



Delivering on the promise  
of Prime Editing

**Advances in Prime Editing enable *in vivo* therapeutic correction of the *ATP7B* p.H1069Q and p.R778L mutations causing Wilson's Disease**

**Jeremy S. Duffield MD PhD FRCP**  
Chief Scientific Officer, Prime Medicine

American Association for the Study of Liver Diseases  
November 18<sup>th</sup>, 2024

On behalf of the team at Prime Medicine

## Disclosures

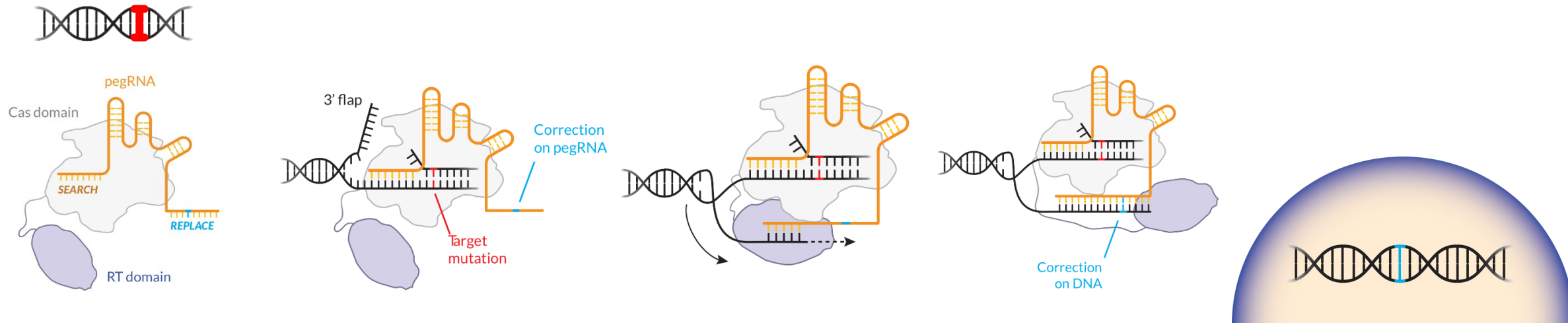
---

Jeremy Duffield declares he is currently an employee of Prime Medicine, Inc. and owns equity in Prime Medicine.

# Prime Editing is programmable for both search and replace

The PE technology utilizes a Prime Editor protein and a Prime Editing guide RNA (pegRNA) to directly write new genetic information into a targeted DNA site without requiring a DSB

Gene with mutation



## SEARCH

Prime editor complex initiates search for target DNA

## FIND & NICK

Prime editor complex finds DNA with target mutation, nicks one strand

## PRIME

Nicked DNA strand primes the RT domain for DNA synthesis

## REPLACE

Prime editor complex copies in corrective DNA sequence

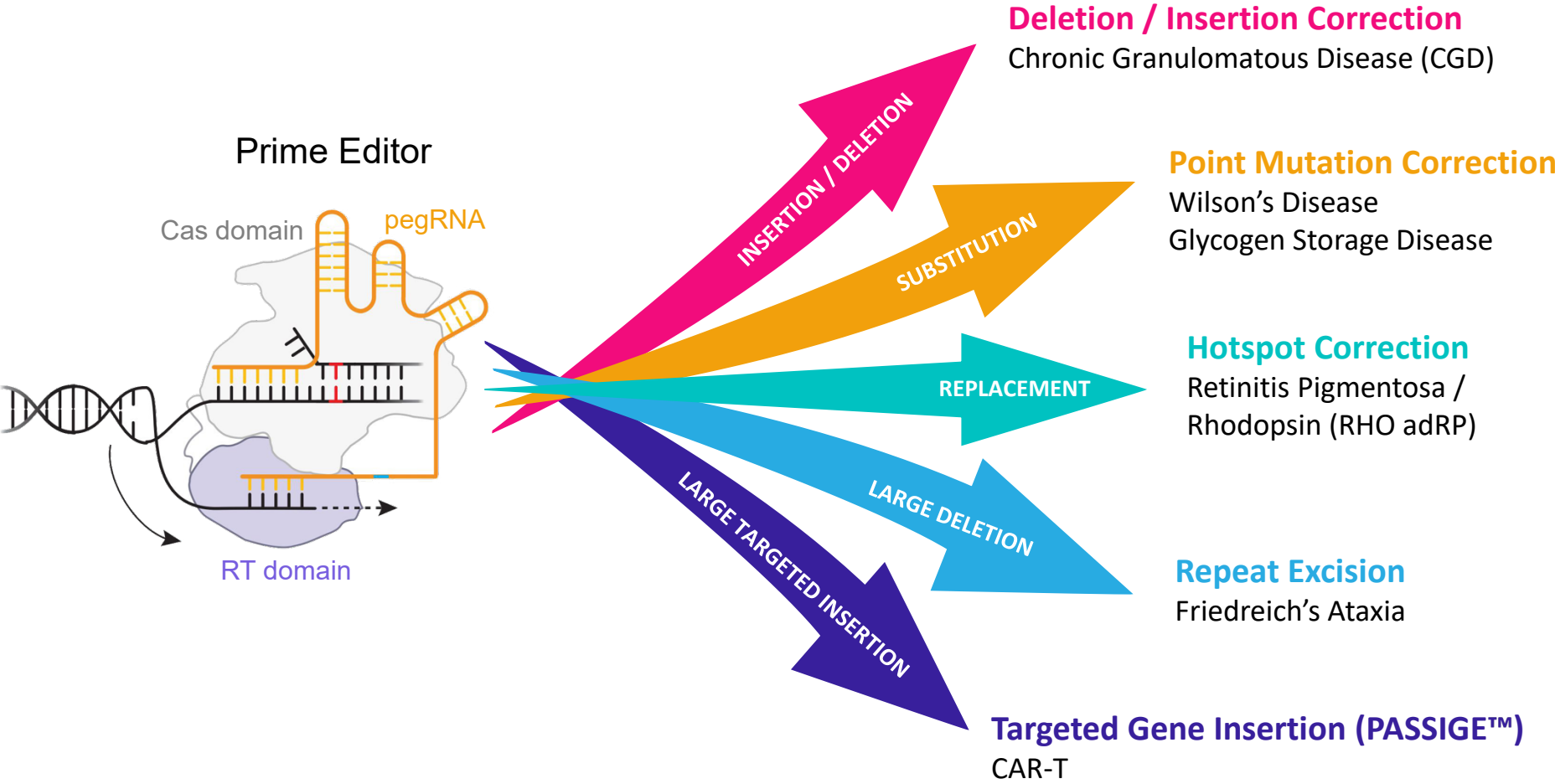
## GENE CORRECTED

3' flap preferentially incorporated<sup>1</sup>, excess flap repaired, gene fully corrected

<sup>1</sup> Completion of an edit requires 3 "edit checks"; pegRNA = Prime Editing guide RNA; RT = reverse transcriptase; Cas = CRISPR associated protein; DSB = Double-stranded break

# We believe Prime Editing is the only gene editing technology that can edit, correct, insert and delete DNA sequences in any target tissue

Corrects mutations across many tissues, organs and cell types, in dividing and non-dividing human cells

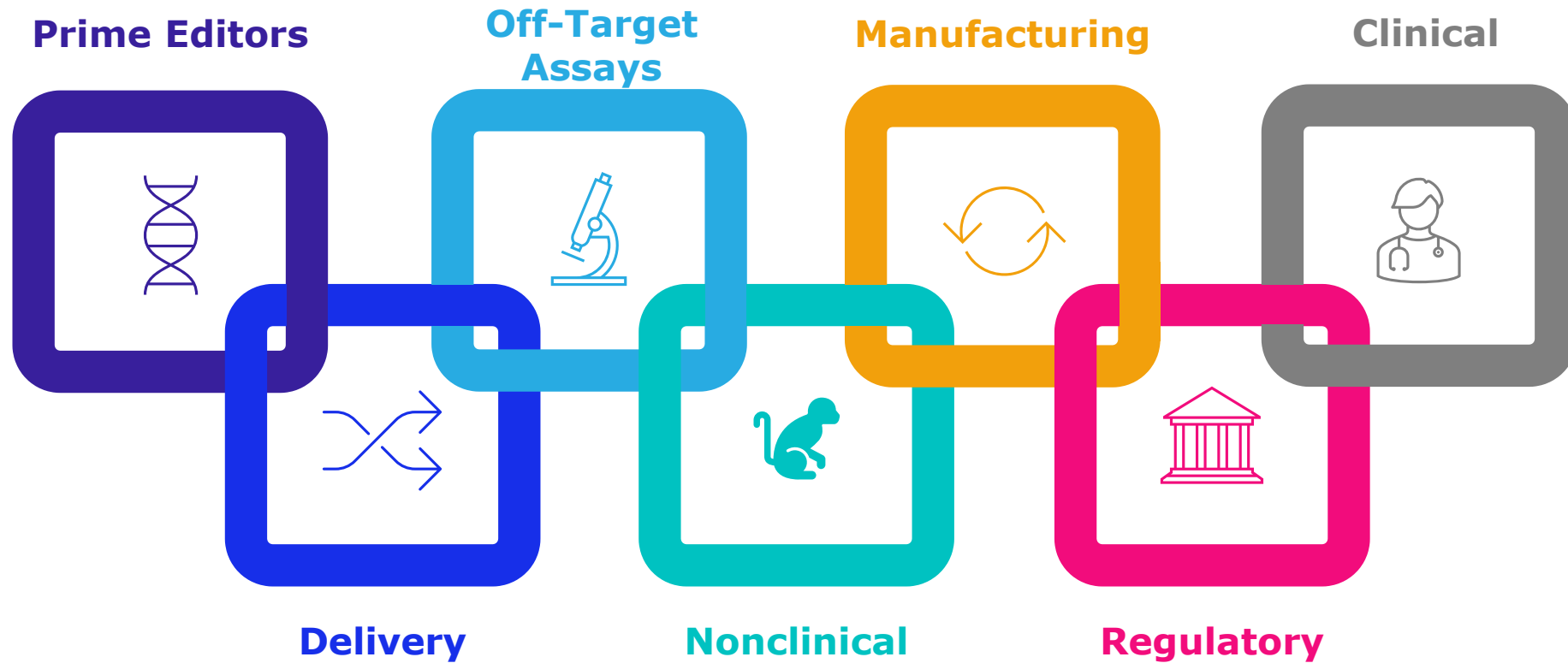


Broad and versatile editing capabilities unlock opportunities across **thousands of indications**, including genetic diseases, infectious diseases, cancers and immunological diseases

CAR-T = chimeric antigen receptor (CAR)-T cell therapy

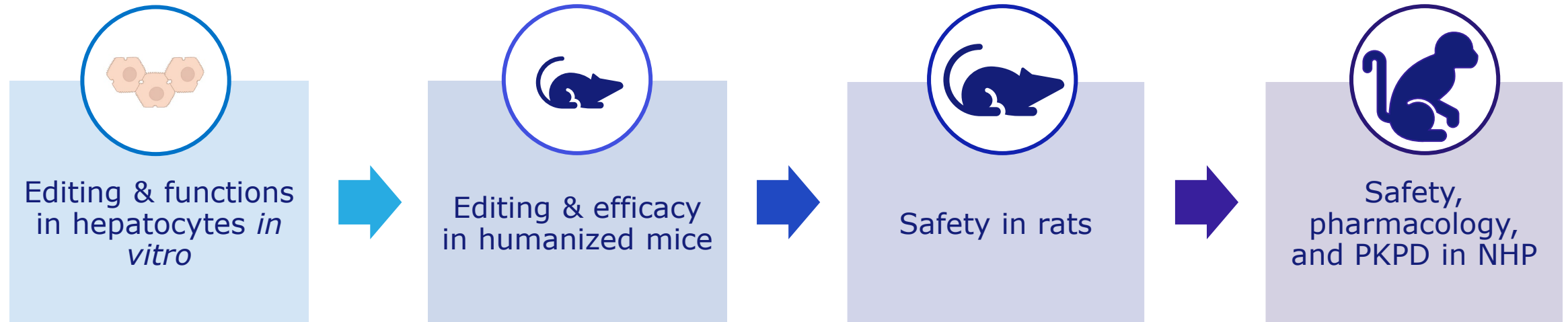
Prime Editing platform modularity accelerates and de-risks ongoing efforts, enabling rapid generation of new product candidates

Core components can be readily leveraged to accelerate pipeline growth, efficiency and execution



# Prime Medicine's approach to developing Prime Editors to treat liver and metabolic diseases

Prime Editors are specific to *human* patient DNA sequence and designed for the correction of *human* mutations

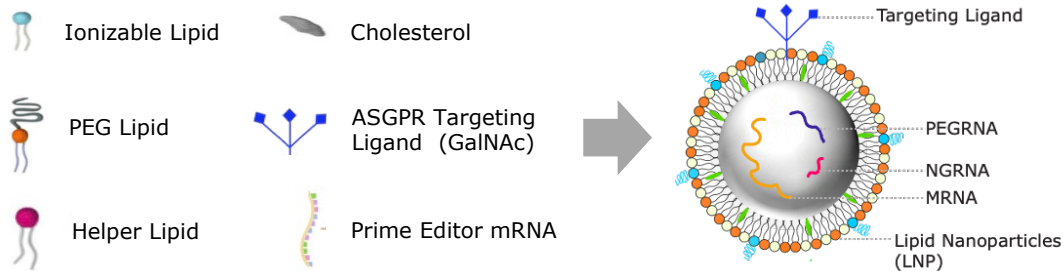


- Establish potency of lead Prime Editor drug candidates
- Establish genotype to phenotype correlation and off-target profile
- Establish pharmacology, safety, tolerability
- Determine biodistribution, drug pharmacokinetics
- Determine PK/PD relationships, human dose projections

# Prime Medicine has developed a universal LNP for our liver and metabolic programs

## Prime Medicine's Universal LNP contains a novel GaINAc targeting ligand

### Shared LNP/PE components

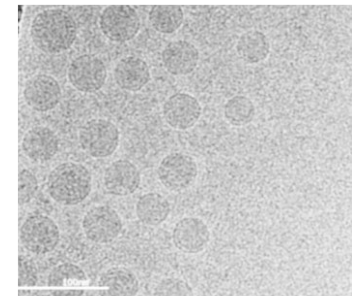
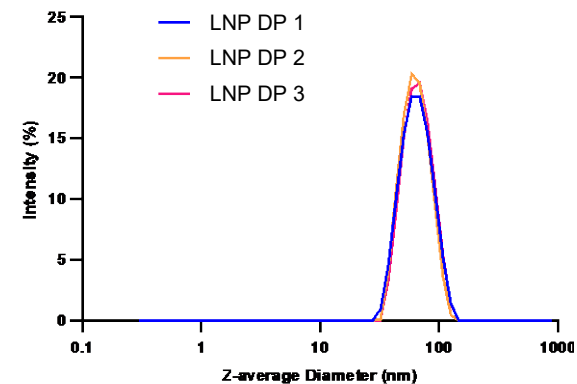


Compared to LNPs without a targeting ligand, Prime's Universal LNP\*:

- ✓ Increases potency
- ✓ Improves safety profile
- ✓ Improves biodistribution

## Multiple different drug product candidates (DP) from one LNP composition

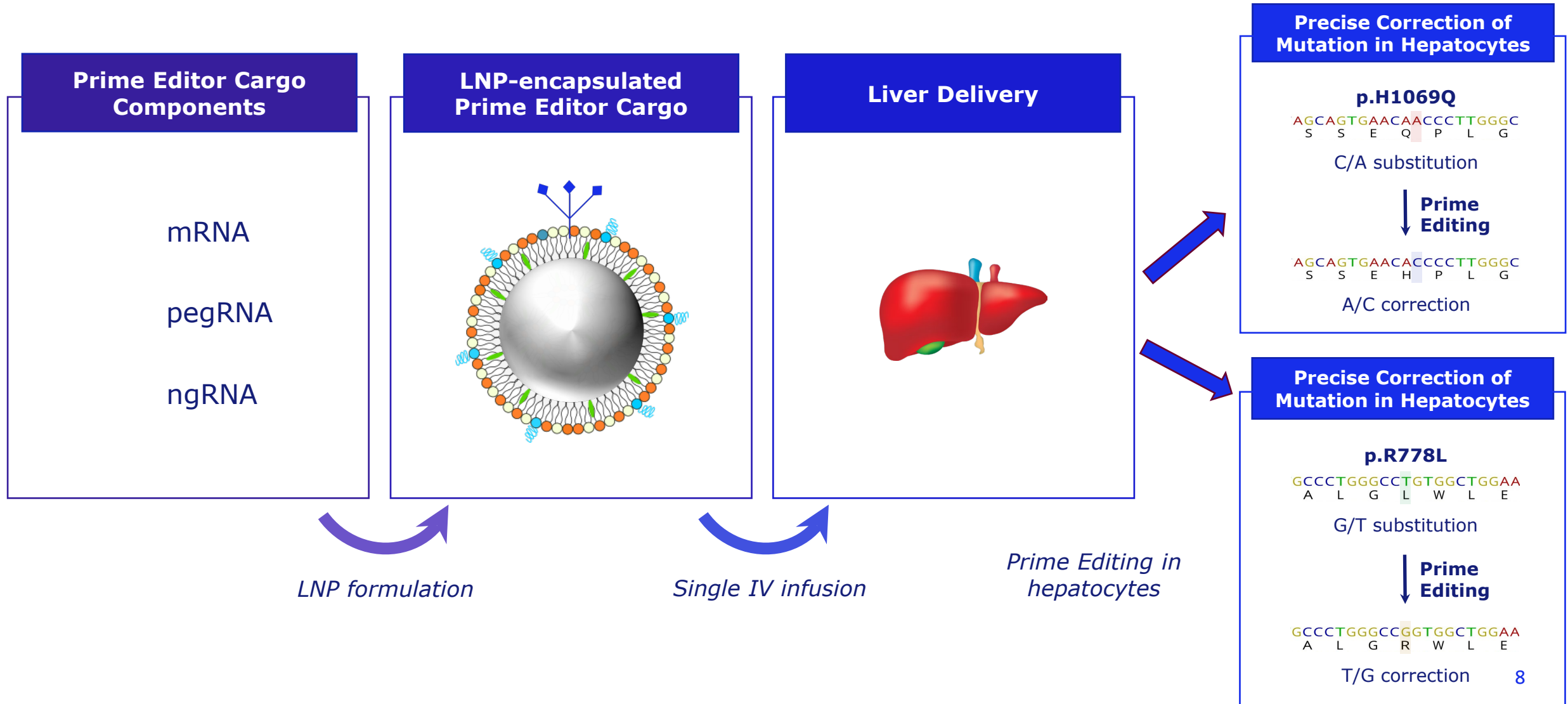
LNP DP	Avg. size (nm)	PDI	% EE
1	60	0.056	99.3
2	59	0.045	97.9
3	60	0.052	98.8



**By swapping only the guide RNAs while keeping the other components constant, we have a new drug product with the potential for the same critical quality attributes**

# Therapeutic approach: LNP-mediated delivery of Prime Editor components to liver

One-time delivery of LNP Prime Editor cargo with potential to correct pathogenic mutations in the liver





# LNP-Formulated Prime Editors to correct common pathogenic mutations causing Wilson's Disease (WD)

## Wilson's Disease

### Disease severity

- Common liver and systemic disease presenting in teens to 20's (prevalence approx. 1:30,000)
- Leads to liver failure, neurocognitive decline and premature death

### Unmet need

- Many patients die without liver transplant. No approved disease-modifying therapies

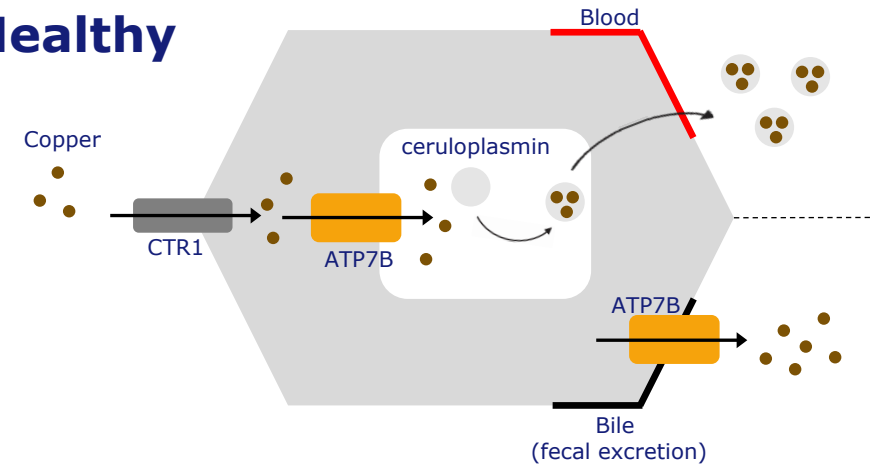
### Human biology

- Autosomal recessive due to loss of function mutations in *ATP7B*
- Affects copper homeostasis, leading to toxic accumulation of copper in liver and brain
- H1069Q and R778L are two prevalent mutations found in up to 50% of patients
- Correction of 20-30% of hepatocytes may be curative

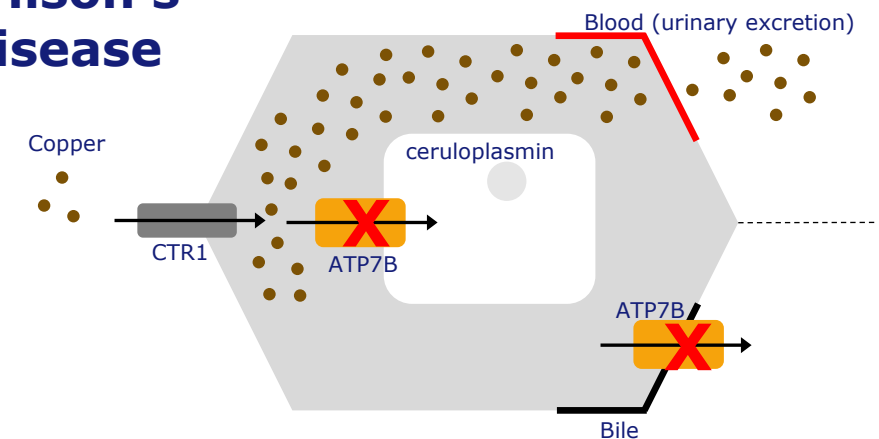
### Prime Medicine's therapeutic approach

- Prime's universal liver-targeted LNP to deliver RNA Prime Editors to patient liver to correct mutations in *ATP7B* to restore copper metabolism

## Healthy



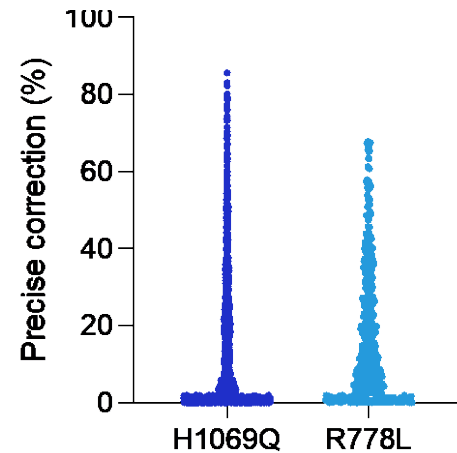
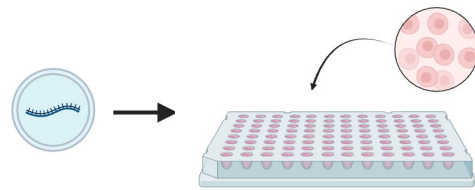
## Wilson's Disease



# Identification of lead Prime Editors for correction of *ATP7B* H1069Q and R778L

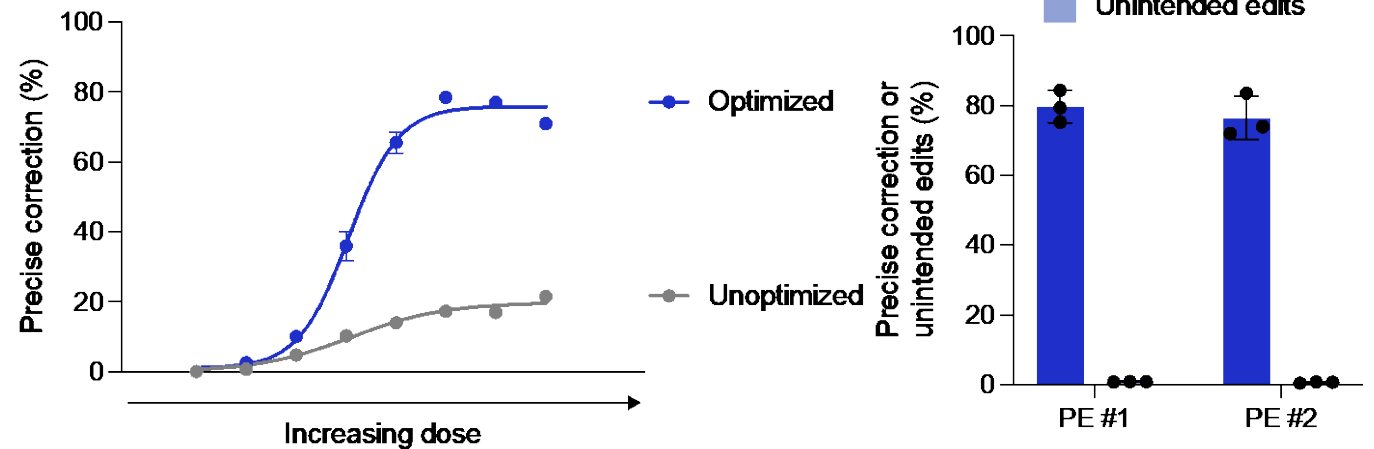
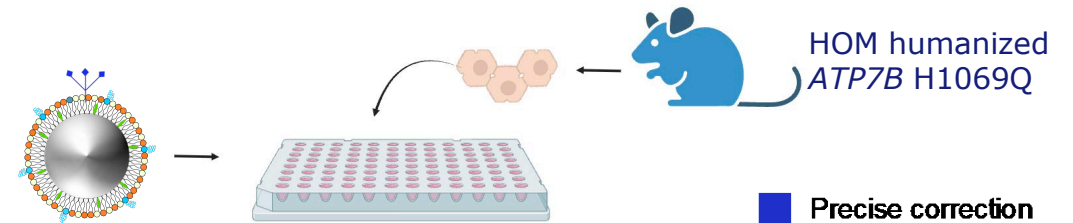
## High throughput screening in hepatocyte cell line

Delivery of RNA Prime Editing components



## Validation in primary humanized mouse hepatocytes

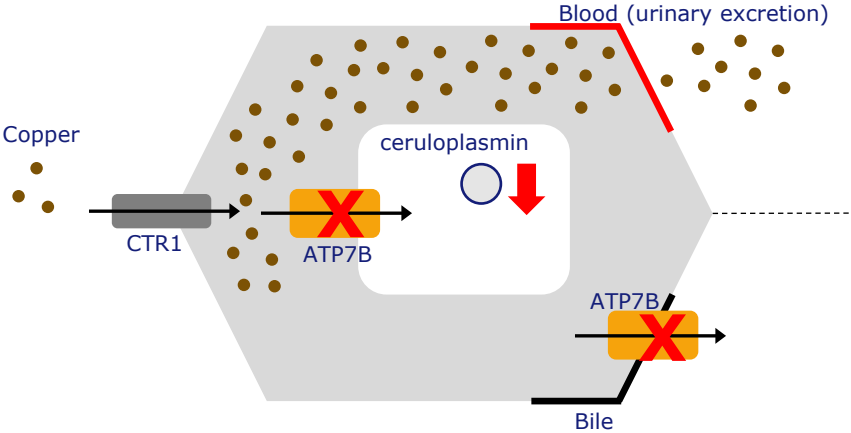
LNP delivery of RNA Prime Editing components



Prime Medicine's high throughput screening platform identifies multiple Prime Editors capable of efficient correction of H1069Q and R778L in hepatocyte cell lines and primary mouse hepatocytes isolated from WD humanized mice

# Prime editing restores ceruloplasmin abundance in patient-derived H1069Q iHeps *in vitro*

## Reduction in ceruloplasmin in WD hepatocytes



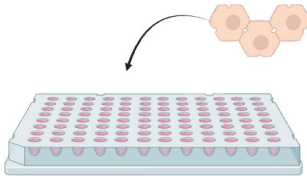
Failure to transport copper in WD reduces ceruloplasmin protein abundance

## Restoration of ceruloplasmin levels *in vitro*

RNA Prime Editing components



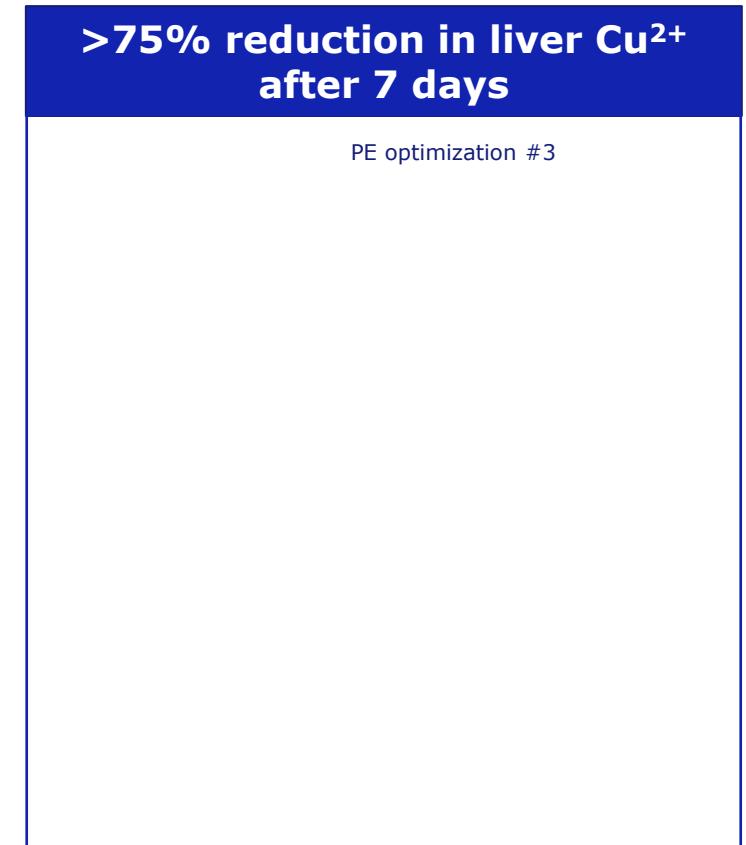
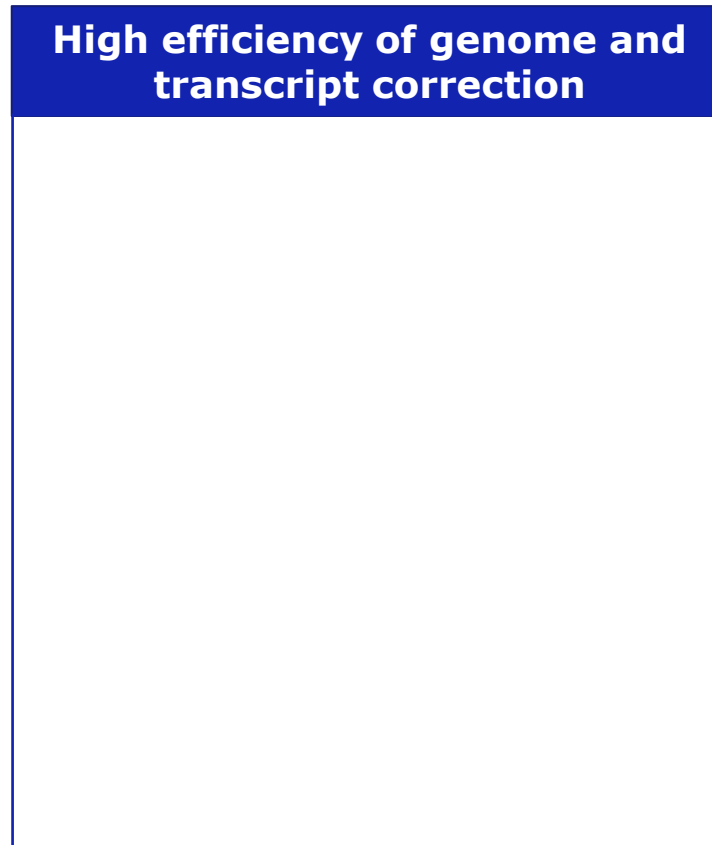
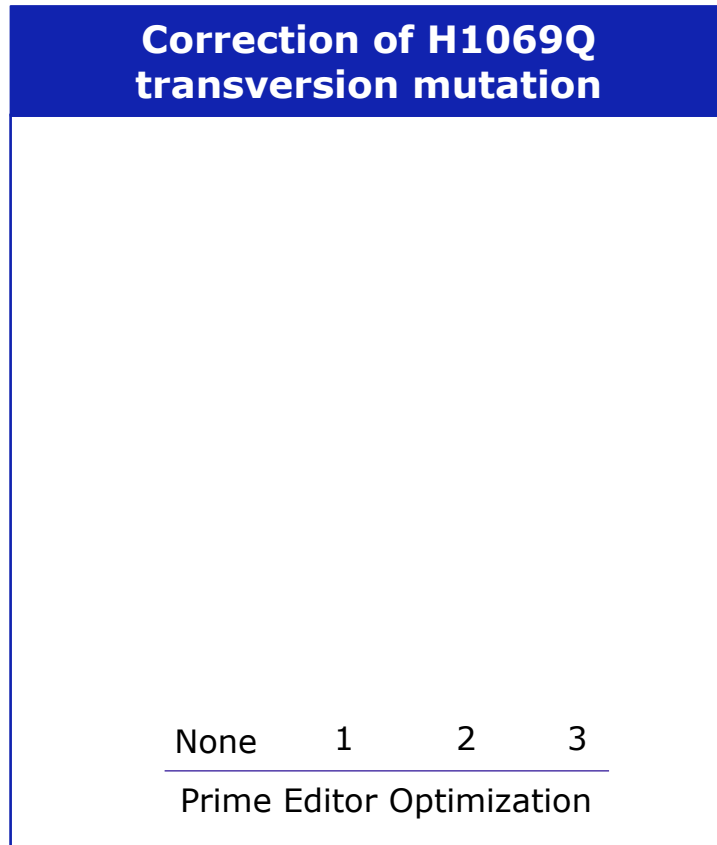
Patient-derived H1069Q iHeps



**Prime Editing in WD patient-derived induced hepatocytes restores ceruloplasmin abundance to levels similar to WT**

# Prime Editors demonstrate efficient DNA and mRNA correction and reduce liver copper in humanized Wilson's Disease mouse model

Fully humanized homozygous p.H1069Q *ATP7B* mouse model



Long term studies ongoing with Prime Editor Optimization #3

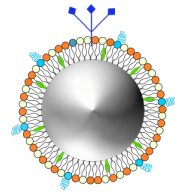
**PE optimizations enable efficient genome and transcript correction as well as liver copper reduction in humanized mice**

\*Based on PK/PD relationships and quantification of cell types in liver: Wang et al Sci. Rep. (2021) 11:19396; MacParland et al Nat Commun. (2018) 9:4383; Hansel et al, Curr Protoc Toxicol (2014) 62:14.12.1; Kmiec, Adv Anat Embryol Cell Biol. (2001) 161:III-XIII. 1-151. Calculation based on 60% of cells in whole liver are hepatocytes; PE = Prime Editor; RNA = ribonucleic acid; mRNA = messenger RNA

# Proof of concept for Wilson's Disease H1069 surrogate Prime Editor in NHP using Prime Medicine's Universal LNP

Initial *in vivo* WD NHP studies show up to 51% *ATP7B* p.H1069 precise hepatocyte editing (interim data)

IV injection of PE  
RNA encapsulated in  
Universal LNP



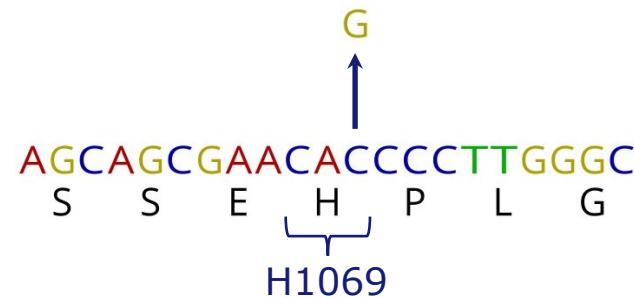
mRNA, pegRNA & ngRNA  
Formulated in Prime  
Universal liver LNP



*Cynomolgus  
macaque*

*ATP7B*\* *in vivo* Prime Editing in NHP  
Precise Editing at H1069 with no  
detectable unintended edits

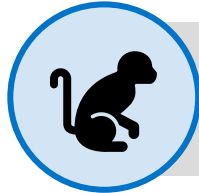
Prime Editing of  
*ATP7B* H1069 in NHP  
liver hepatocytes



- Excellent *in vitro* to *in vivo* translation
- Further surrogate Prime Editor optimizations ongoing

LOD = 0.25%

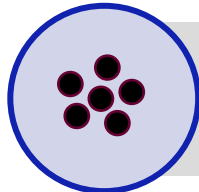
# Prime Medicine's Universal LNP exhibits an excellent safety profile in cynomolgus monkey (NHP)



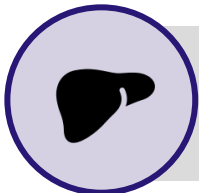
- Well-tolerated with no acute reactions, clinical observations, or body weight changes
- Animals healthy at 54 weeks



- No observed change in platelets, coagulation time or blood count
- No observed change in blood biochemistry panel



- Minimal changes in serum IL-6 levels
- No other observed cytokine changes

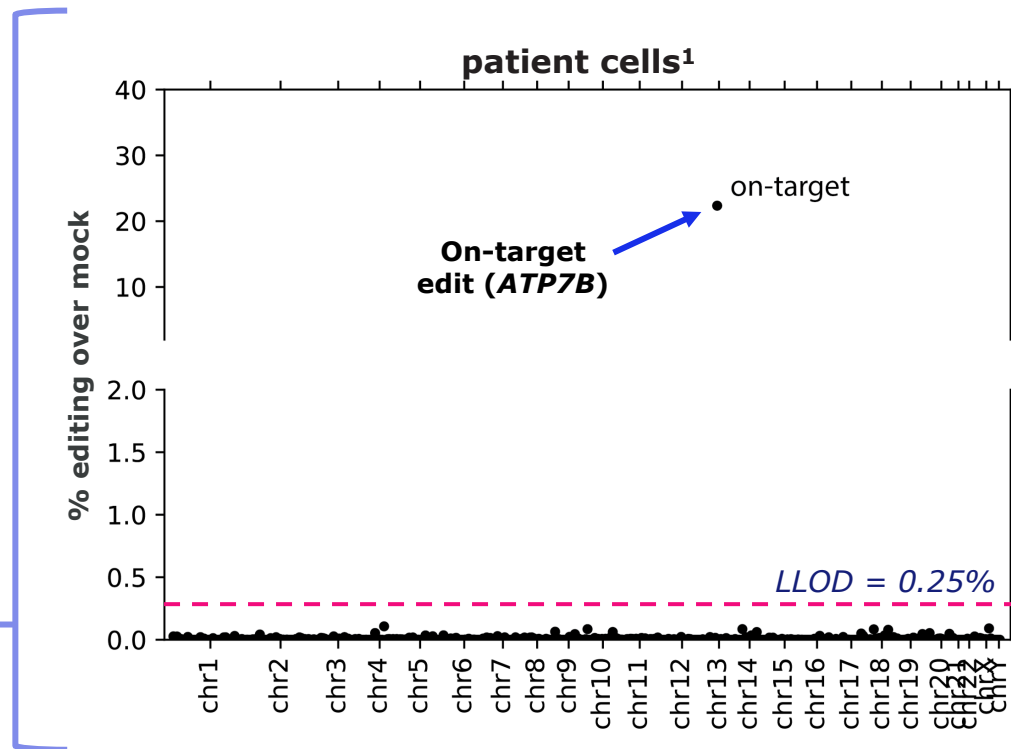
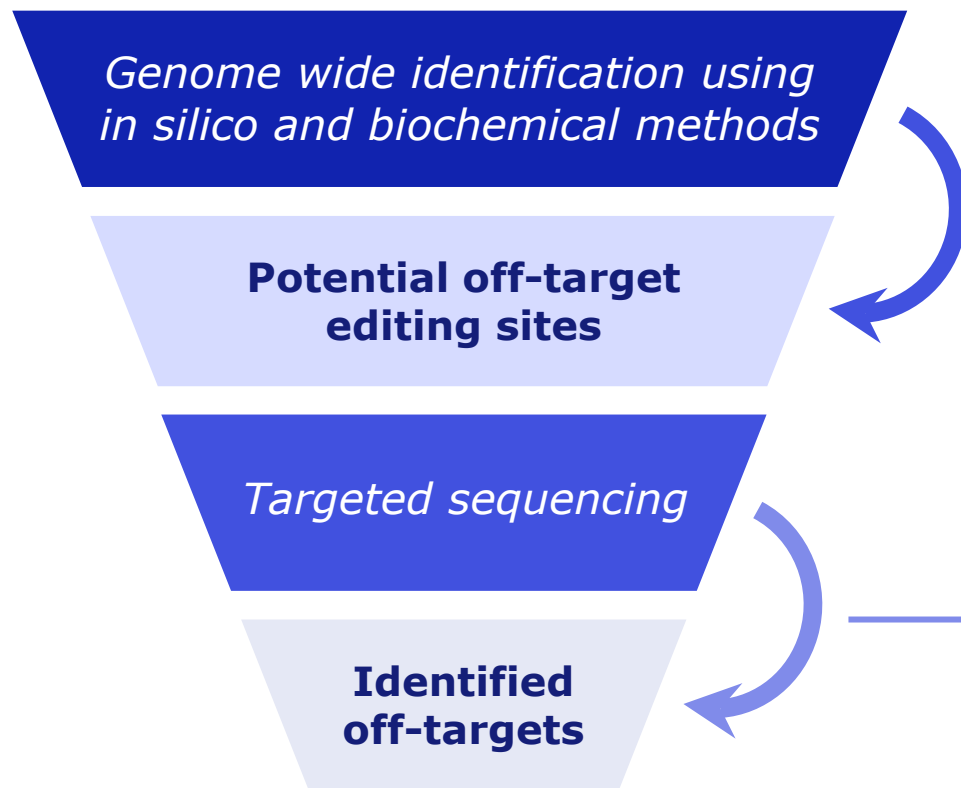


- No changes observed in liver histopathology (H&E)
- Minimal transient LFT elevations

*Benchmarked against other LNPs in clinical development*

# Preliminary analysis: no detectable off-target editing in patient cells treated with Wilson's Disease

"IND ready" set of comprehensive off-target assays



<sup>1</sup>Targeted Analysis of potential off-target sites using targeted deep sequencing in Prime Edited human patient iPSC cells. ATP7B = ATPase copper transporting beta; LLOD = lower limit of detection

## Summary

# Advances in Prime Editing enable *in vivo* therapeutic correction of the *ATP7B* p.H1069Q and p.R778L mutations causing Wilson's disease

### Modular LNP platform

- Prime Medicine has developed a universal liver targeted LNP-PE platform with the potential to deliver Prime Editors to precisely correct disease-causing mutations
- GalNAc targeting ligand improves dose potency, editing, and biodistribution compared to LNPs without targeting ligand
- Excellent and differentiated safety profile in large animal and rodent studies

### Wilson's Disease

- Prime Medicine's Universal LNP-formulated Prime Editors for Wilson's Disease precisely correct the p.H1069Q mutation, with up to 80% precise correction *in vivo*, restore wild-type mRNA expression, and reduce hepatic copper levels in p.H1069Q Wilson's Disease humanized mice at clinically relevant doses
- Results from the initial NHP study demonstrated up to 51% precise hepatocyte editing of *ATP7B* at p.H1069 using an unoptimized surrogate NHP Prime Editor at a dose that was safe and well tolerated

### Off-target editing

- No off-target editing was detected in human cells derived from Wilson's Disease patients





Delivering on the promise  
of Prime Editing

Thank you!



[primemedicine.com](https://primemedicine.com)