

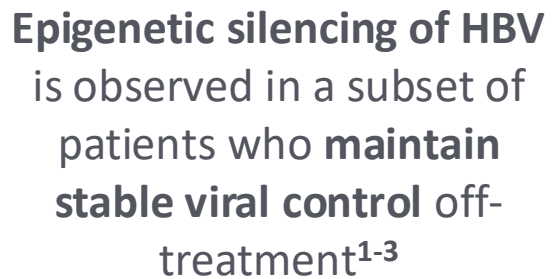
**Tune-401:  
a First-in-class Epigenetic Silencer  
Developed for the Treatment of  
Chronic Hepatitis B**

**TUNE THERAPEUTICS**

AASLD 18<sup>th</sup> November 2024



**TUNE**  
THERAPEUTICS



<sup>1</sup> *Suslov et al., J Hepatology (2021)*

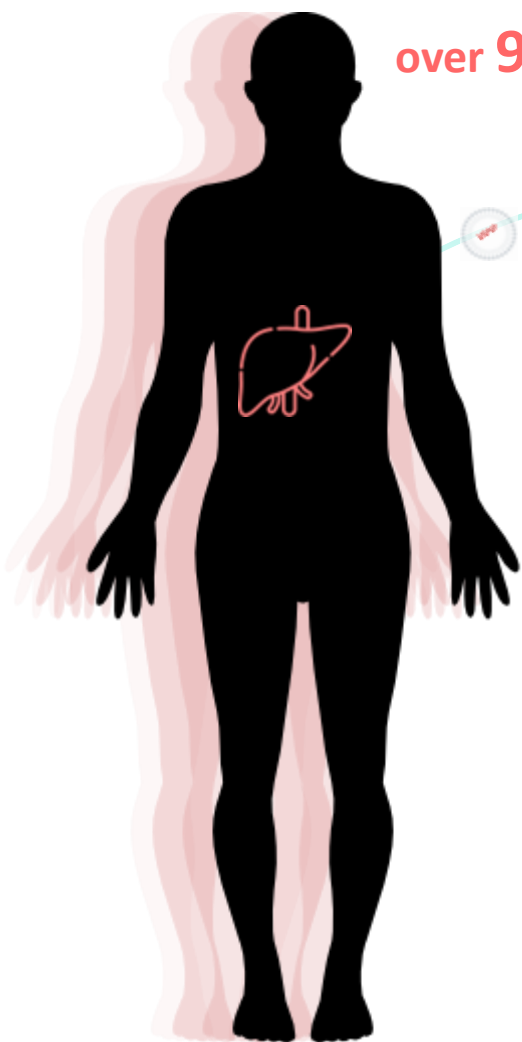
<sup>2</sup> *Pollicino et al., Gastroenterology (2006)*

<sup>3</sup> *Lebosse et al., Scientific Reports (2020)*

# Tune-401: a precise and durable epigenetic silencer of HBV DNA



Tune-401 target site found in  
**over 98%** of HBV genomes



5

Durable off-treatment  
reductions in:

- HBV proteins
- pgRNA
- viremia

= core elements  
of **functional cure**

4

Corresponding **reduction**  
in **HBV RNA** (all sources)

1

**Tune-401** is a liver-targeted LNP encapsulating **mRNA** (encoding the epi-silencer) and **gRNA** (specifying the HBV target)

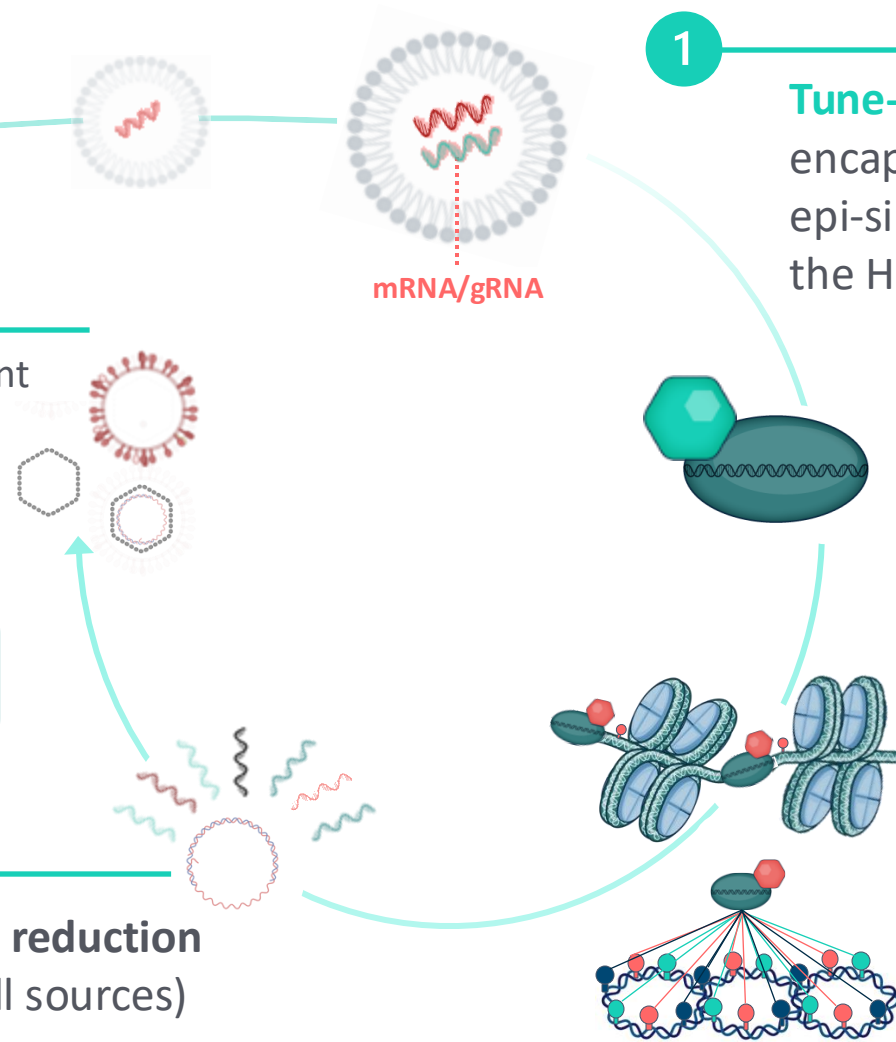
2

**mRNA** is **translated** into HBV epi-silencing **protein**, which complexes with gRNA to find HBV targets

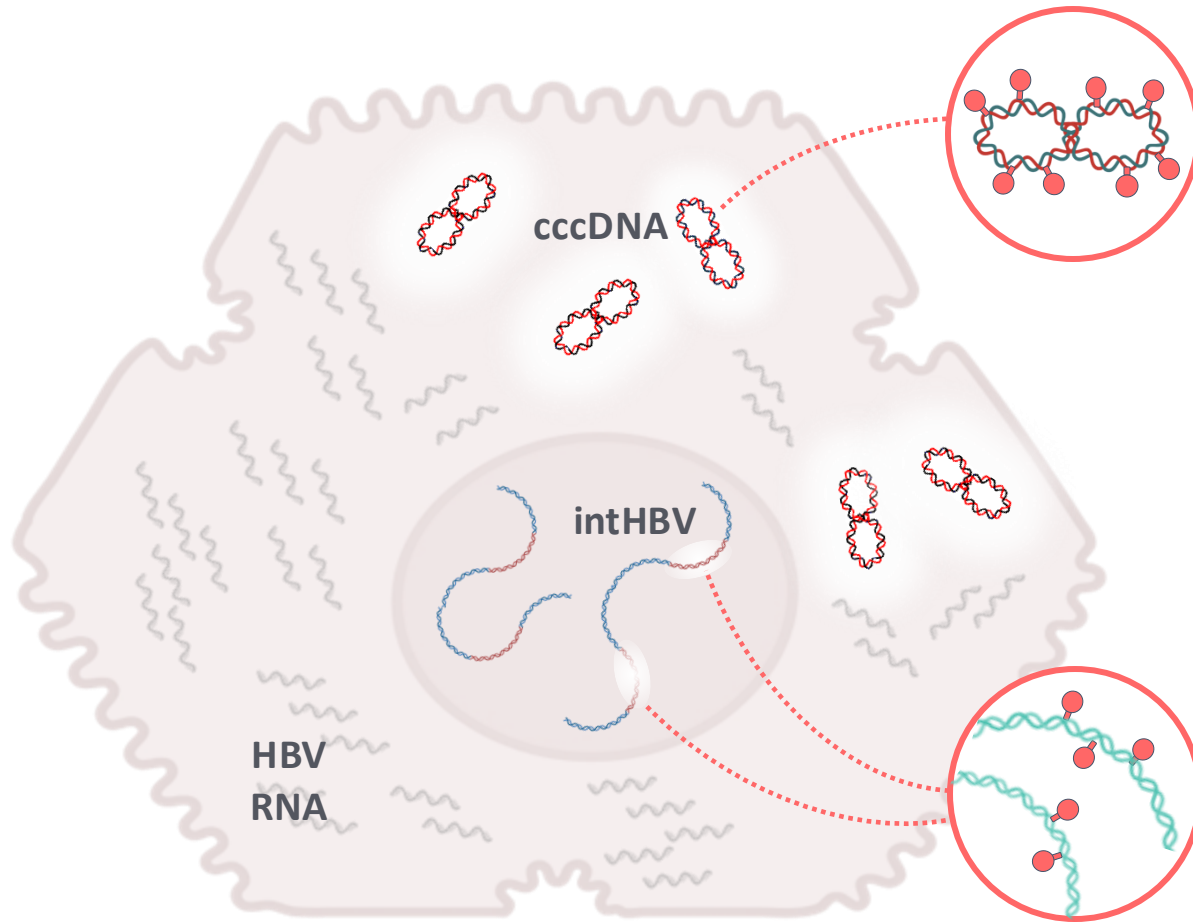
3

**Methylation of integrated HBV DNA**

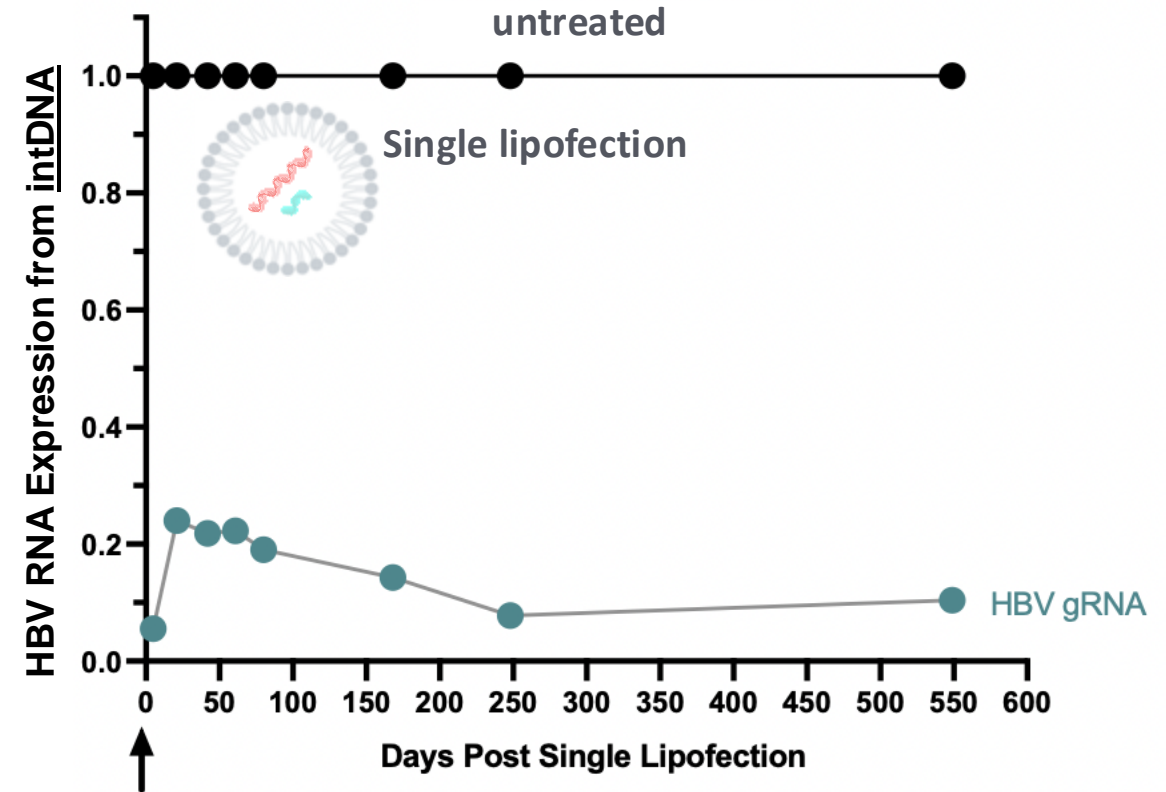
**Methylation of HBV cccDNA**



# Why is Tune-401 different than other HBV therapies?

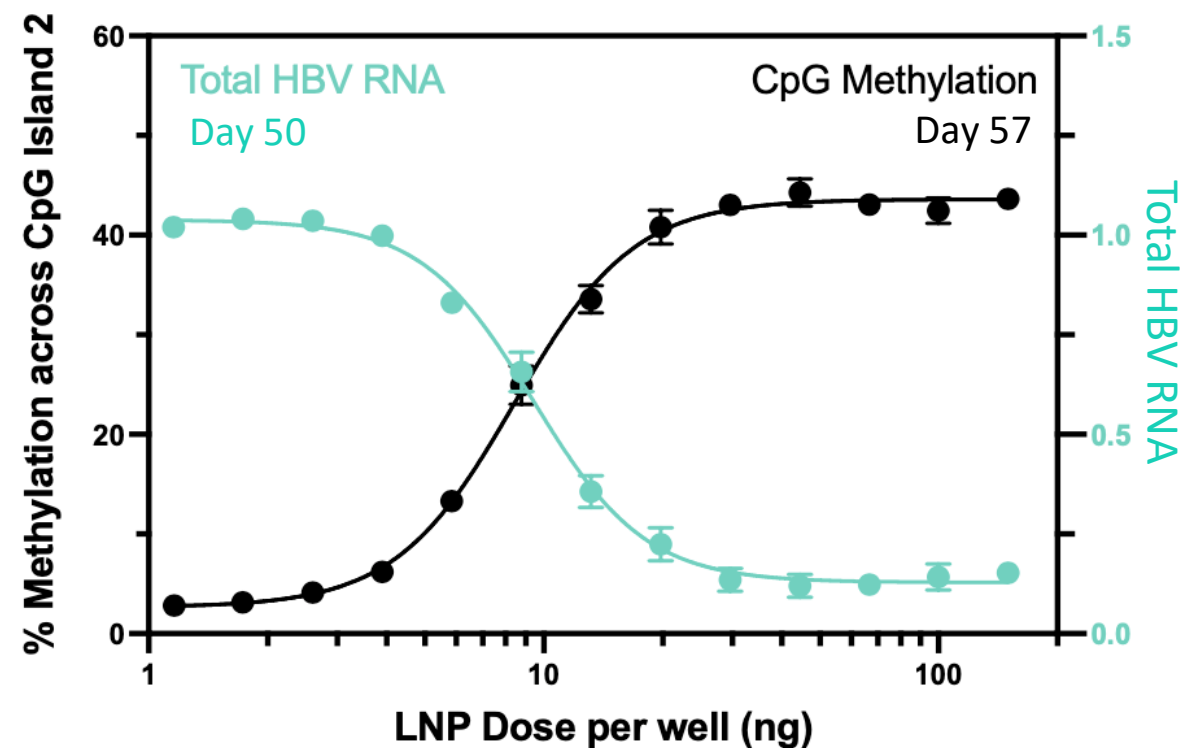
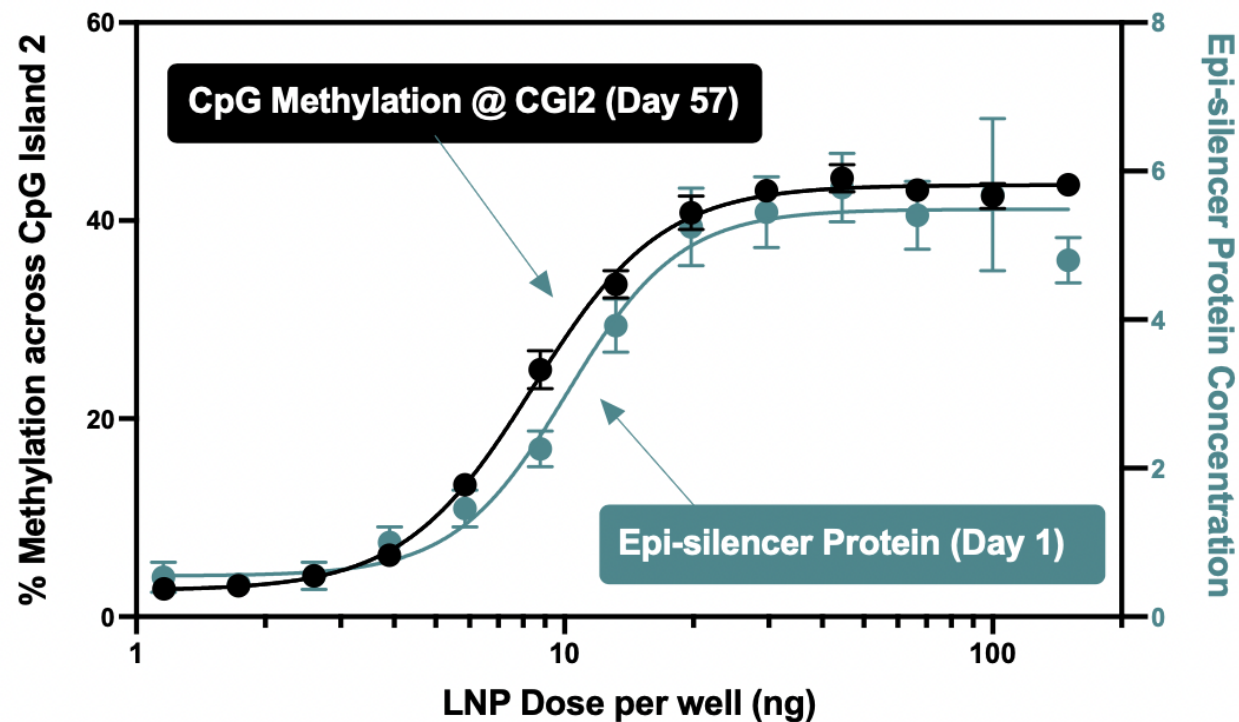
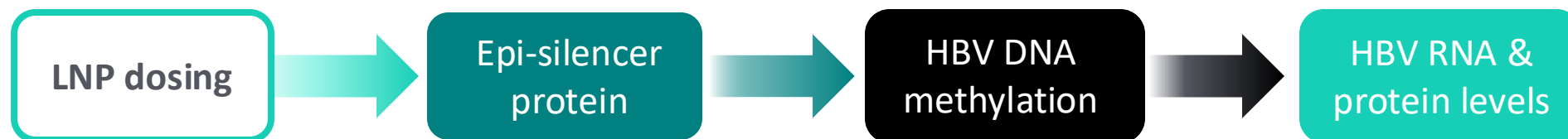


First therapy to **silence transcription** from intHBV + cccDNA **without cutting genome**



**High durability** following transient LNP delivery (without **host immune clearance**)

# Strong, durable silencing of **integrated HBV DNA** in Hep3B cells



# Near complete repression of 3.5kb HBV RNA transcription from **cccDNA** in human hepatocytes



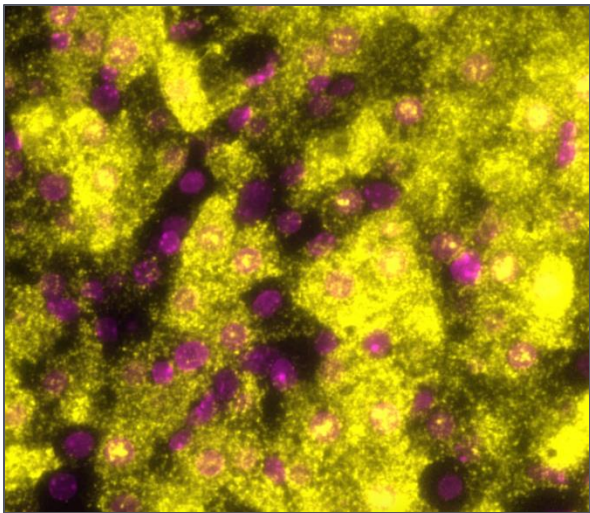
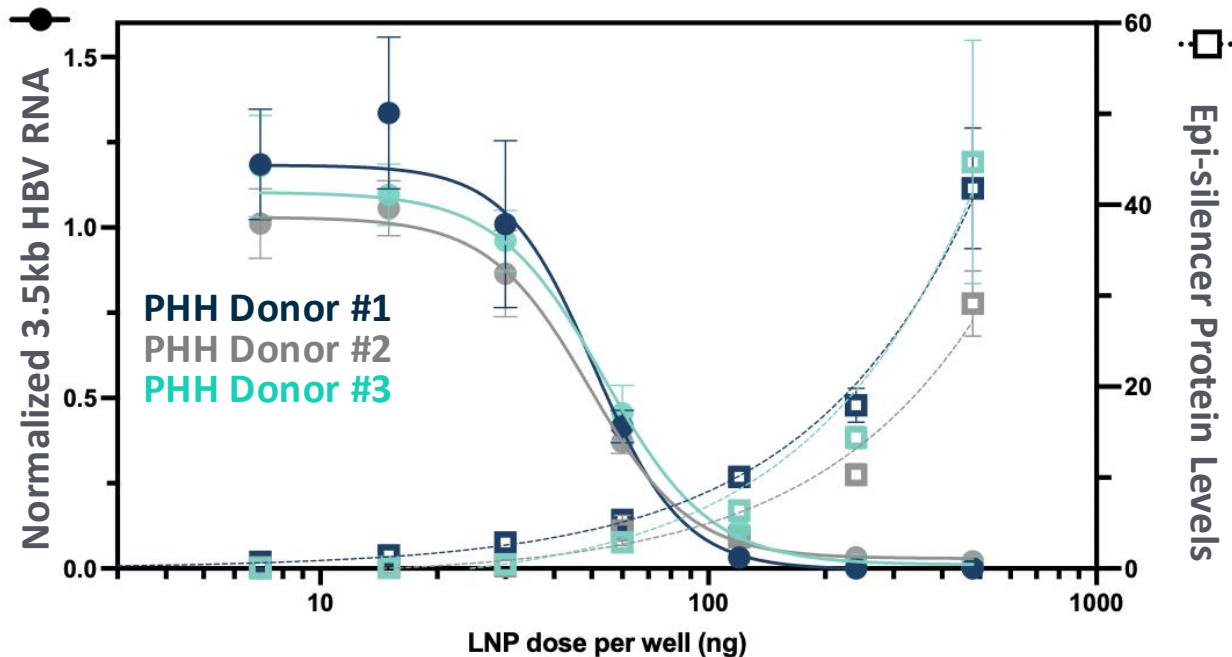
PHH + HBV

A single dose of Tune-401 repressed as much as **99.99%** of 3.5kb HBV RNA from physiologic ranges of cccDNA per cell

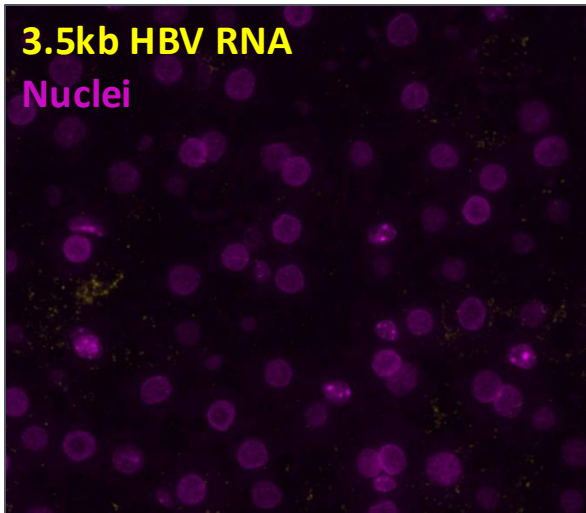


FRG Human  
Chimeric Liver  
Mouse + HBV

Near complete 3.5kb HBV RNA repression found in huHeps that received LNP. ISH quantification showed average population shift from 67 copies/cell at baseline to 0 copies/cell in cells that received Tune-401 (N= >17k huHeps/mouse)



Non-Targeting Epi-silencer



Tune-401 Epi-silencer

Baseline values: 90% humanization, ~3 cccDNA/cell, 45k IU/mL HBsAg

# Assessing the specificity of Tune-401 for the HBV genome



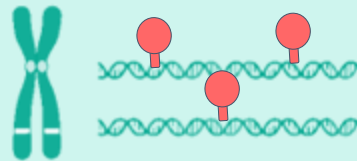
## *In Silico* Analysis



### Computer prediction for off-target sites

Identifies sites similar to the Tune-401 target site in **1,000 Genomes Db**

## Methyl-seq Analysis



Human methylome capture panel (3.98M CpGs)  
Identifies CpGs with differential **methylation**

## RNA-seq Analysis



### Transcriptome-wide sequencing (28k transcripts)

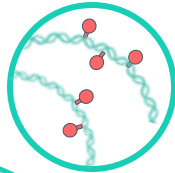
Identifies transcripts with differential **expression**

# Assessing the specificity of Tune-401 for the HBV genome



**0 sites**

< 3 mismatches



**0.025%**

of CpGs with diff.  
methylation  
compared to  
untreated

**2**

transcripts with  
differential expression  
compared to  
untreated

**0**

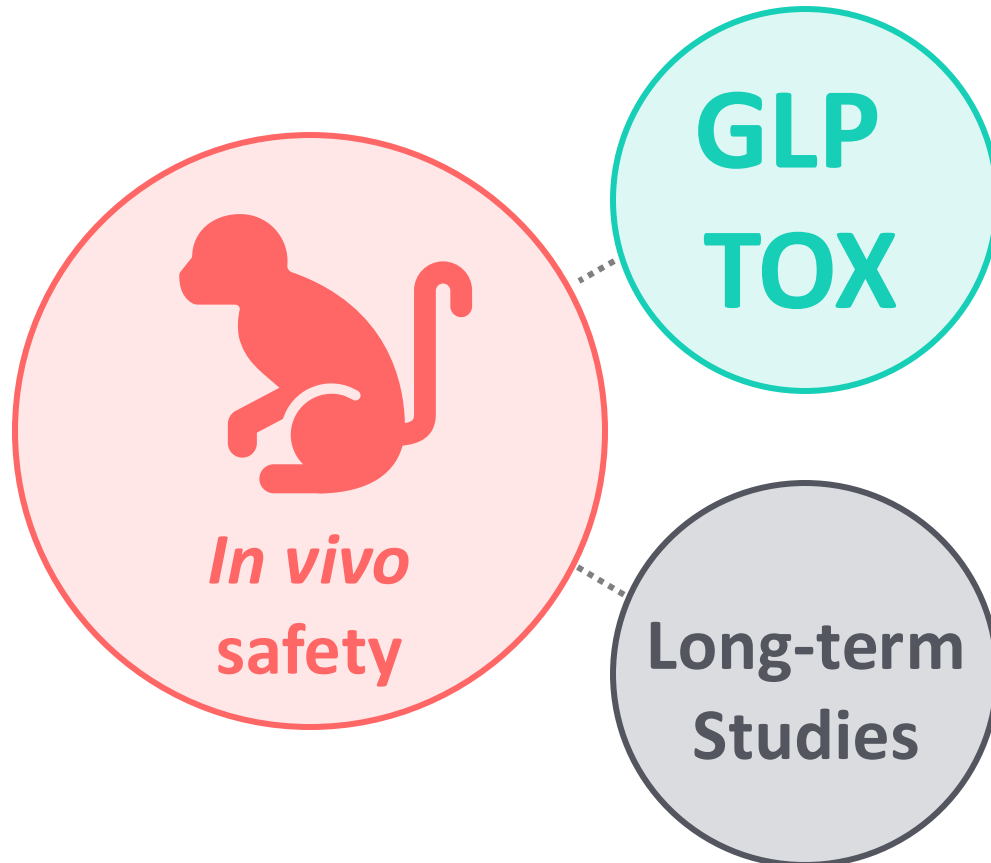
No overlap between genes with differential gene  
expression and regions with differential methylation  
*Even at 3xIC90 dose, this overlap is 2 genes*

## Sequencing Readouts

- Performed across 3 PHH donors
- 3 dose levels: IC50, IC90, 3xIC90
- Compared to untreated cells
- High degree of coverage: ~4,800 reads per unique transcript, 165 reads per unique CpG.
- Also performed sequencing analysis in primary cells from tissues with LNP biodistribution

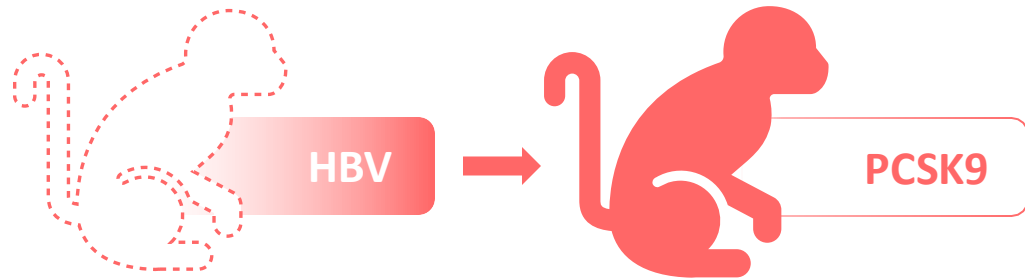
IC90

# Assessing the safety of Tune-401 in large animal models



- NOAEL determined to be highest dose tested (3 mg/kg)
  - Transient ALT/AST elevations that resolved by 14d post-dosing
  - No cage-side observations
  - No adverse histopathology findings
  - No Tune-401 detected in germline cells (RNA and protein)
- 
- No adverse clinical observations on NHPs dosed with surrogate research epi-editor (now at ~600d post-dosing)

# Demonstrating the Tune-401 MoA in a large animal model

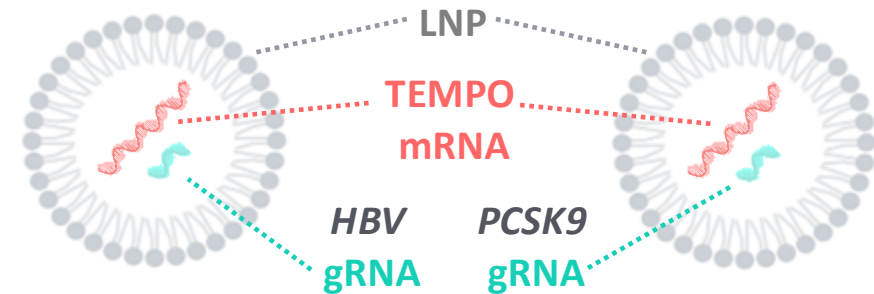


No large animal  
model available  
for HBV

Surrogate liver target  
in established  
NHP model

## Same mode of epigenetic action

Methylation of CpG islands leads to  
repression of transcription initiation



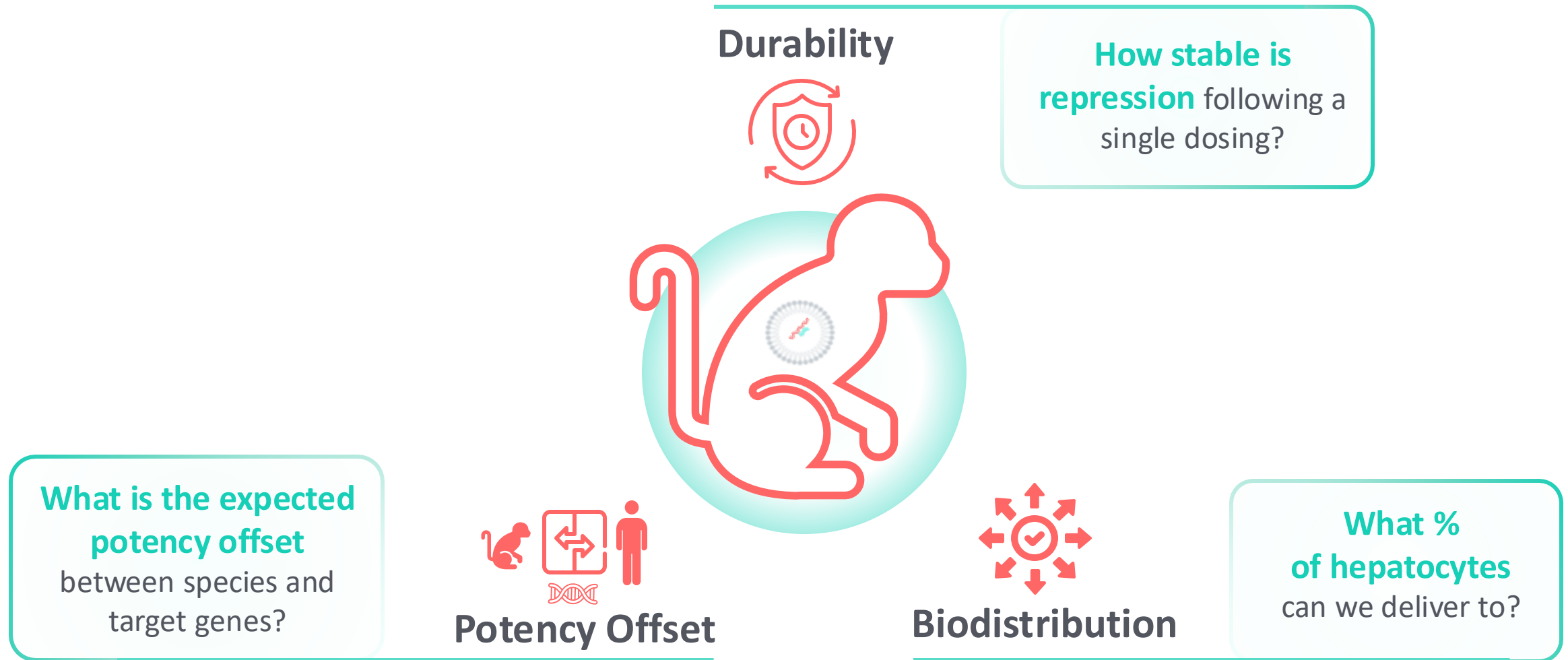
Tune-401

PCSK9  
Surrogate

## Same delivery vehicle and effector protein

LNP and epi-silencing protein identical –  
only the targeting guide RNA differs

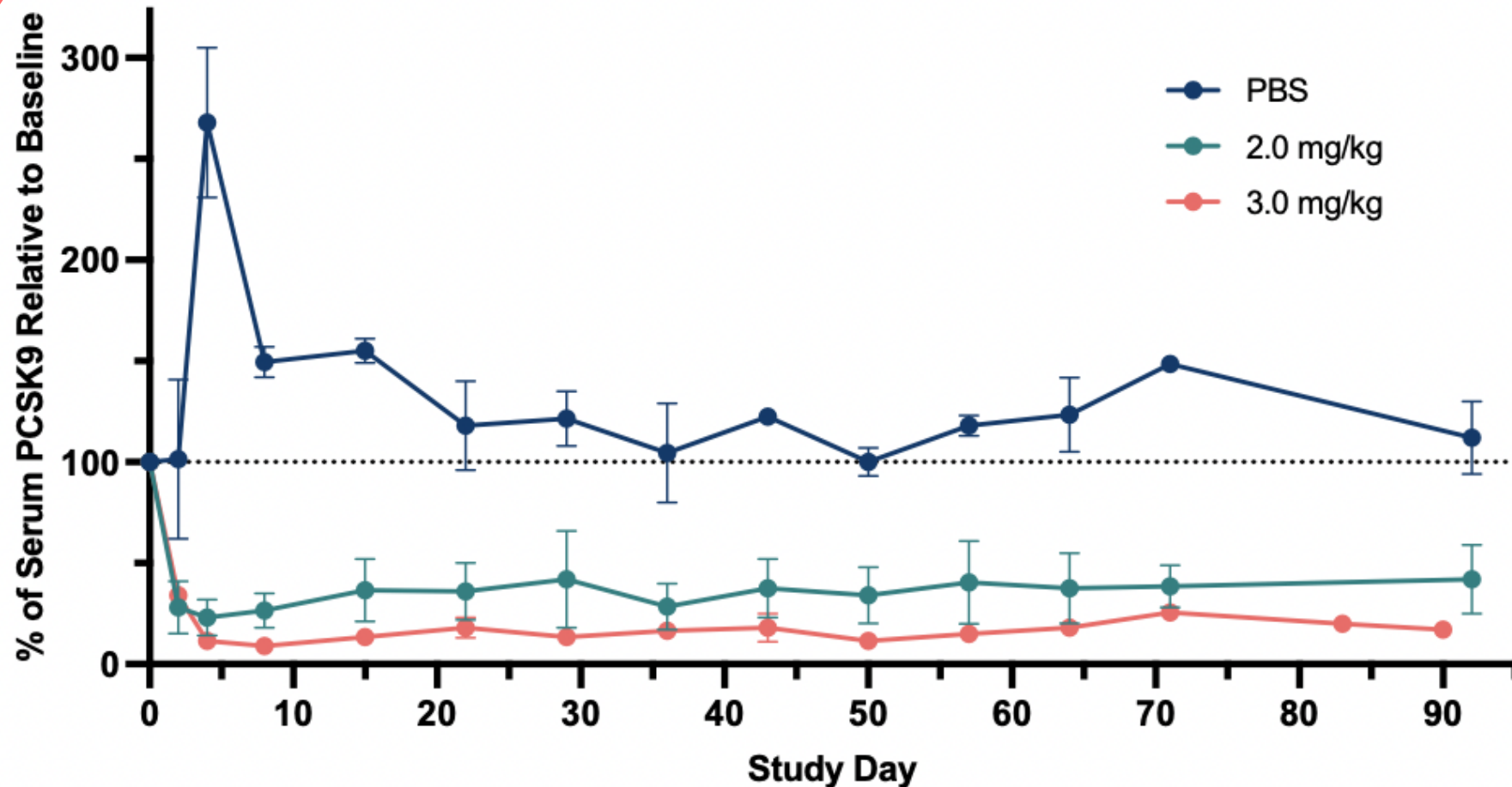
# Demonstrating the Tune-401 MoA in a large animal model



# Surrogate PCSK9 epi-silencer shows strong and dose-dependent repression of PCSK9 in NHP



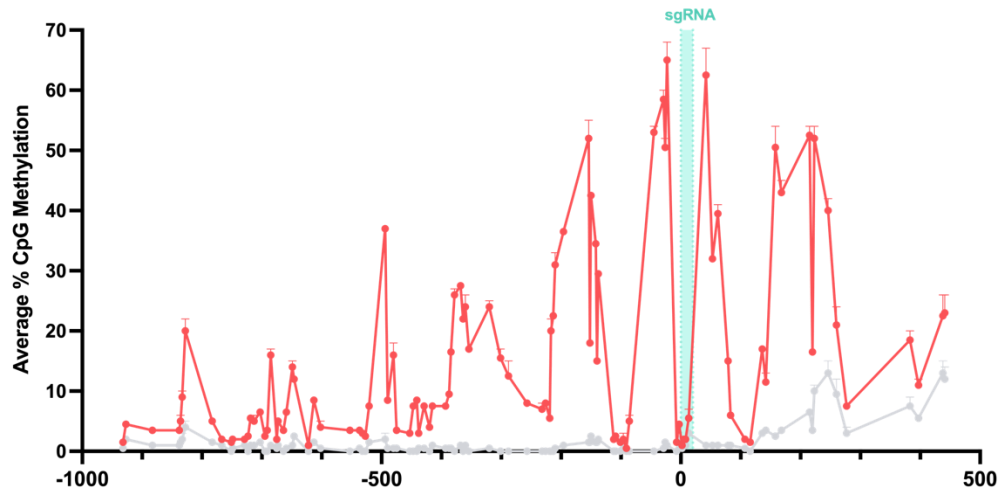
## PCSK9



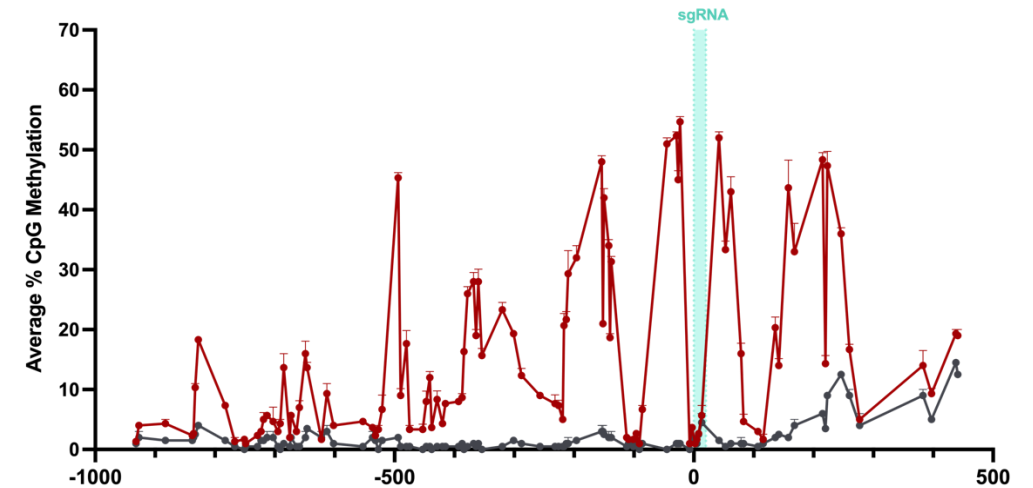
# Strong durability of deposited CpG Methylation over time



**Methylation @ 8 days post dose**



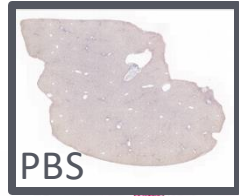
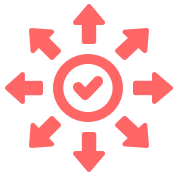
**Methylation @ 365 days post dose**



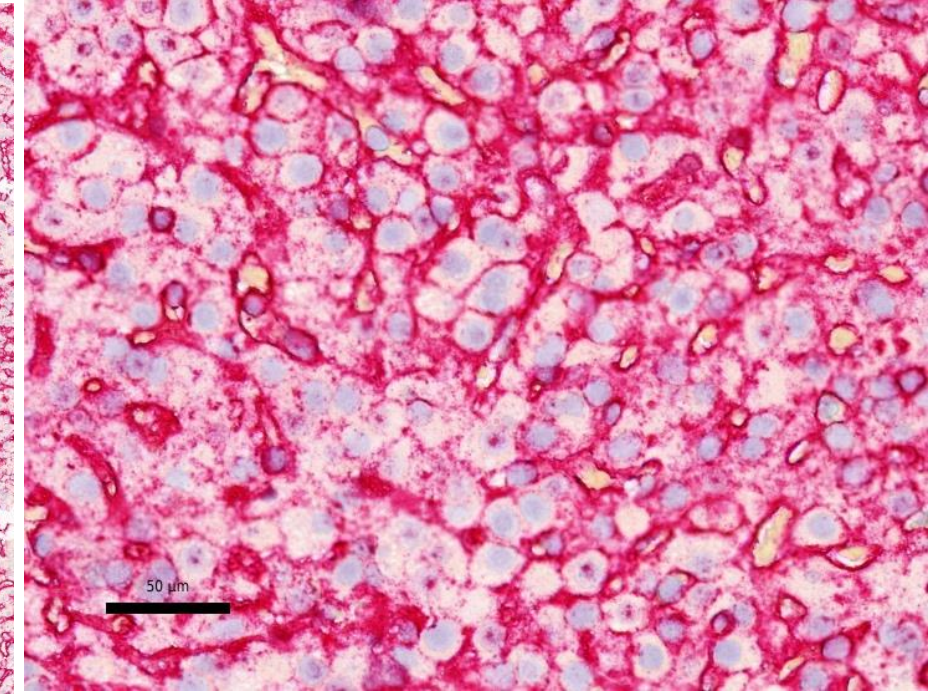
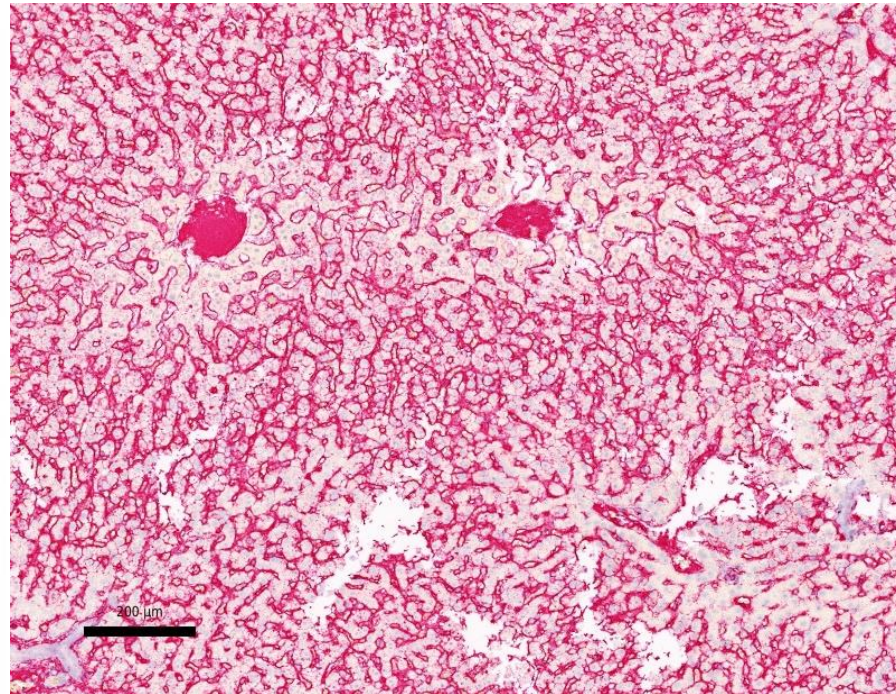
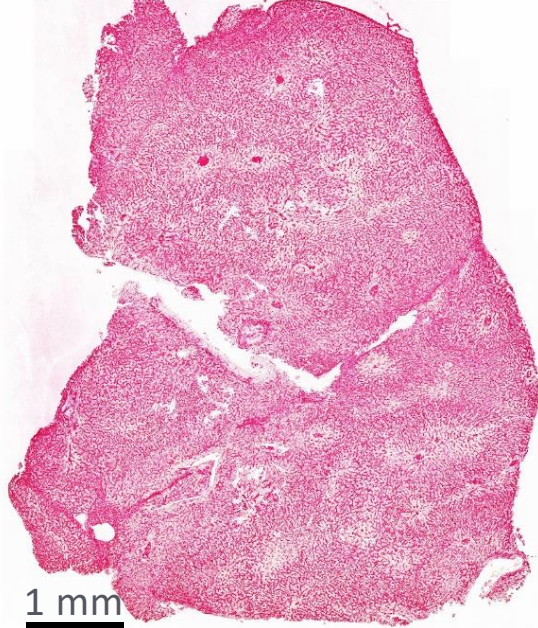
**PCSK9**

Deposited CpG Methylation around the PCSK9 proxy target locus was highly stable between **8d post-dose** and **365d post-dose**

# Strong biodistribution of PCSK9 epi-silencer in NHP livers shows functional delivery to nearly all hepatocytes

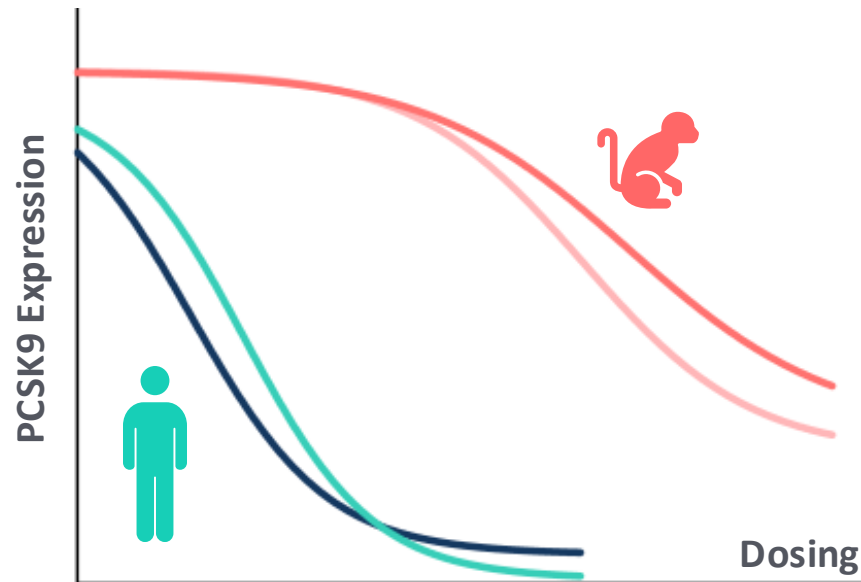


Epi-silencer mRNA ISH @ 6hr post-delivery



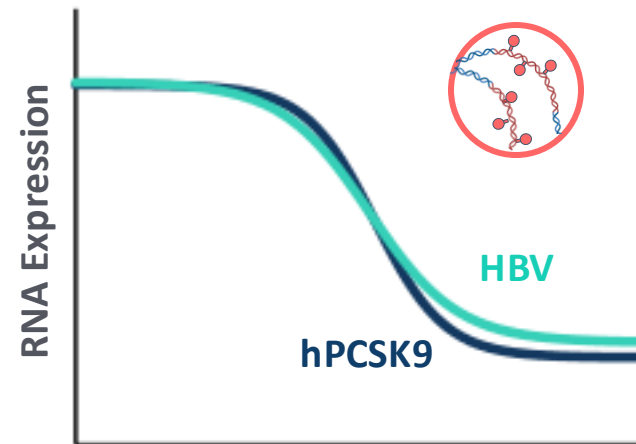
ISH, IHC, and serum PCSK9 repression all support that LNP delivery reaches nearly every hepatocyte in the NHP liver

# Estimation of HBV epi-silencer potency offset using data from the cyno PCSK9 epi-silencer



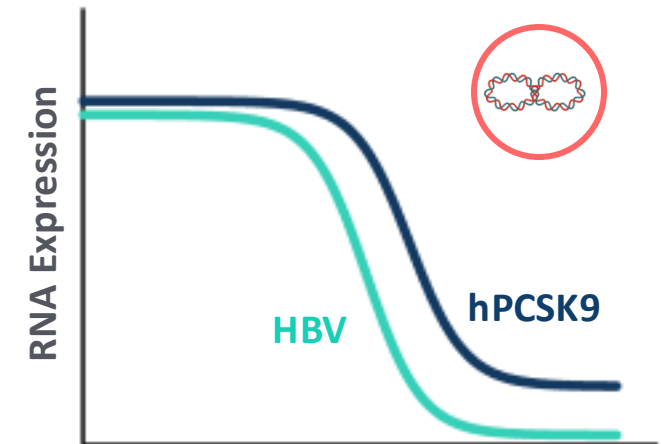
**4.3x** species potency offset

intDNA/Hep3B



**0.9x** potency offset

cccDNA/PHH



**1.5x** potency offset

Total Offset **3.87-fold** ↗

**6.45-fold** ↗

# Tune-401 entering human clinical trials in 2024





THANK YOU

LIPID NANOPARTICLE TECHNOLOGY PROVIDED BY

