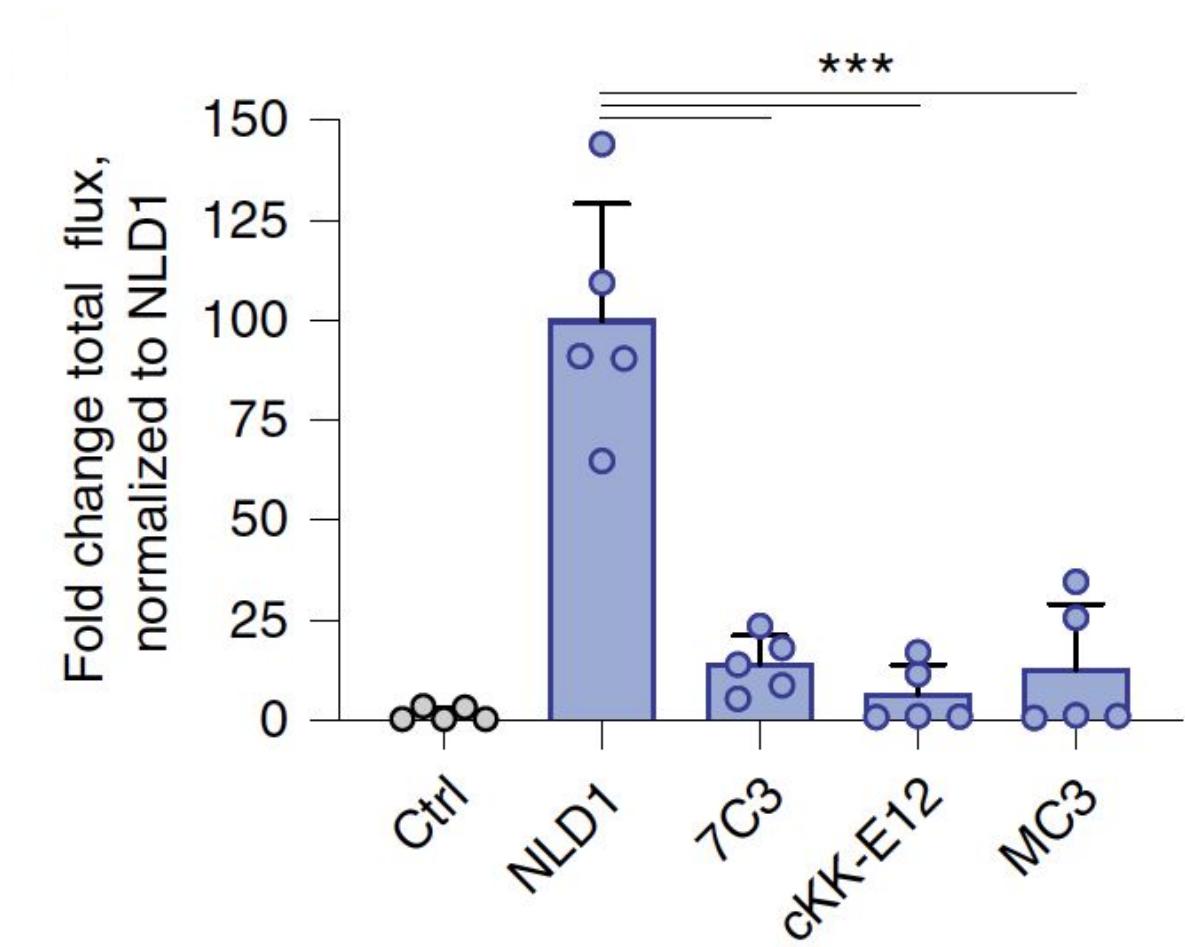
110 LNPs formulated

54 nebulized/in vivo

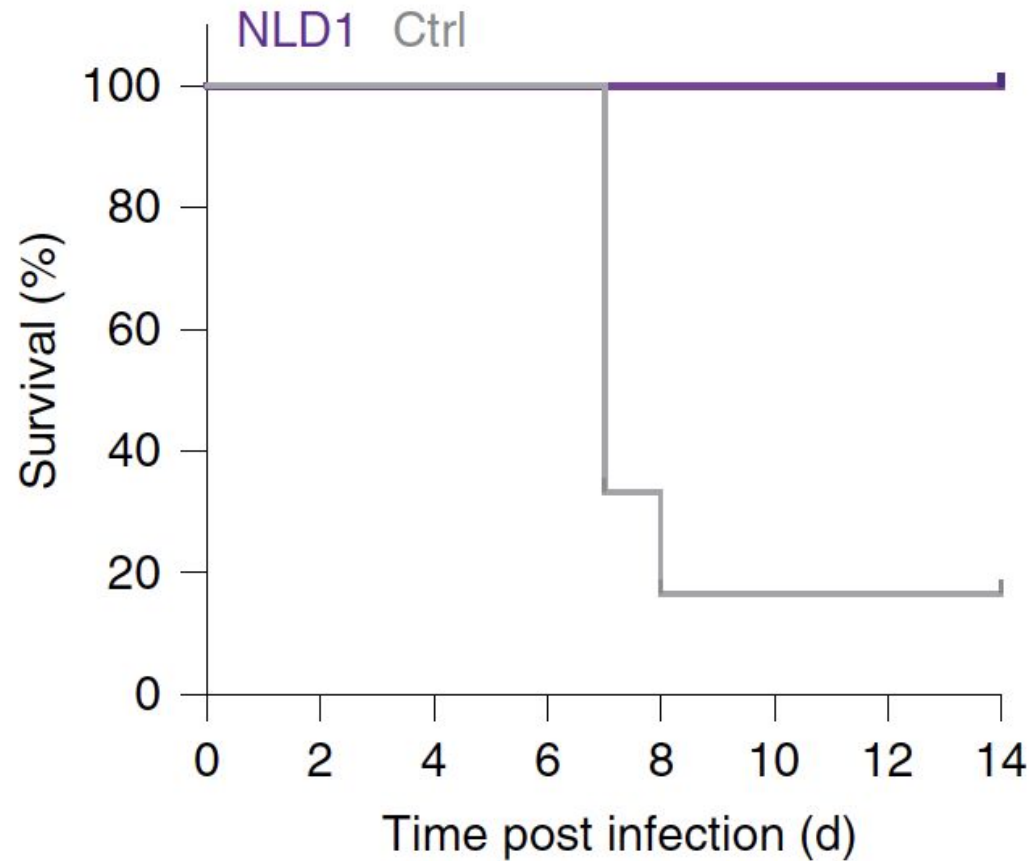
PEG is important for
nebulized lung LNP delivery

- 1) PEG is necessary for LNP stability
- 2) Helper lipids and PEG interact
 - (a) Neutral lipid + low PEG
 - (b) Cationic lipid + high PEG

Nebulized LNPs designs is not the same as systemic LNPs



NLD1 can deliver mRNA protecting against fatal influenza



What have we learned?

Most lipids are unexplored.

Keep LNP structure as simple as possible.

You find what you look for, so run predictive assays.

Sincere thanks to collaborators, funders, trainees!



Phil Santangelo



Eric Sorscher



Hanjoong Jo



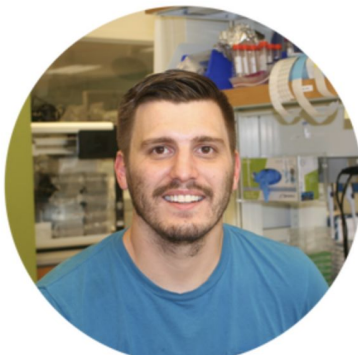
Jeffrey Glenn



Kalina Paunovska



Melissa Lokugamage



Curtis Dobrowolski



Marine Hatit



CMT
Research
Foundation



Two additional delivery papers

nature chemistry

Article

<https://doi.org/10.1038/s41557-023-01138-9>

Nanoparticle stereochemistry-dependent endocytic processing improves in vivo mRNA delivery

nature
biomedical engineering

ARTICLES

<https://doi.org/10.1038/s41551-022-00847-9>



Augmented lipid-nanoparticle-mediated in vivo genome editing in the lungs and spleen by disrupting Cas9 activity in the liver