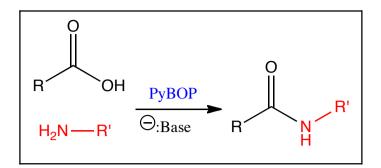
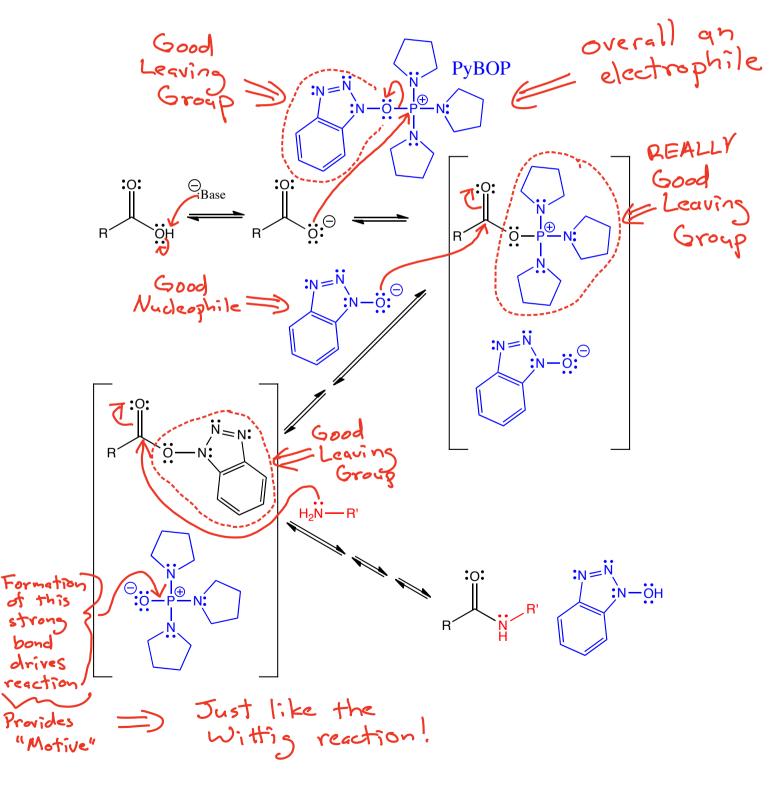


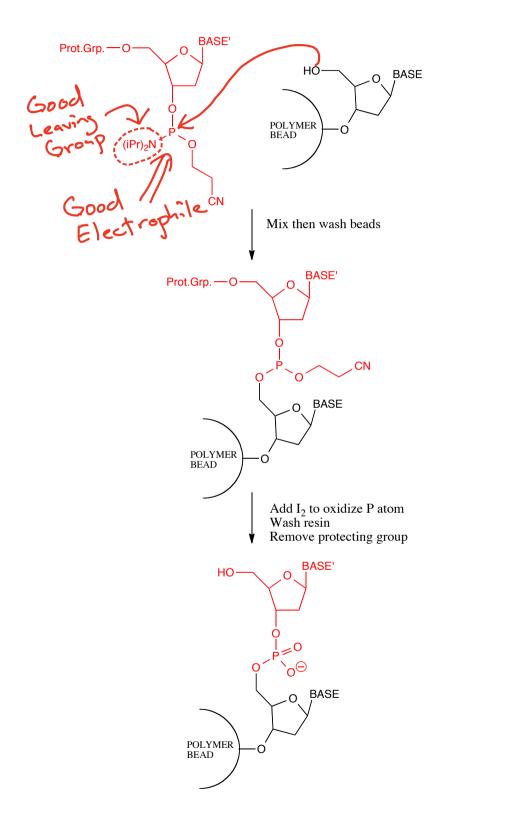
Repeat as necessary then remove from resin. Can add up to 100 amino acids this way.



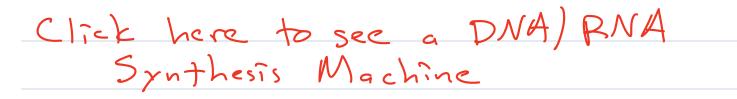


ঠ

Solid Phase Synthesis of DNA



• •

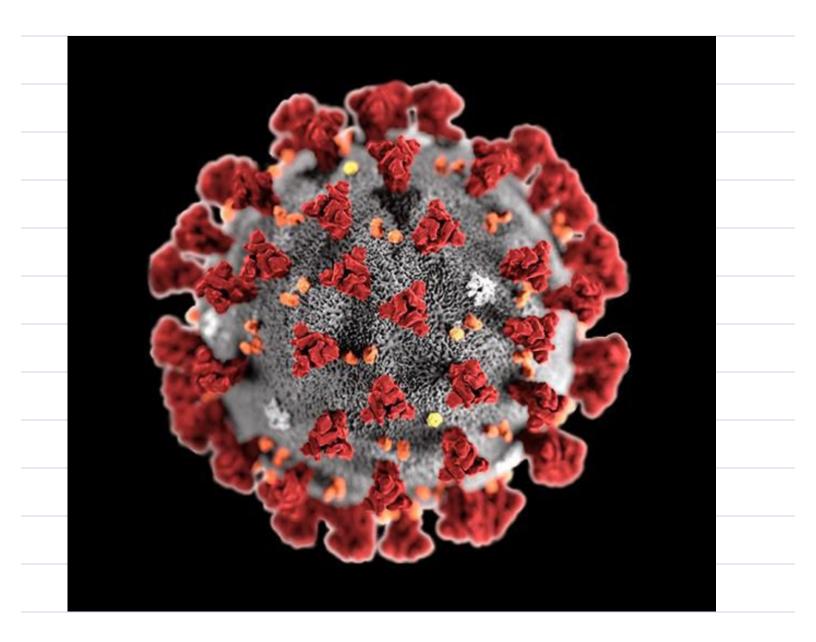


https://shop.biosearchtech.com/nucleic-acid-chemistry-reagents-and-instruments/dna-and-rna-synthesis/dna-and-rna-

oligonucleotide-synthesizers/mermade-6-and-12/c/mermade-6-and-12

Click here to see how to order DNA

https://www.idtdna.com/pages/products/custom-dna-rna/dna-oligos



New UT cases

The University of Texas said two people in its community, an unnamed student and the Dean of **Undergraduate Studies** Brent Iverson, had tested positive for the virus. It is unclear if the pair are included in Travis County's count of confirmed cases.



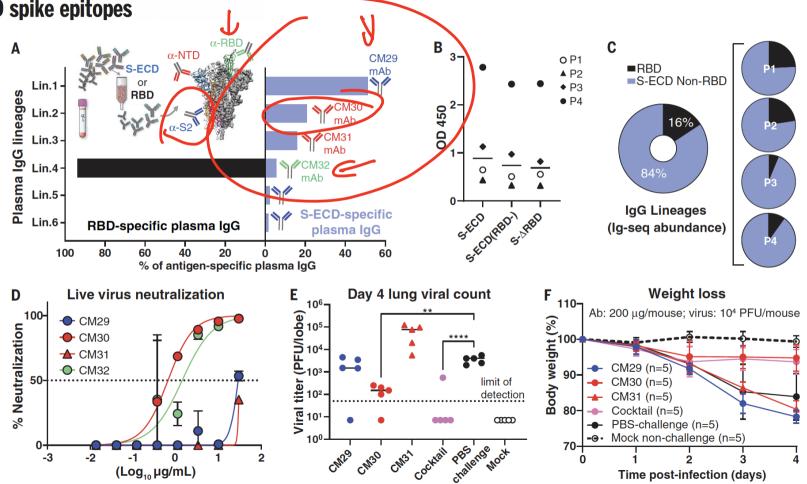
CORONAVIRUS

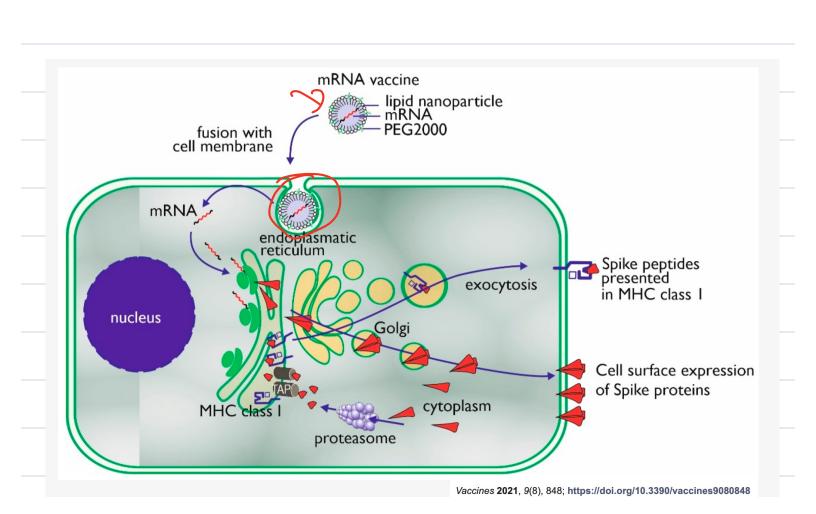
Prevalent, protective, and convergent IgG recognition of SARS-CoV-2 non-RBD spike epitopes

William N. Voss¹, Yixuan J. Hou^{2,#}, Nicole V. Johnson^{1,#}, George Delidakis³, Jin Eyun Kim⁴, Kamyab Javanmardi¹, Andrew P. Horton¹, Foteini Bartzoka¹, Chelsea J. Paresi⁵, Yuri Tanno³, Chia-Wei Chou¹, Shawn A. Abbasi⁶, Whitney Pickens¹, Katia George¹, Daniel R. Boutz^{1,7}, Dalton M. Towers³, Jonathan R. McDaniel⁸, Daniel Billick¹, Jule Goike¹, Lori Rowe^{9,10}, Dhwani Batra⁹, Jan Pohl⁹, Justin Lee⁹, Shivaprakash Gangappa¹¹, Suryaprakash Sambhara¹¹, Michelle Gadush¹², Nianshuang Wang¹, Maria D. Person¹², Brent L. Iverson⁵, Jimmy D. Gollihar^{1,7,13}, John M. Dye⁶, Andrew S. Herbert⁶, Ilya J. Finkelstein¹, Ralph S. Baric^{2,14}, Jason S. McLellan¹, George Georgiou^{1,3,4,15}, Jason J. Lavinder^{1,3*}, Gregory C. Ippolito^{1,13,15*} Voss *et al.*, *Science* **372**, 1108–1112 (2021) 4 June 2021

CORONAVIRUS

Prevalent, protective, and convergent IgG recognition of SARS-CoV-2 non-RBD spike epitopes





Structure-based design of prefusion-stabilized SARS-CoV-2 spikes

Ching-Lin Hsieh¹, Jory A. Goldsmith¹, Jeffrey M. Schaub¹, Andrea M. DiVenere², Hung-Che Kuo¹, Kamyab Javanmardi¹, Kevin C. Le², Daniel Wrapp¹, Aliso<u>n G. Lee¹</u>, Yutong Liu², Chia-Wei Chou¹, Patrick O. Byrne¹, Christy K. Hjorth¹, Nicole V. Johnson¹, John Ludes Meyers¹, Annalee W. Nguyen², Juyeon Park¹, Nianshuang Wang¹, Dzifa Amengor¹, Jason J. Lavinder^{1,2}, Gregory C. Ippolito^{1,3}, Jennifer A. Maynard^{2*}, Ilya J. Finkelstein^{1,**}, Jason S. McLellan^{1*}

۶

¹Department of Molecular Biosciences, University of Texas, Austin, TX 78712, USA. ²Department of Chemical Engineering, University of Texas, Austin, TX 78712, USA. ³Department of Oncology, Dell Medical School, University of Texas, Austin, TX 78712, USA. ⁴Center for Systems and Synthetic Biology, University of Texas, Austin, TX 78712, USA. ⁴Center for Systems and Synthetic Biology, University of Texas, Austin, TX 78712, USA. ⁴Center for Systems and Synthetic Biology, University of Texas, Austin, TX 78712, USA. ⁴Center for Systems and Synthetic Biology, University of Texas, Austin, TX 78712, USA. ⁴Center for Systems and Synthetic Biology, University of Texas, Austin, TX 78712, USA.

*Corresponding author. Email: maynard@she.utexas.edu (J.A.M.); Ilya@finkelsteinlab.org (I.J.F.); jmclellan@austin.utexas.edu (J.S.M.)

- The COVID-19 pandemic has led to accelerated efforts to develop therapeutics and vaccines. A key target
 of these efforts is the spike (S) protein, which is metastable and difficult to produce recombinantly. Here, we characterized 100 structure-guided spike designs and identified 26 individual substitutions that increased protein yields and stability. Testing combinations of beneficial substitutions resulted in the
 identification of HexaPro, a variant with six beneficial proline substitutions exhibiting ~10-fold higher expression than its parental construct and the ability to withstand heat stress, storage at room temperature, and three freeze-thaw cycles. A 3.2 Å-resolution cryo-EM structure of HexaPro confirmed
- that it retains the prefusion spike conformation. High-yield production of a stabilized prefusion spike protein will accelerate the development of vaccines and serological diagnostics for SARS-CoV-2.

Incorporation of Pseudouridine Into mRNA Yields Superior Nonimmunogenic Vector With Increased Translational Capacity and Biological Stability

Katalin Kariko¹, Hiromi Muramatsu¹, Frank A Welsh¹, János Ludwig², Hiroki Kato³, Shizuo Akira³, and Drew Weissman⁴

¹Department of Neurosurgery, University of Pennsylvania, Philadelphia, Pennsylvania, USA

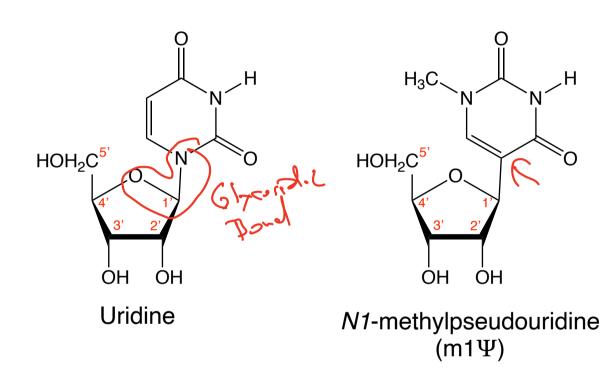
²Laboratory of RNA Molecular Biology, The Rockefeller University, New York, New York, USA

³Department of Host Defense, Research Institute for Microbial Diseases, Osaka University, Osaka, Japan

⁴Department of Medicine, University of Pennsylvania, Philadelphia, Pennsylvania, USA

Abstract

In vitro-transcribed mRNAs encoding physiologically important proteins have considerable potential for therapeutic applications. However, in its present form, mRNA is unfeasible for clinical use because of its labile and immunogenic nature. Here, we investigated whether incorporation of naturally modified nucleotides into transcripts would confer enhanced biological properties to mRNA. We found that mRNAs containing pseudouridines have a higher translational capacity than unmodified mRNAs when tested in mammalian cells and lysates or administered intravenously into mice at 0.015–0.15 mg/kg doses. The delivered mRNA and the encoded protein could be detected in the spleen at 1, 4, and 24 hours after the injection, where both products were at significantly higher levels when pseudouridine-containing mRNA was administered. Even at higher doses, only the unmodified mRNA was immunogenic, inducing high serum levels of interferon- α (IFN- α). These findings indicate that nucleoside modification is an effective approach to enhance stability and translational capacity of mRNA while diminishing its immunogenicity *in vivo*. Improved properties conferred by pseudouridine make such mRNA a promising tool for both gene replacement and vaccination.





Careers in Chemistry

Amy Rhoden Smith, PhD works at Precision BioSciences, a gene editing therapeutics company. As a program development leader, Dr. Rhoden Smith works with multiple teams to lead cell and gene editing therapy programs through preclinical and early clinical development. She loves working in biotechs and startups, where she can think about how to use cutting-edge science to bring meaningful therapies to patients. Dr. Rhoden Smith earned her BS in chemistry at the College of Charleston. She received her PhD in organic chemistry from the University of Texas at Austin.

Dr. Rhoden Smith has always been interested in applying her chemical knowledge to understand biological problems, so she started her career in biotechnology by joining Moderna Therapeutics. Her team used chemistry to create novel therapeutic mRNA conjugates for improved mRNA in vivo half-life and protein expression. She then moved to Intellia Therapeutics, where she led a team focused on generating Cas9 mRNA and synthetic guide RNAs for in vivo gene editing therapeutic applications. After working on projects that were discovery-based, she moved to Precision BioSciences, where she focuses on bringing products into the clinic.

Dr. Rhoden Smith says that organic chemistry gave her an understanding of how molecules interact with each other, which applies not only to small molecules, but also much larger ones like proteins, nucleic acids, and even cells. Just as importantly, it also taught her the value of problem-solving and gave her a passion for understanding complex problems, which she uses every day.



A Novel Amino Lipid Series for mRNA Delivery: Improved Endosomal Escape and Sustained Pharmacology and Safety in Non-human Primates

Staci Sabnis,¹ E. Sathyajith Kumarasinghe,¹ Timothy Salerno,¹ Cosmin Mihai,¹ Tatiana Ketova,¹ Joseph J. Senn,¹ Andy Lynn,¹ Alex Bulychev,¹ Iain McFadyen,¹ Joyce Chan,¹ Örn Almarsson,¹ Matthew G. Stanton,^{1,2} and Kerry E. Benenato¹

¹Moderna Therapeutics, 200 Technology Square, Cambridge, MA 02139, USA

The success of mRNA-based therapies depends on the availability of a safe and efficient delivery vehicle. Lipid nanoparticles have been identified as a viable option. However, there are concerns whether an acceptable tolerability profile for chronic dosing can be achieved. The efficiency and tolerability of lipid nanoparticles has been attributed to the amino lipid. Therefore, we developed a new series of amino lipids that address this concern. Clear structure-activity relationships were developed that resulted in a new amino lipid that affords efficient mRNA delivery in rodent and primate models with optimal pharmacokinetics. A 1-month toxicology evaluation in rat and non-human primate demonstrated no adverse events with the new lipid nanoparticle system. Mechanistic studies demonstrate that the improved efficiency can be attributed to increased endosomal escape. This effort has resulted in the first example of the ability to safely repeat dose mRNA-containing lipid nanoparticles in non-human primate at therapeutically relevant levels.

INTRODUCTION

mRNA-based therapies have the potential to revolutionize the way we treat diseases. The surging interest in mRNA as a drug modality stems from the potential to deliver transmembrane and intracellular proteins, targets that standard biologics are unable to access due to their inability to cross the cell membrane.¹ One major challenge to making mRNA-based therapies a reality is the identification of an optimal delivery vehicle. Due to its large size, chemical instability, and potential immunogenicity, mRNA requires a delivery vehicle that can offer protection from endo- and exo-nucleases, as well as shield the cargo from immune sentinels. Lipid nanoparticles (LNPs) have been identified as a leading option in this regard.² Moderna Therapeutics has recently validated this approach by demonstrating safe and effective delivery of an mRNA-based vaccine formulated in LNPs.³

Key performance criteria for an LNP delivery system are to maximize cellular uptake and enable efficient release of mRNA from the endosome. At the same time, the LNP must provide a stable drug product and be able to be dosed safely at therapeutically relevant levels. LNPs are multi-component systems that typically consist of an amino lipid, phospholipid, cholesterol, and a PEG-lipid.² Each component is required for aspects of efficient delivery of the nucleic acid cargo and stability of the particle. The key component thought to drive cellular uptake, endosomal escape, and tolerability is the amino lipid. Cholesterol and the PEG-lipid contribute to the stability of the drug product both *in vivo* and on the shelf, while the phospholipid provides additional fusogenicity to the LNP, thus helping to drive endosomal escape and rendering the nucleic acid bioavailable in the cytosol of cells.

Several amino lipid series have been developed for oligonucleotide delivery over the past couple of decades.⁴ The literature highlights direct links between the structure of the amino lipid and the resultant delivery efficiency and tolerability of the LNP. The amino lipid MC3 (DLin-MC3-DMA) is the most clinically advanced oligonucleotide delivery system, as siRNA formulated in MC3-based LNPs has progressed to phase III for the treatment of transthyretin-mediated amyloidosis.^{5,6} More recently, literature reports have demonstrated the effectiveness of MC3-based LNPs to deliver mRNA.7 LNPs of this class are quickly opsonized by apolipoprotein E (ApoE) when delivered intravenously (i.v.), which enables cellular uptake into hepatocytes by the low-density lipoprotein receptor (LDLr).⁸ Concerns remain that MC3's long tissue half-life could contribute to unfavorable side effects hindering its use for chronic therapies.⁹ In addition, LNPs can induce activation of the immune system resulting in complement activation-related pseudoallergy (CARPA), an acute immunological response that can lead to anaphylactic-like shock.¹⁰

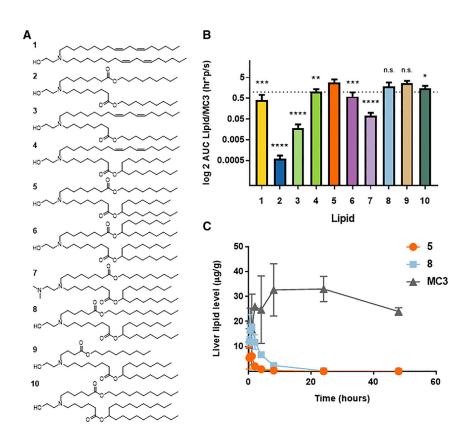
To unleash the potential of mRNA therapies for humans, we required a class of LNPs with increased delivery efficiency along with a metabolic and toxicity profile that would enable chronic dosing in humans.

Correspondence: Kerry E. Benenato, Moderna Therapeutics, 200 Technology Square, Cambridge, MA 02139, USA.

E-mail: kerry.benenato@modernatx.com

Received 20 November 2017; accepted 9 March 2018; https://doi.org/10.1016/j.ymthe.2018.03.010.

²Present address: Generation Bio, 215 First Street, Suite 150, Cambridge, MA 02142, USA



We took a rational medicinal chemistry approach to amino lipid optimization aiming to identify structural motifs that provide chemical stability, optimal tissue clearance, and mRNA delivery efficiency. Our initial rodent screens led to the identification of a lead lipid with good delivery efficiency and pharmacokinetics. The lead LNP was profiled further in non-human primate for efficiency of delivery after single and repeat dosing. Finally, the optimized LNPs were evaluated in 1-month repeat dose toxicity studies in rat and non-human primate.

RESULTS

Initial screening of a broad chemical space identified ethanolamine as an amino lipid head group that could effectively drive mRNA encapsulation and provide LNPs with superior physiochemical properties. Combining the ethanolamine head group with di-linoleic lipid tails (lipid 1; Figure 1; Table 1) generated an LNP with high encapsulation of luciferase mRNA, small particle size, and low polydispersity index (PDI). The LNP with lipid 1 had a surface pK_a (apparent value for the particle) in the range that has been shown to be optimal for siRNA delivery.^{11,12} To evaluate the efficiency of the new amino lipids, LNPs using the novel lipids were tested in vivo in mice using firefly luciferase mRNA as a reporter. An MC3 LNP was included as a control in each experiment, enabling us to compare LNPs from experiment to experiment. Measured luciferase activity also enabled us to determine protein bio-distribution. i.v. delivery of 0.5 mg/kg (mRNA dose level) of lipid 1-based LNPs to mice resulted in luciferase activity two-fold lower than an MC3 LNP control (Figure 1B).

1510 Molecular Therapy Vol. 26 No 6 June 2018

Figure 1. Optimization of Efficiency and Clearance of Amino Lipid

(A) Structures of amino lipids. (B) Whole-body luciferase bioluminescence AUC of novel LNPs versus MC3 LNPs, measured in CD-1 mice (n = 6 at 3 and 6 hr, n = 3 at 24 hr), 0.5 mg/kg dose firefly luciferase (ffLuc) mRNA, i.v. bolus, error bars indicate SD of the ratio of novel lipid AUC versus MC3 AUC. *p < 0.05, **p < 0.01, ***p < 0.001, ***p < 0.001, n.s. = not statistically significant. (C) Parent amino lipid levels measured in liver tissue from Sprague-Dawley rats (n = 3 per time point), 0.2 mg/kg dose hEPO mRNA, mean \pm SD, p < 0.05 for lipids 5 and 8 AUC relative to MC3.

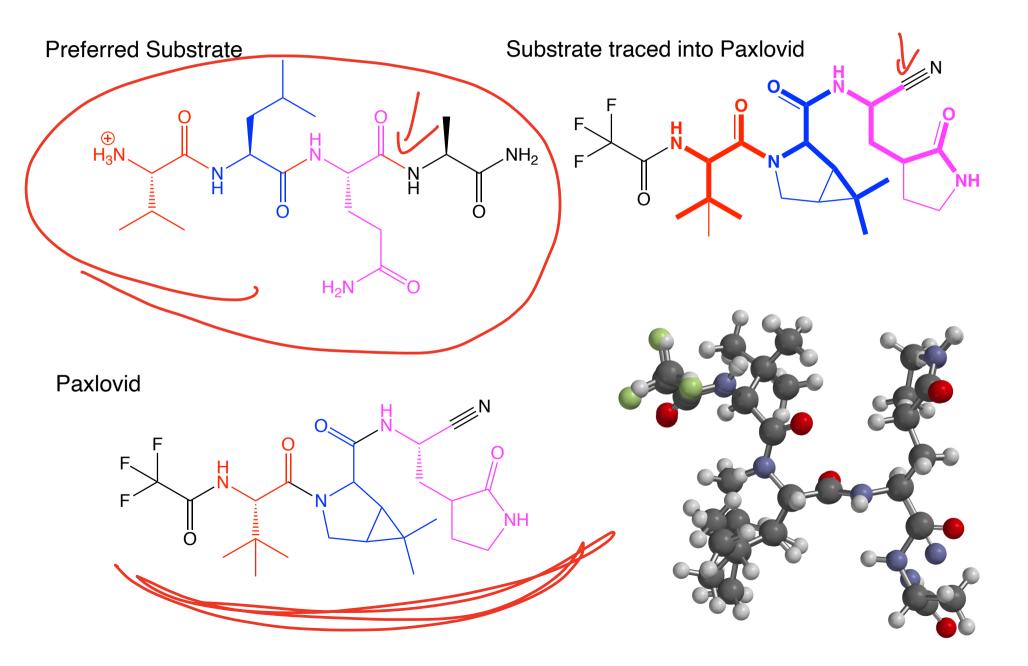
Whole-body imaging clearly demonstrated that the majority of protein expression was localized in the liver (Figure S1). We found that the lipid had similar clearance to MC3 from liver tissue with 66% of the original dose remaining in liver tissue of mice 24 hr post-dose (Table 1).

To improve tissue clearance, we introduced ester linkages in the lipid tails (lipids 2 and 3; Figure 1A), which are reported to trigger metabolism by esterases *in vivo*.¹³ This has been shown to be a viable strategy to improve lipid

clearance in a MC3-based lipid structure.⁹ First, we established that the lipids were chemically stable by measuring ethanol stability at room temperature and 37° C (Table S1). We observed less than 1% change in purity for all lipids tested. LNPs formed with lipid 2 were significantly larger with a surface $pK_a > 7$ (Table 1). Removal of one ester (lipid 3) afforded LNPs with improved physiochemical characteristics and lower LNP surface pK_a . *In vivo*, neither lipid demonstrated efficient mRNA delivery (Figure 1B); however, we did observe rapid tissue clearance, with no lipid detected at 24 hr (Table 1).

Improvement in protein expression was observed when a secondary ester was introduced (lipid 4; Figures 1A and 1B). We observed equivalent expression to MC3 LNPs, but the clearance rate was slower than lipids 2 and 3 (67% lipid remaining; Table 1). Replacement of the linoleic tail with a primary ester-containing lipid tail (lipid 5; Figure 1A) provided increased expression (3-fold higher than MC3; Figure 1B) and optimal tissue clearance (no lipid detected at 24 hr; Table 1). To further increase expression an additional secondary ester was introduced (6), but this resulted in a lowering of the surface pK_a to 6.00 and lower luciferase activity. In addition, the lipid had a significantly slower tissue clearance with 68% remaining at 24 hr.

With an optimal lipid tail structure identified we re-visited the ethanolamine head group. Lipid 7 is one representative example (Figure 1A) where the alcohol functionality is replaced with a dimethylamine. This generated an LNP with comparable physiochemical properties, but complete loss of delivery efficiency (Figure 1B).



<u>Organic Chemistry</u> is the study of carbon-containing molecules. This class has two points.

The first point of the class is to understand the organic chemistry of living systems. We will teach you how to think about and understand the most amazing molecules on the planet!!

You will learn how MRI scans work. 1/18/24

You will learn the basic principles of pharmaceutical science and how many drugs work. $\sqrt{23/24}$

You will learn about the special bond that holds carbohydrates such as glucose in six-membered rings, connects carbohydrate monomers together to make complex carbohydrate structures and is critical to DNA and RNA structure. $\gamma/(/24)$

You will learn how soap is made from animal fat and how it works to keep us clean. 2/27/24

You will learn the important structural reason proteins, the most important molecular machines in our bodies, can support the chemistry of life. 2/20/24 A mide Day!

You will learn how important antibiotics like penicillins work, including ones that make stable covalent bonds as part of their mode of action. 4/4/24

You will learn why carrots are orange and tomatoes are red. 4/4/24

You will learn the very cool reason that the DNA and RNA bases are entirely flat so they can stack in the double helix structure. 4/23/24

You will learn even more about why fentanyl is such a devastating part of the opioid problem and how Naloxone is an antidote for a fentanyl overdose. $\frac{1}{23}$

You will learn even more details about why Magic Johnson is still alive, decades after contracting HIV, and how the same strategy is being used to fight COVID. $\frac{4}{23}{24} \rightarrow \frac{4}{25}{24}$

You will learn about the surprising chemical reason the Pfizer and Moderna mRNA vaccines elicit strong immune responses. 4/25/24

The second point of organic chemistry is the synthesis of complex molecules from simpler ones by making and breaking specific bonds, especially carbon-carbon bonds.

You will learn how carbon-metal bonds lead to new carbon-carbon bonds. 1/18/24

You will learn how most reactions of carbonyl compounds involve only the four common mechanistic elements operating in only a few common patterns. $\frac{1}{23}$

You will learn how, by simply adding a catalytic amount of base like HO⁻ to aldehydes or ketones, you can make new carbon-carbon bonds, giving complicated and useful products. $\frac{3}{7}$

You will learn a reaction that can convert vinegar and vodka into a common solvent. 2/13/24 Esterificities

You will learn why molecules with six-membered rings and alternating double bonds are stable. $\frac{\gamma}{9}$

You will learn a reaction that can turn model airplane glue into a powerful explosive. $\frac{1}{23}$

Most important, you will develop powerful critical thinking skills:

- 1. You will learn how to look at a molecule and accurately predict which atoms will react to make new bonds, and which bonds will break during reactions.
- 2. You will learn how to analyze a complex molecule's structure so that you can predict ways to make it via multiple reactions starting with less complex starting molecules.

Oh Lord, Make Me A Doctor With A Mercedes-Benz

Oh Lord, make me a doctor with a Mercedes-Benz My friends are all prelaw, I must make amends. Worked hard in organic, never thought it would end. So Lord, make me a doctor, with a Mercedes-Benz.

Oh Lord, won't you give me an "A" on the test, I have to show med-schools that I am the best. Tried hard all semester, the TA says I'm a pest, So Lord, won't you give me an "A" on the test.

Oh Lord, won't you give me a high MCAT score. My friends all got 520's, I have to get more. I went to Kaplan, oh man what a bore, So Lord, won't you give me a high MCAT score.

Oh Lord, won't you get me into Johns Hopkins U. Baylor would be nice, I'd like an interview O.K. let's be serious, A&M will do. So Lord, won't you get me into TAMU.

Oh Lord, make me a doctor with a Mercedes-Benz My friends are all prelaw, I must make amends. Worked hard in organic, never thought it would end. So Lord, make me a doctor, with a Mercedes-Benz.



Ring of Pi

Benzene is a funny thing 'Cause it is a special ring Hückel's rules, they do apply For a ring, a ring of pi

Electrons in a big old ring of pi They go round, round, round. And stability is high. As we learn, learn, learn, The ring of pi-i The ring of pi.

The structure, it is sweet When 4N and then 2 meet Sp2, and flat required Then the 2p's, they are wired

Electrons in a big old ring of pi They go round, round, round. And stability is high, As we learn, learn, learn, The ring of pi-i The ring of pi.

Electrons in a big old ring of pi They go round, round, round. And stability is high, As we learn, learn, learn, The ring of pi-i The ring of pi.

"Make it B"

On the final When I'm having trouble Inspiration comes to me Speaking words of wisdom Make it B

A tetrahedral intermediate And a leaving group you see Here are words of wisdom Make it B,

Make it B, Make it B, Make it B, Make it B Mechanism B Make it B

The Claisen is a great reaction Chemists of the world agree For Claisen its the answer Make it B

And when my mind is cloudy There is still a light that shines on me With a leaving group, Make it B

Make it B, Make it B, Make it B, Make it B With a leaving group Make it B

"Organic Labville"

There must be a mistake, What did I just make Those aren't crystals, it looks more like oil. Looks like another screw up, Like the last one that blew up, Uh oh now its beginning to boil.

Wasting my time again in my organic lab. Looking for that last product I lost. Some people say that my lab partners to blame, But I know, its my T.A.'s fault.

Don't know the reason, I always get C's in My lab notebook it looks good to me. I copy it from Suzy, She's a real cutie, Why she doesn't like me I haven't a clue.

Wasting my time again in my organic lab. Looking for that last product I lost. Some people say that my bad technique is to blame, But I know, its my T.A.'s fault.

There's another fire in Welch, AFD just can't squelch. They said it wasn't safe so they sent us all home. But there's chemicals in my blender And I hope it will render, That crystalline product I need to be done.

Wasting my time again in my organic lab. Looking for that last product I lost. Some people say that the professor is to blame, But I know, its my T.A.'s fault. Yes and Some people say that the professor is to blame, But I know, its my T.A.'s fault.

We All Love Organic Chemistry

In the town where I was born, Lived a man of chemistry. And he told us of his life In the organic laboratory.

Making molecules to fight disease Coming up with their syntheses. So we sit in 320N Learning organic chemistry.

Refrain: We all love organic chemistry Synthetic chemistry Molecules with "C" We all love organic chemistry Synthetic chemistry Molecules with "C"

All our friends think we're a bore Our grade point averages begin with 4. But we await graduation day To work in lab for meager pay.

But its OK, who else can say They're curing cancer or fighting AIDS. We hope that you in 320N Respect organic chemistry

Refrain: We all love organic chemistry Synthetic chemistry Molecules with "C" We all love organic chemistry Synthetic chemistry Molecules with "C"