Prime Editing advancements enable in vivo therapeutic correction of ATP7B p.H1069Q and p.R778L mutations in Wilson's disease

Healthy hepatocytes

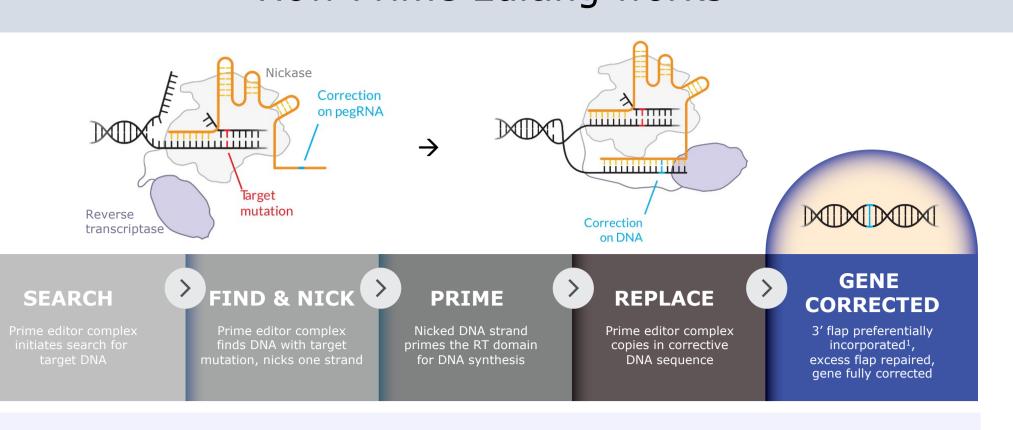
Wilson's disease



David P. Waterman, Shivangi Modi, Alicia Y. Volmar, Michelle S. O'Connor, Celia Chang, Justin Darcy, Chaitali Dutta, Weiyi Li, Marine Hatit, Rowshon Alam, Mallik Putta, Serge Kyrychenko, Jacob Stewart-Ornstein, Seth Alexander, Jonathan Winnay, Andrea De Erkenez, Jon Levy, Andrew V. Anzalone, Vivian W. Choi, John R. Hadcock, Jeremy S. Duffield Prime Medicine, Inc. 60 First Street, Cambridge, MA, USA

Background

How Prime Editing works



Prime Editing for safe and effective treatment of >90% of genetic diseases

Targeted and precise search and replace genome editing enables highly effective disease-causing mutation correction while minimizing unintended edits

Wilson's Disease Overview

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

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

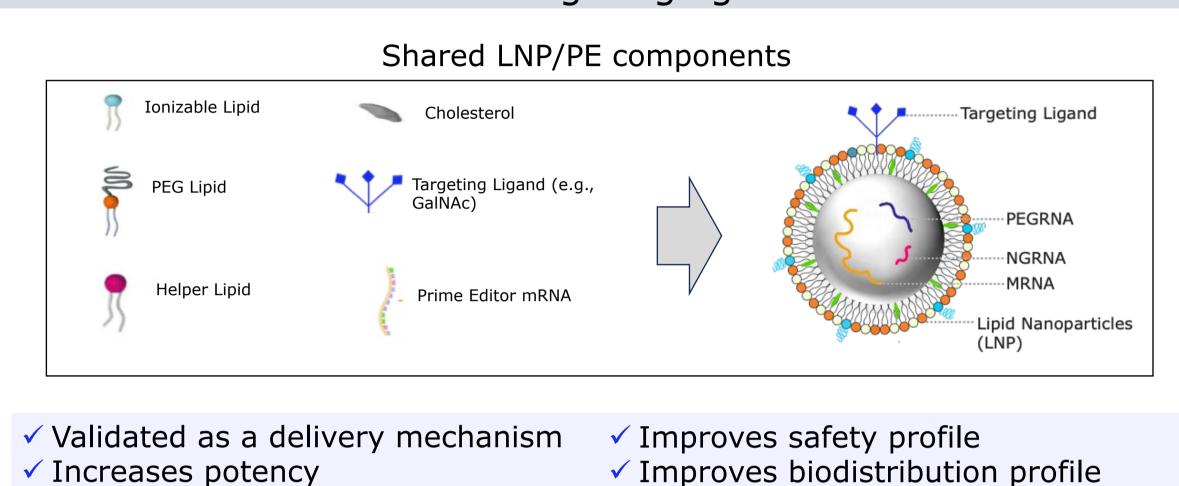
Prime Medicine therapeutic approach

Prime's universal liver targeted LNP, to deliver RNA Prime Editors to patient liver to correct mutations in ATP7B

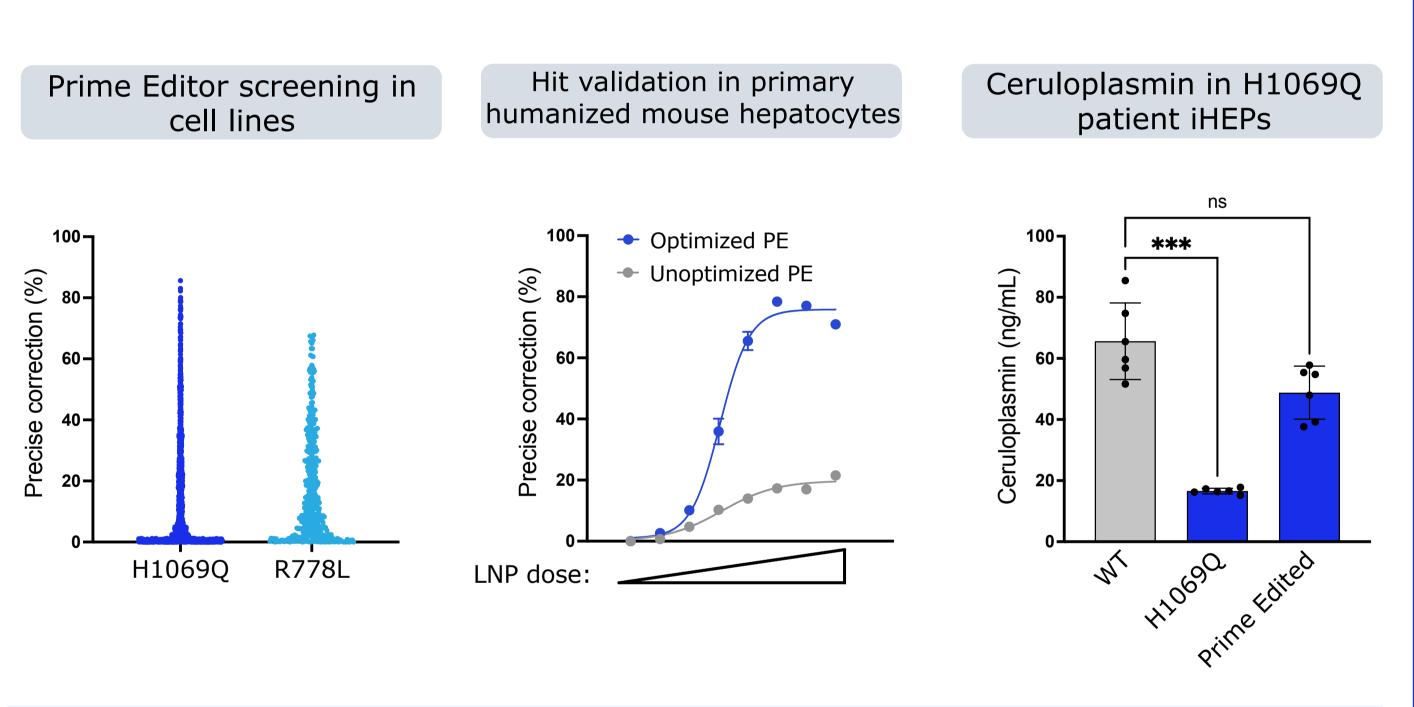
Results

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Prime Medicine's Universal LNP contains a novel GalNAc liver targeting ligand

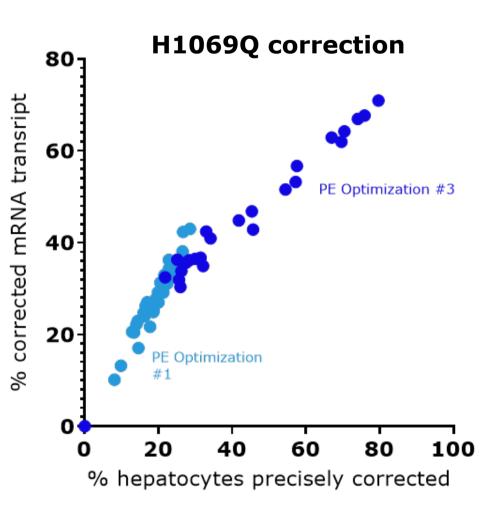


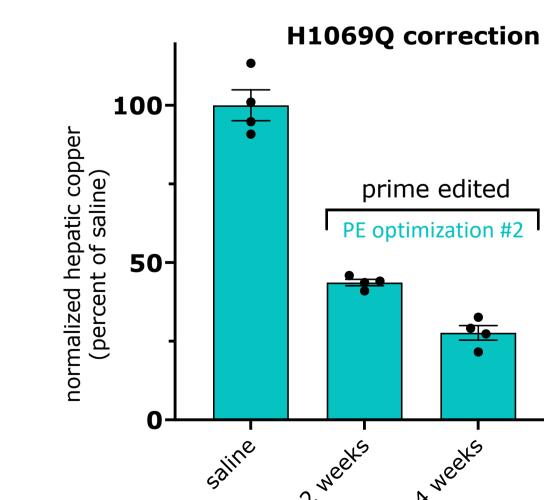
Screening and validation identifies highly active Prime Editors for H1069Q and R778L correction



- High throughput screening in hepatocyte cell lines identifies several candidate Prime Editor components capable of >60% editing for H1069Q and R778L
- Further optimization of PE in ATP7B H1069Q humanized mouse hepatocytes greatly improves Prime Editor potency and efficacy
- Prime Editor normalizes ceruloplasmin abundance in H1069Q patient induced human hepatocytes (iHeps)

Correction of H1069Q in humanized mouse models produces corrected mRNA transcripts and reduces liver copper





- DNA correction of H1069Q by Prime Editing correlates with corrected mRNA
- Time-dependent 75% reduction in liver copper using optimization 2 Prime Editor
- Further copper studies ongoing with Prime Editor Optimization #3

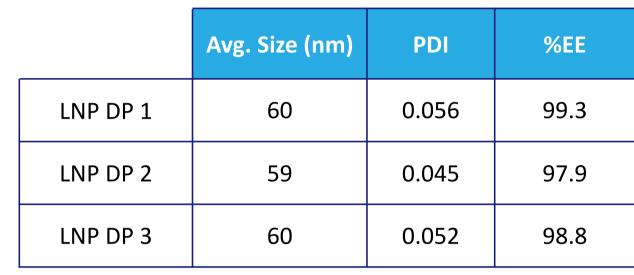
Universal LNP-formulated surrogate H1069 Prime Editor in vivo in NHP achieves up to 51% precise editing in liver hepatocytes

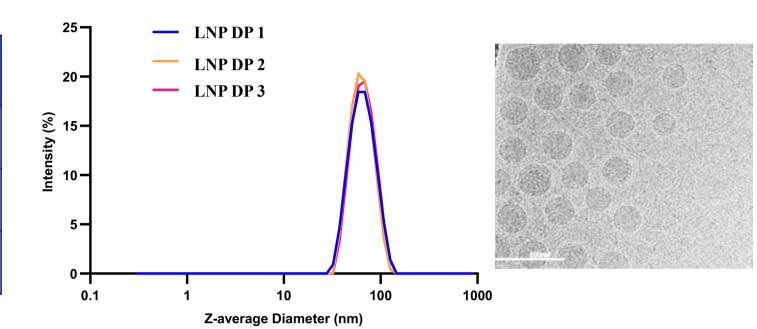
Prime's Universal LNP has an excellent safety profile in NHP

- Well-tolerated with no acute reactions, clinical observations,
- or body weight changes Minimal LFT abnormalities
- No change in platelet, coagulation, or blood count
- No change in blood biochemistry
- Minimal change in IL6 production No other cytokine changes
- No change in liver histopathology (H&E)

ATP7B* H1069 editing in NHP GCGGAGGCCAGCGAACACCCCTTGGGCGTGGCAGTC 60₇ H1069 წ **50**-∙50 % 40-40 No detectable **30** unintended edits 20 **20**· 10-10 *unoptimized pegRNA

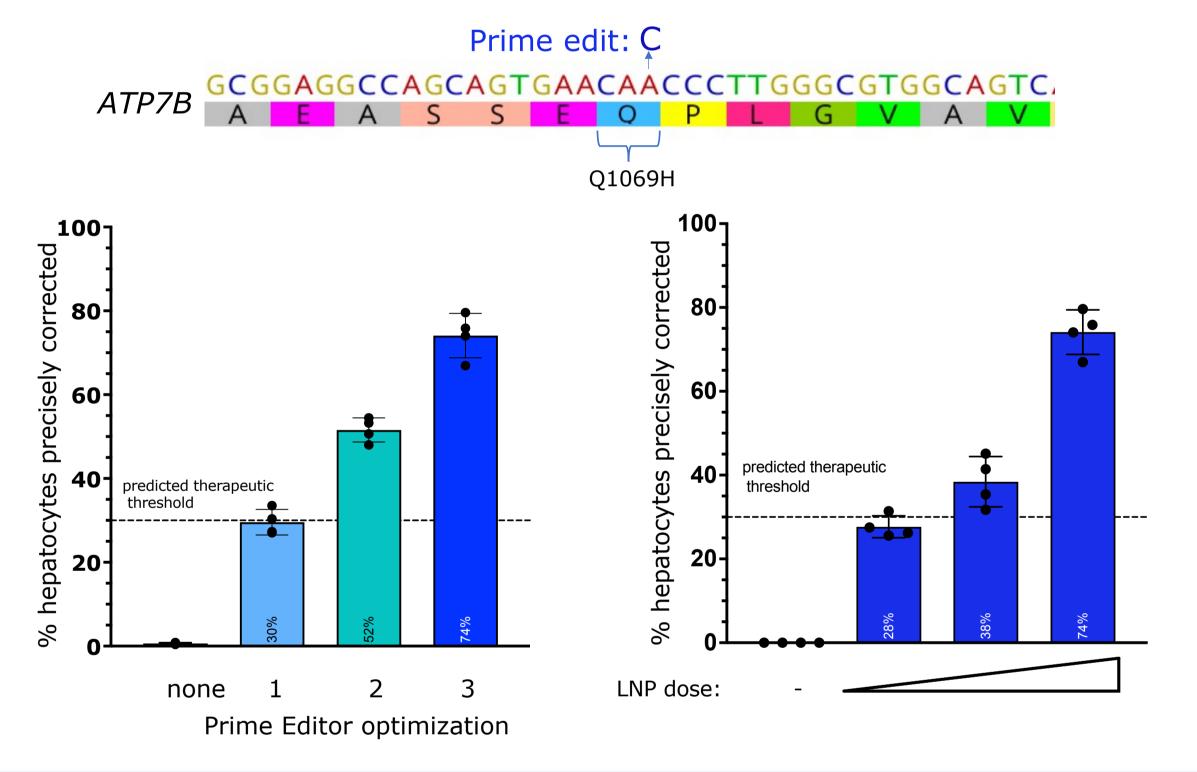
Prime Medicine's Universal LNP can be used to generate multiple drug products





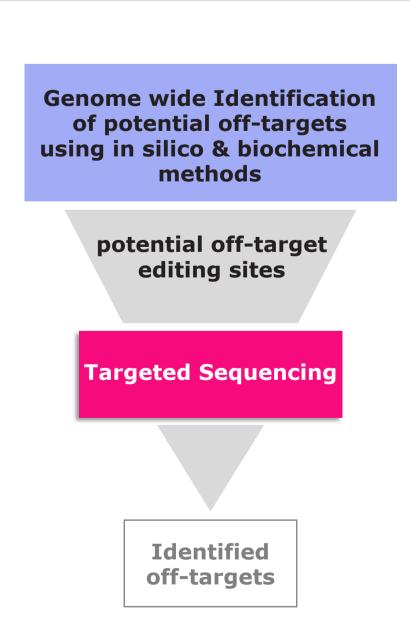
- Prime Medicine's Universal LNP keeps constant 6 out of the 8 components of the LNP and PE cargo for all liver programs
- By swapping out only the guide RNAs new drug products are made

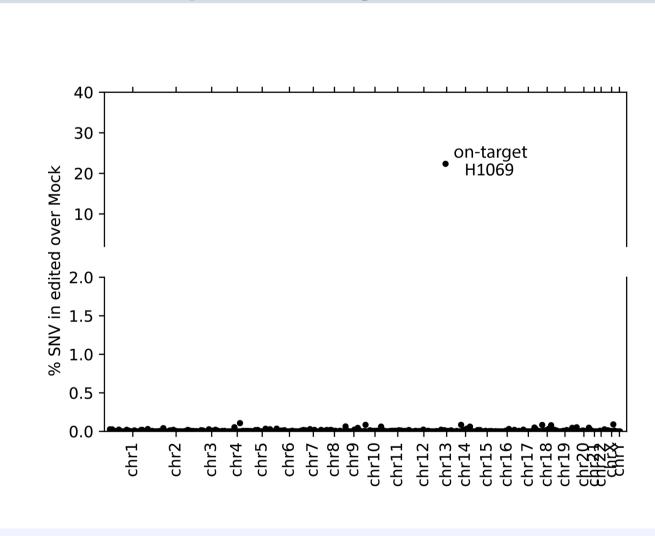
Improvements to Prime Editor cargo in Universal LNP enables therapeutic correction in fully humanized ATP7B H1069Q WD mouse models



- Optimization of LNP components improve Prime Editing to up to 80% of hepatocytes in humanized ATP7B H1069Q mouse models
- Dose-dependent editing was observed for H1069Q correction, exceeding 30% therapeutic threshold at clinically relevant dose

Preliminary off-target analysis for H1069Q Prime Editor does not identify off-target (SNVs or indels) events genome-wide





 Targeted sequencing analysis of off-target events following Prime Editing of H1069Q in patient-derived iPSC

Conclusions

- Successfully identified several candidate Prime Editor components capable of >60% editing for ATP7B p.H1069Q and p.R778L
- LNP-formulated Prime Editors using Prime Medicine's Universal liver targeted LNP efficiently correct (up to 80%) the H1069Q mutation in a fully humanized ATP7B Wilson's disease mouse model without detectable unintended edits
- Preliminary off-target analysis demonstrated H1069Q Prime Editors do not result in
- detectable off-target events in patient-derived iPSC LNP-formulated Prime Editors resolve copper accumulation in humanized mouse livers
- Using Prime Medicine's Universal LNP with a surrogate Prime Editor RNA cargo, up to 51% editing at the H1069 locus was observed in NHP in vivo
- Prime Medicine's Universal LNP is well-tolerated in NHP with a favorable safety profile
- These data support the advancement of a potential one-time, curative approach for Wilson's disease patients H1069Q or R778L mutations