



In vivo gene knockout followed by targeted gene insertion results in simultaneous reduced mutant protein levels and durable transgene expression

Anthony Forget, Ph.D. | October 25, 2019

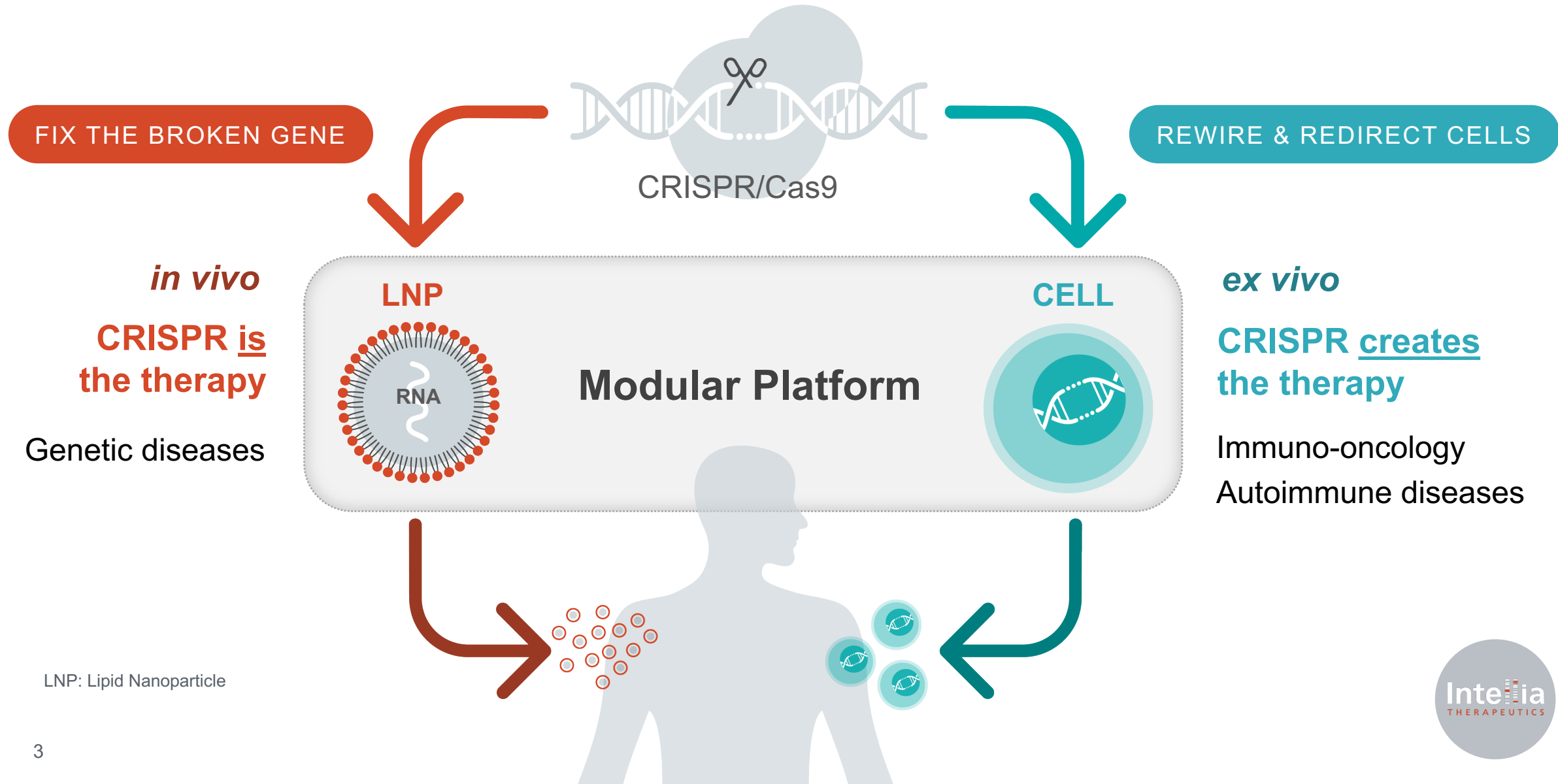
Disclosure: Employee of Intellia Therapeutics, Inc.

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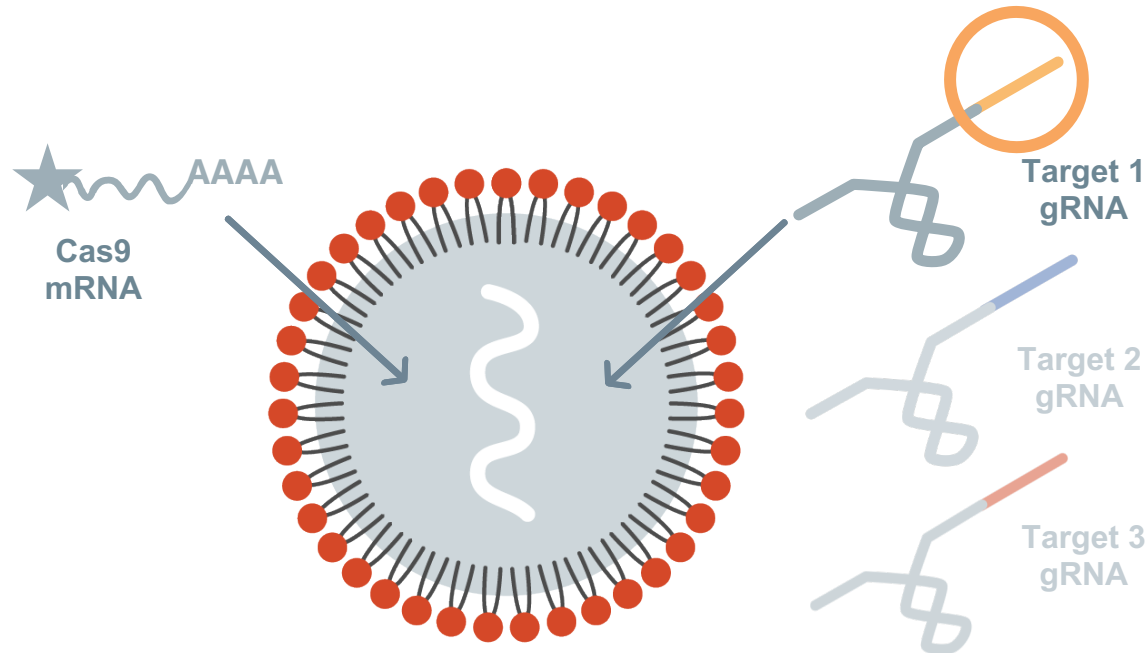
The Full Spectrum of Genome Editing for Rare and Genetic-Based Diseases



LNP: Lipid Nanoparticle

Intellia's Modular Non-Viral Delivery of CRISPR/Cas9 Addresses Disease at the Genetic Level

Lipid Nanoparticles (LNPs)



Variable portion of Intellia's modular LNP-based liver knockout approach limited to 20mer of gRNA

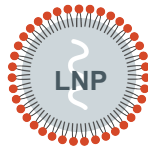
Key Advantages of LNP Delivery

- Redosing capability
- Low immunogenicity
- Transient expression
- Large cargo capacity for CRISPR/Cas9
- Scalable synthetic manufacturing
- Well-tolerated
- Biodegradable
- Adjustable range of tissue tropism

Transthyretin Amyloidosis (ATTR) is Treatable with a Gene Knockout in Liver

KNOCKOUT

Inactivation/deletion of
disease-causing DNA sequence



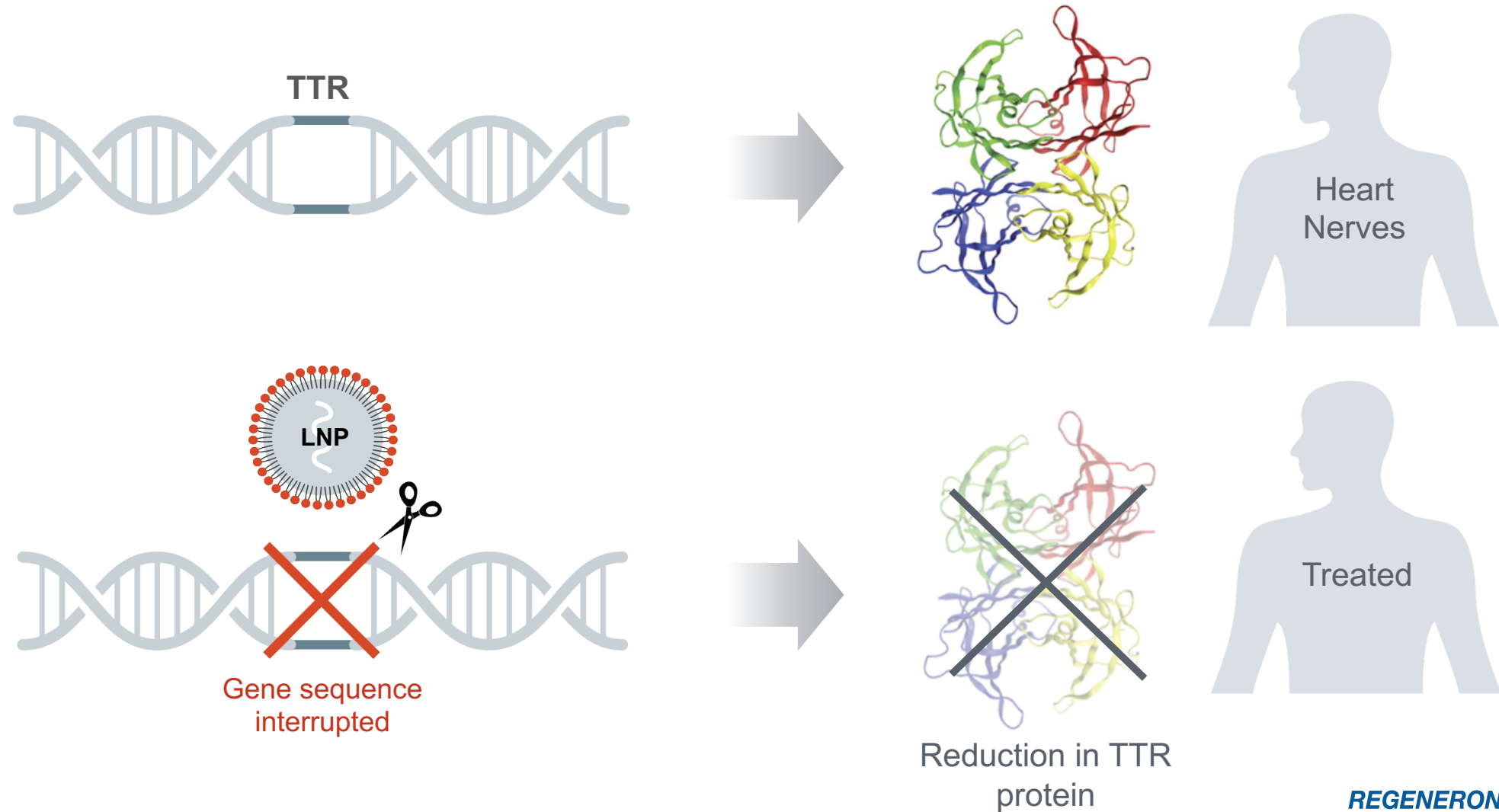
*Caused by accumulation of misfolded transthyretin (TTR) protein, which affects **nerves, heart, kidneys and eyes***

Autosomal dominant; >120 known mutations knock-down of disease causing protein is a clinically validated strategy

Deletion of disease gene in hepatocytes reduces supply of misfolded protein

Efficient Knock-Out Approach Interrupts the TTR Gene Sequence

Reduces TTR Protein Production in Liver



REGENERON

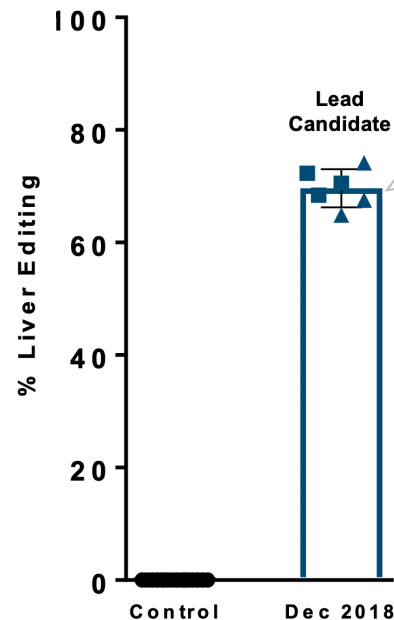


LNP Platform Achieves Sustained Gene Knockout in NHPs After a Single Dose

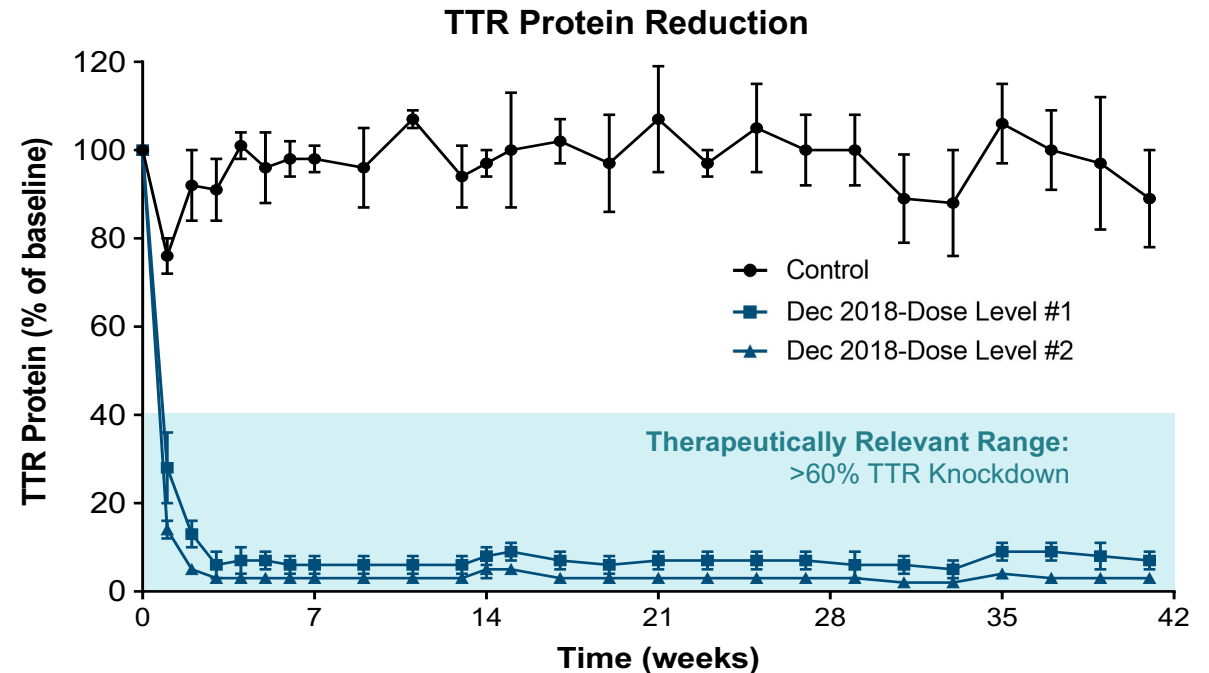
- In vivo knockout is **achievable**- facilitated by combining LNP delivery of Cas9-mRNA and synthetic gRNA
- In vivo knockout is **efficient**- a single dose administration of LNP achieves a durable reduction in circulating protein
- The knockout platform is **modular**- only 20 nucleotides of the gRNA sequence need to be changed to enable editing of a different genomic locus

Single-Dose TTR Editing

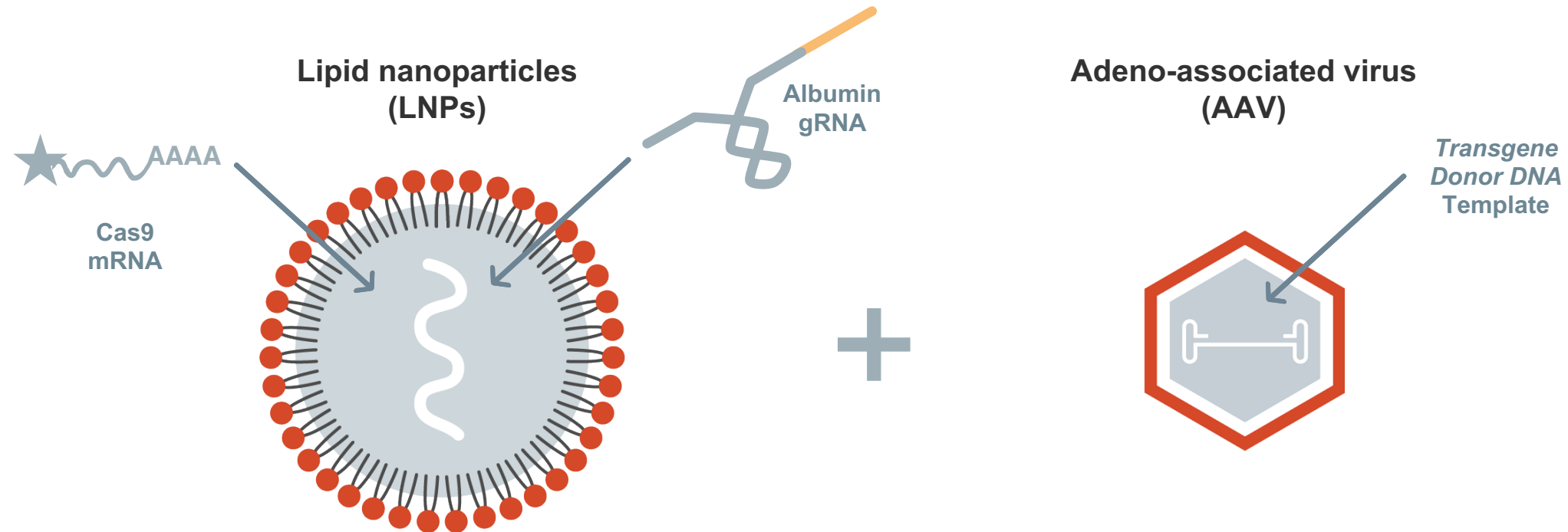
Chart includes single administration within a range of dose levels



>95% Reduction in Circulating Levels of TTR



Hybrid LNP-AAV Delivery of CRISPR and Transgene Template is an Effective Modular Approach for Targeted Gene Insertion



Hybrid LNP-AAV delivery system precisely integrates into the genome, resulting in durable expression, and utilizes the endogenous promoter to drive transgene expression

REGENERON



Hemophilia B is Treatable with Gene Insertion in Liver

INSERT

Insert new DNA sequence to produce therapeutic protein



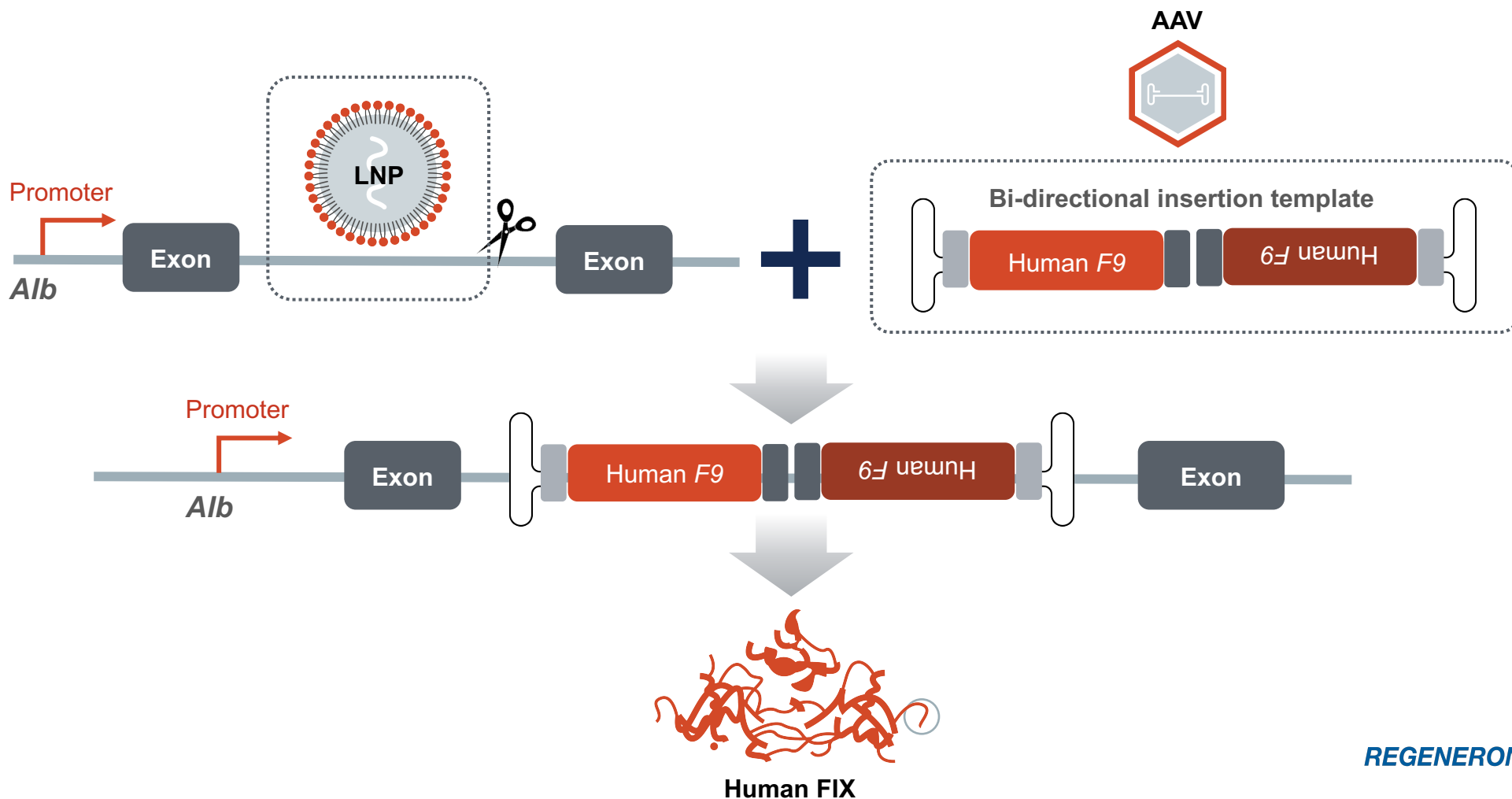
*Rare X-linked genetic disorder caused by missing or defective **Factor IX (FIX)**, a blood-clotting protein encoded by the **F9 gene***

Severe cases often have painful, spontaneous bleeding into joints

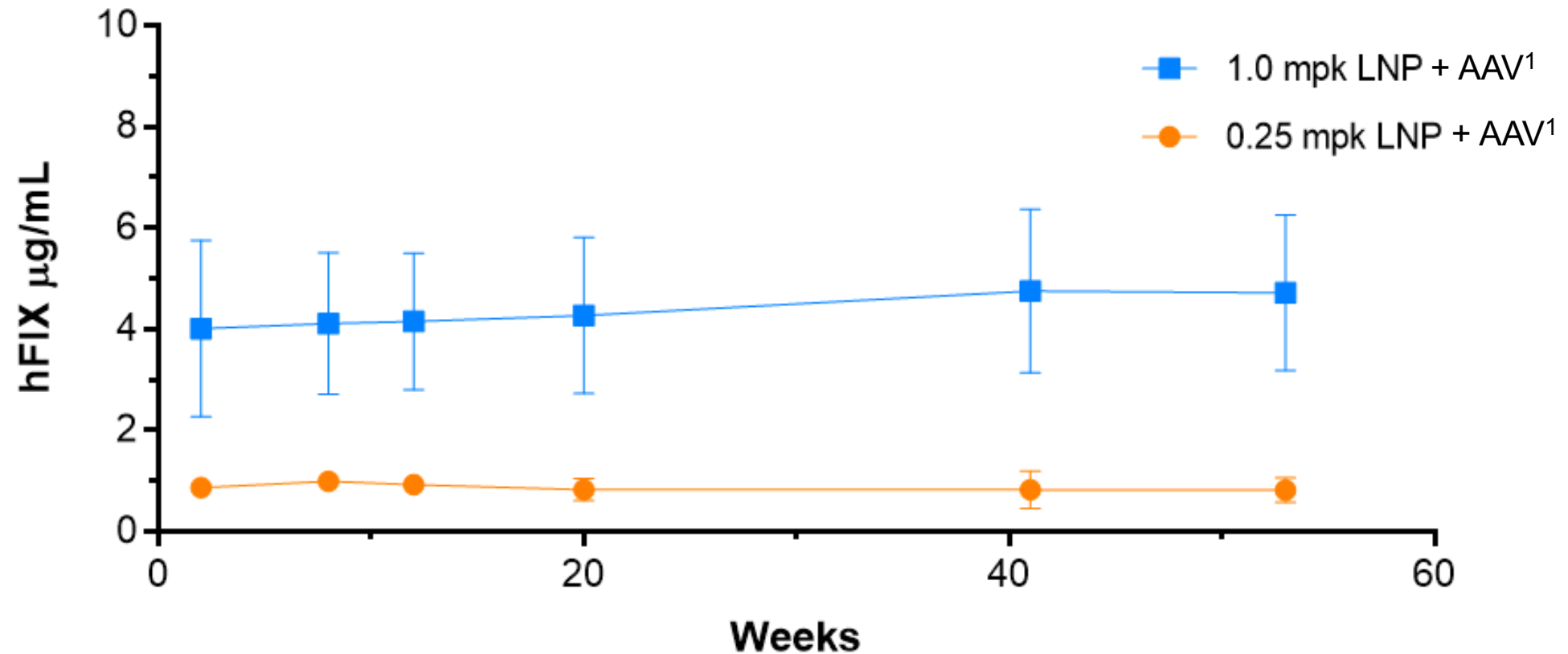
Patients treated chronically with Replacement Factor IX

In Vivo Insertion of Factor 9 Gene at Albumin Intron Safe Harbor Site

Hybrid Delivery System Precisely Integrates Into the Genome



FIX Levels in Adult Mice Are Stable through the completion of a 1-Year Durability Study



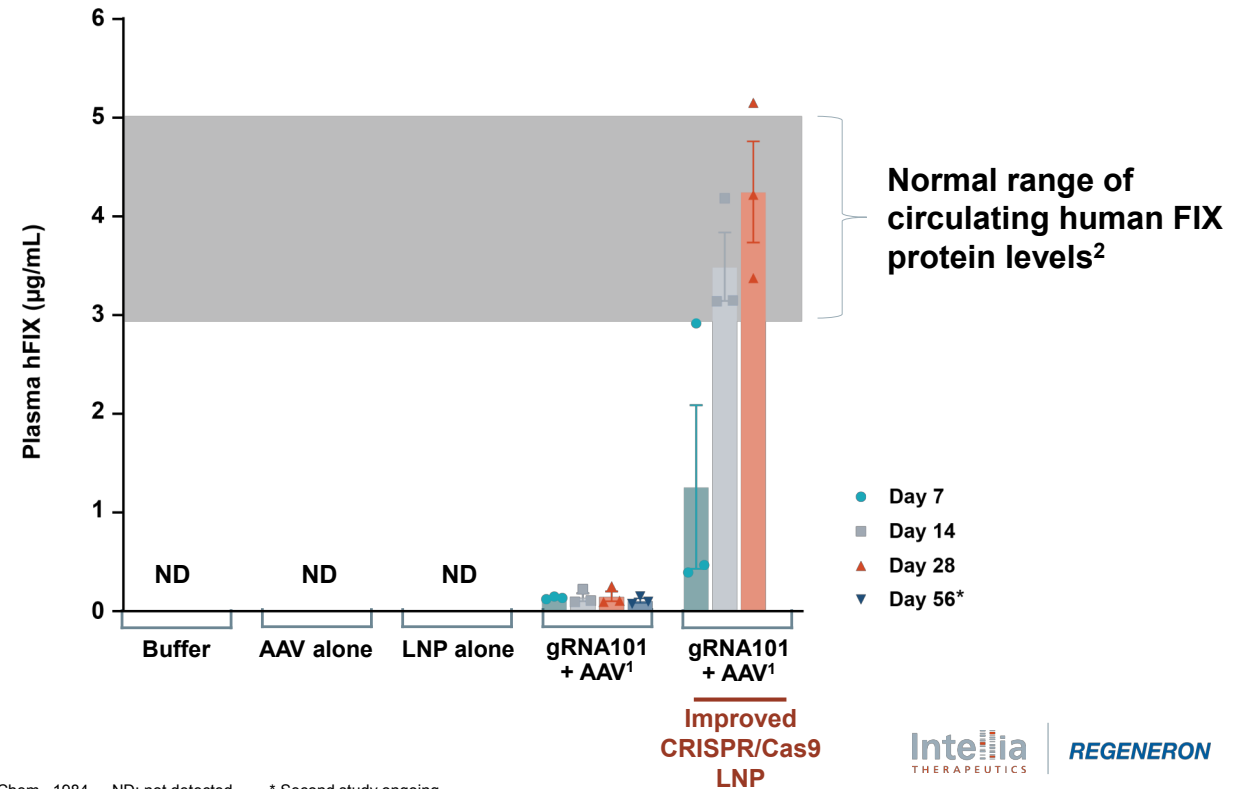
¹AAV MOI 3e11 vg/kg

Effective *In Vivo* Targeted Gene Insertion in NHPs Demonstrated with Hybrid LNP-AAV Delivery Platform

- In vivo insertion is **achievable**-facilitated by combining LNP CRISPR delivery with AAV transgene template delivery
- The insertion platform is **tunable**-protein expression levels can be varied by changing any of three components; gRNA, LNP dose, and/or AAV dose
- The insertion platform is **modular**-only the AAV template sequence needs to be changed to insert other genes of interest

Physiologically Normal Levels of Circulating Human FIX Protein Achieved With Insertion of *F9* in NHPs and Maintained Through Day 28

Baseline albumin levels maintained at day 28



12

¹AAV MOI 3e13 vg/kg

²Amiral et al., Clin. Chem., 1984

ND: not detected

* Second study ongoing

As presented during ASGCT 5/2019

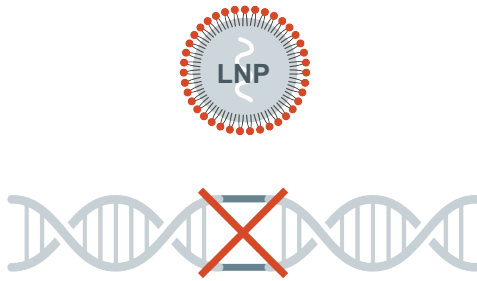
REGENERON



Alpha-1 Antitrypsin Deficiency (AATD) is Treatable with a Combination of Gene Knockout and Gene Insertion

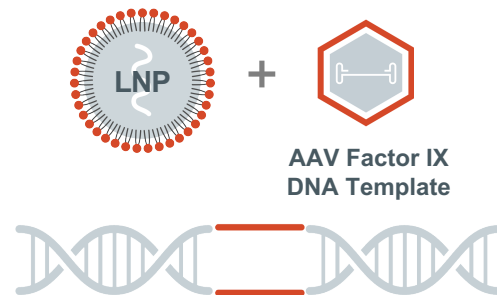
KNOCKOUT

Inactivation/deletion of disease-causing DNA sequence



INSERT

Insert new DNA sequence to produce therapeutic protein



*Caused by mutations in the SERPINA1 gene which encodes Alpha-1 Antitrypsin(AAT) protein, commonly leading to **lung dysfunction** and **liver disease***

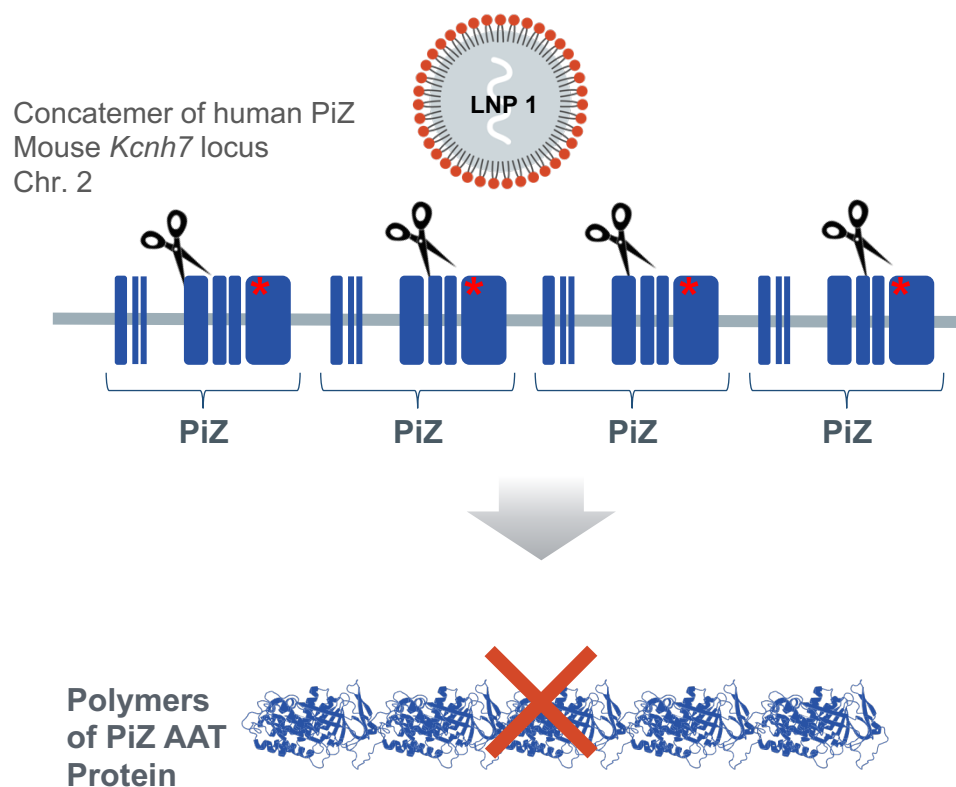
The most severe genetic mutation codes for a single amino acid substitution, E342K, known as the **PiZ allele**

Normally, Alpha-1 Antitrypsin protein (AAT) is produced in the liver, secreted into circulation and acts to inhibit various proteases

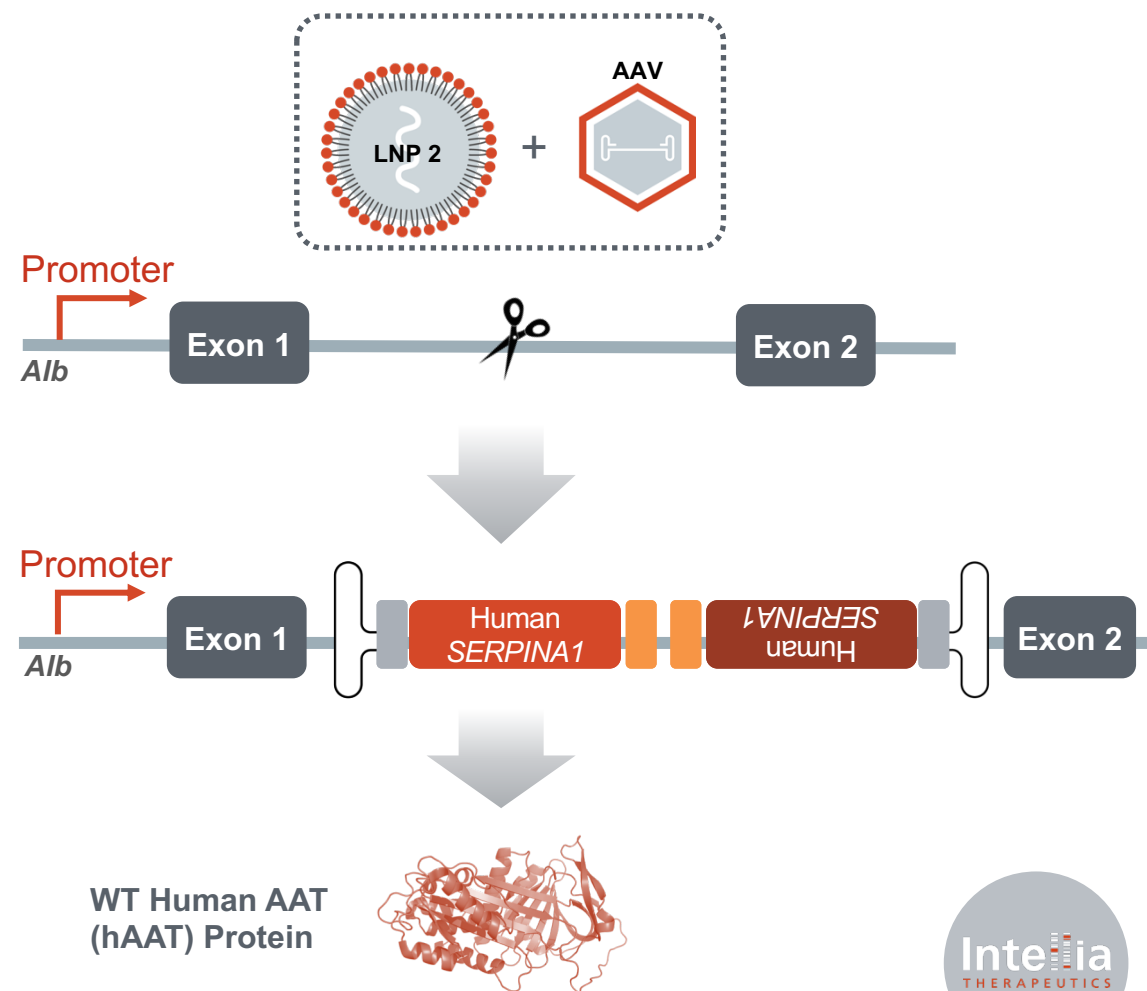
Hepatic accumulation of toxic polymers coded by the PiZ allele can lead to liver disease

Combining the Two Modalities: A Consecutive Knockout and Insertion Approach to Eliminate PiZ Expression and Restore Protease Inhibition Function

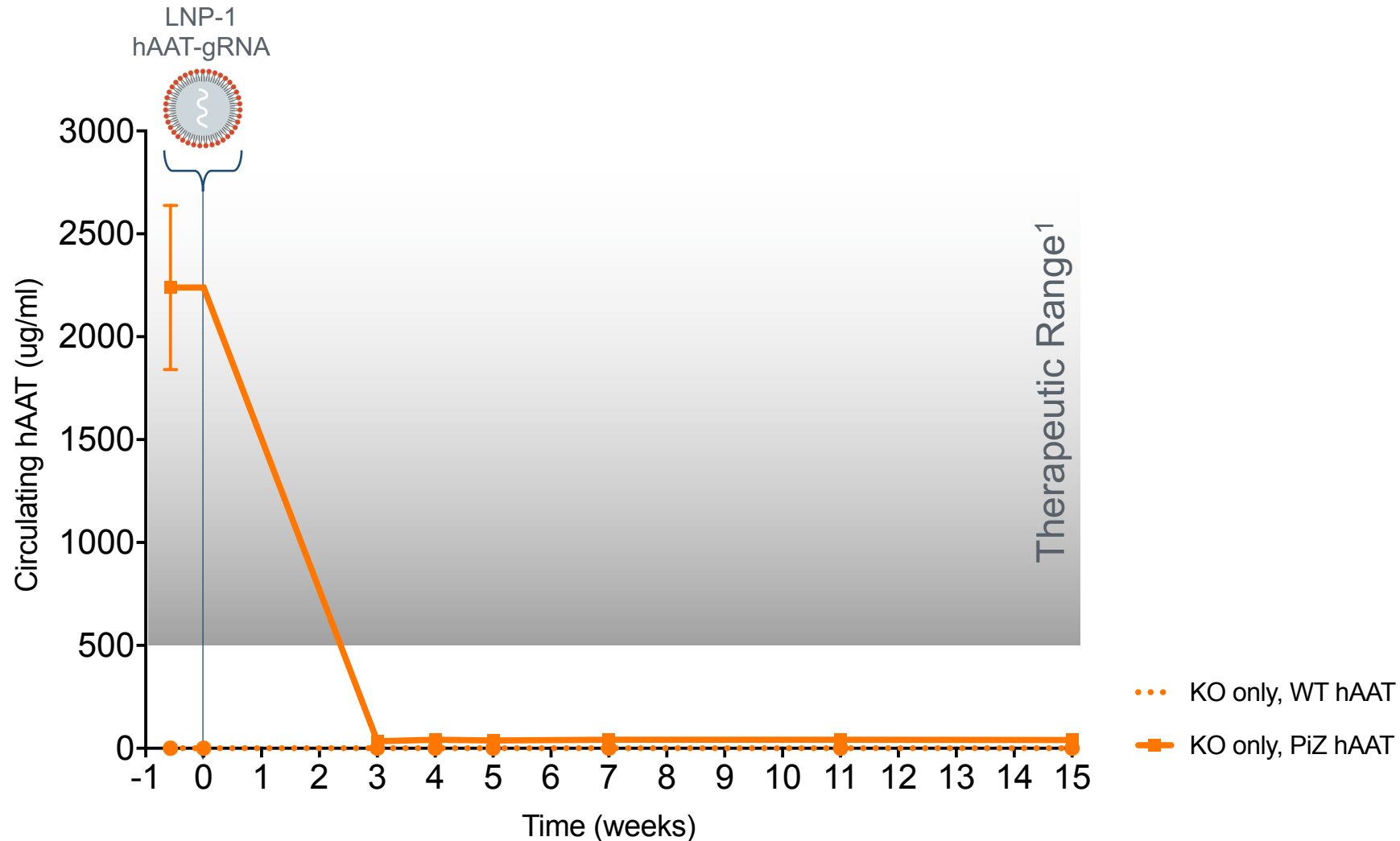
in vivo Knockout of PiZ Allele



in vivo Insertion of *SERPINA1*



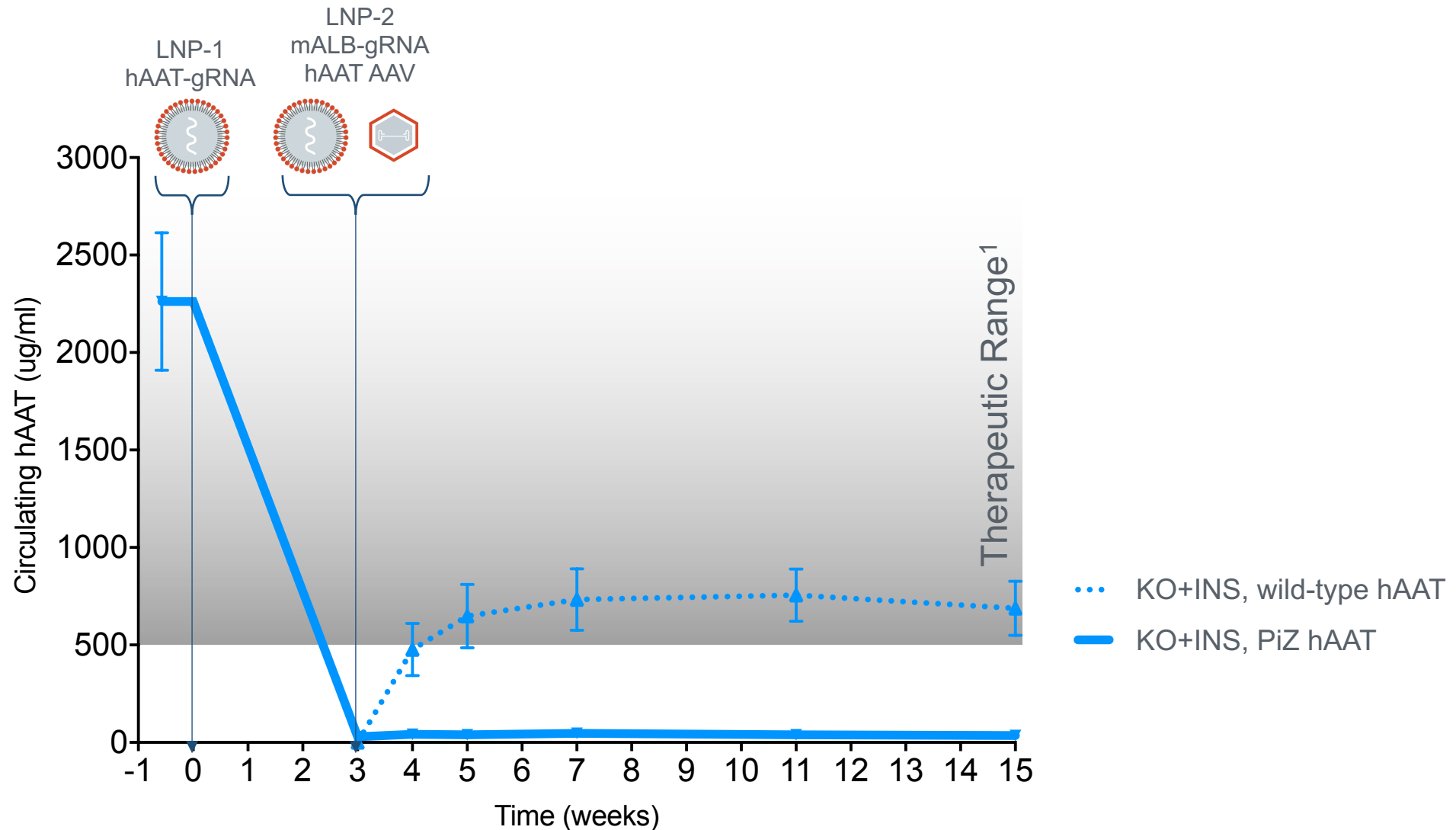
Gene Knockout Results in >98% Reduction in Circulating hAAT PiZ Protein in the PiZ Mouse Model



LC-MS/MS assay that differentiates wild-type from PiZ hAAT

¹Stoller & Aboussouan The Lancet, 2005

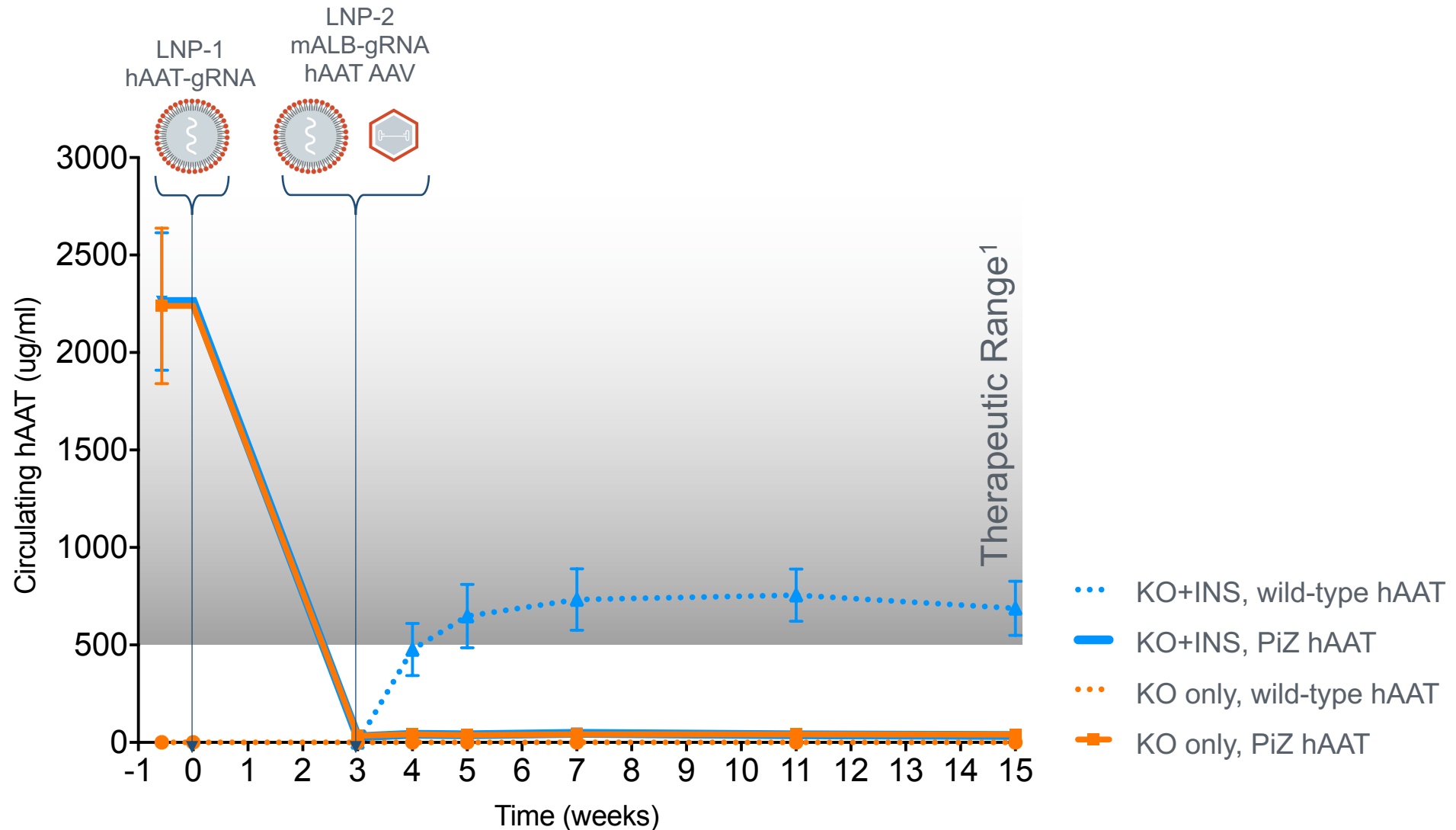
Subsequent Gene Insertion Results in Circulating Therapeutic levels of hAAT Protein



LC-MS/MS assay that differentiates wild-type from PiZ hAAT

¹Stoller & Aboussouan The Lancet, 2005

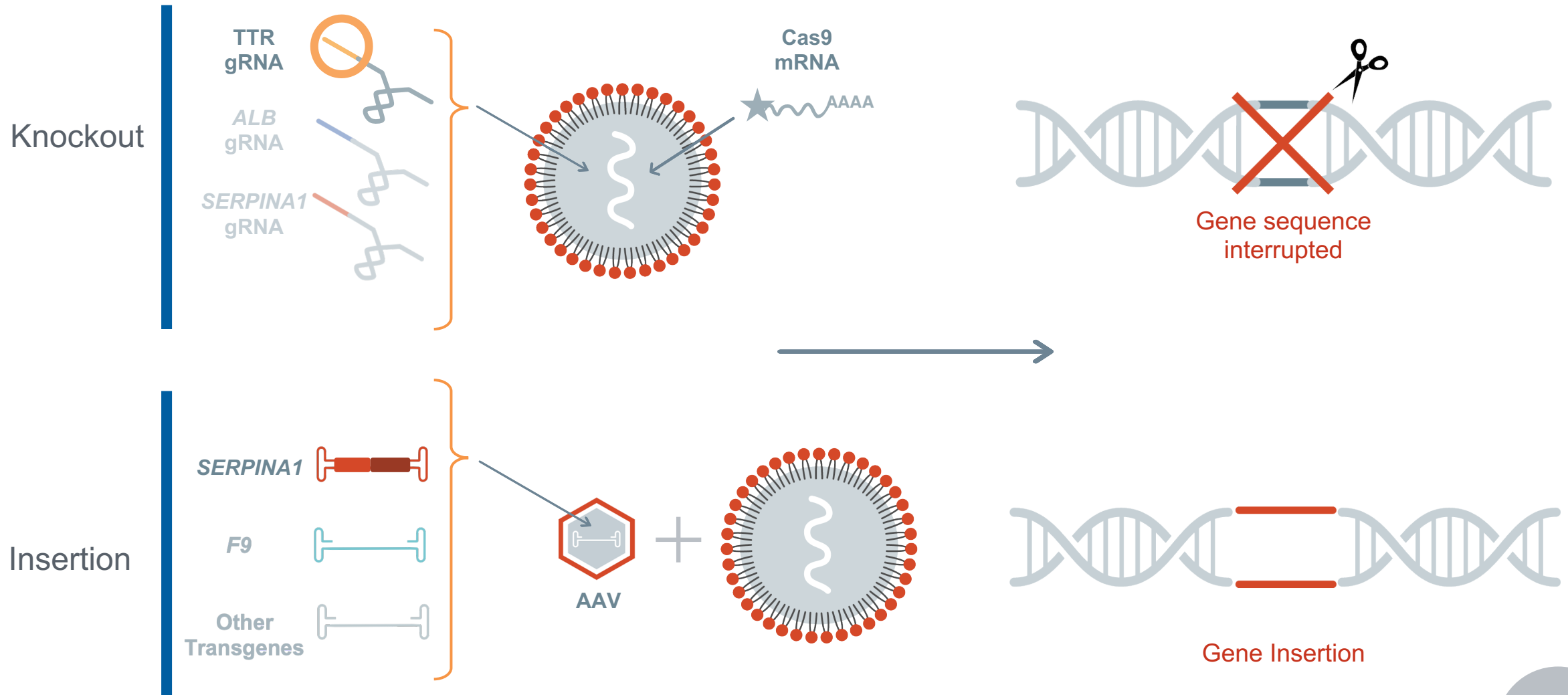
The Modularity of our Gene Editing Platform Enables Consecutive Edits to Achieve Knockout of a Faulty Protein and Expression of a Functional Protein



LC-MS/MS assay that differentiates wild-type from PiZ hAAT

¹Stoller & Aboussouan The Lancet, 2005

Modular Platforms for Gene Knockout and Targeted Gene Insertion can be Applied as Individual Modalities or in Combination



Key Takeaways

- Modalities from Intellia's platform can be combined to broaden the landscape of diseases that can be potentially addressed by gene editing
- First demonstration that consecutive dosing of two LNPs in adult mice can **achieve two distinct targeted gene editing events resulting in reduction of protein expressed from a faulty gene and restoration of activity from insertion of a functional gene**

Acknowledgements

Intellia team

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Intellia
THERAPEUTICS