

# Translational development of CRMA-1001, an epigenetic silencer for the treatment of chronic hepatitis B

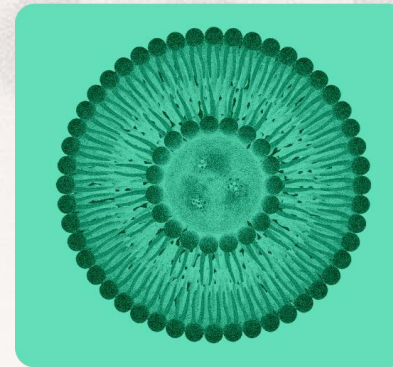
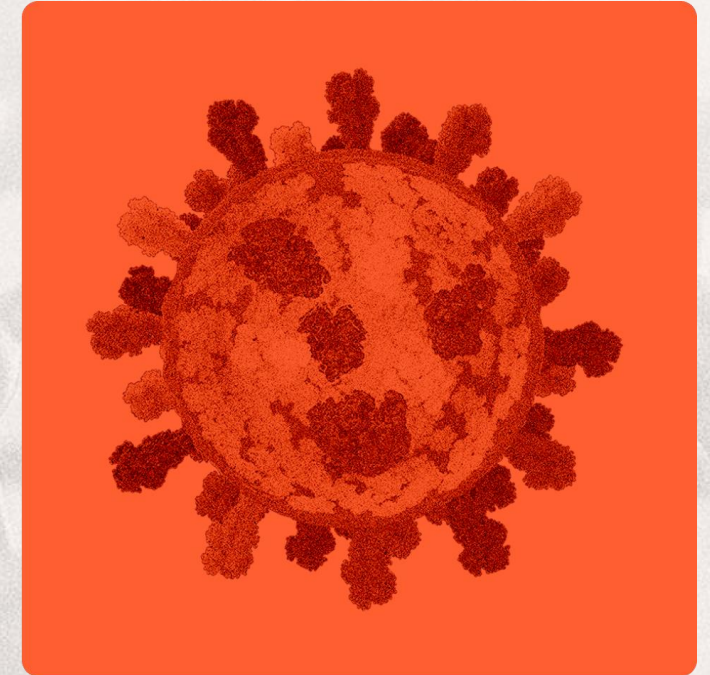
Sarah Voytek, PhD

Vice President, Translational Medicine

Global Program Lead

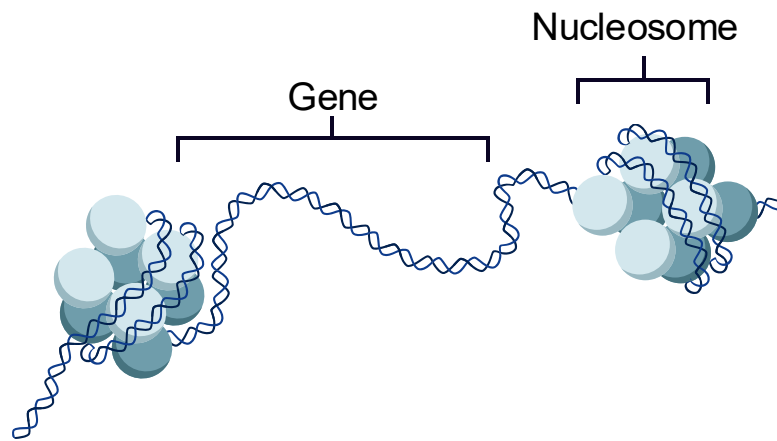
*ASGCT*

*May 12, 2026*



# Epigenetic silencing leverages endogenous mechanisms to precisely control gene expression

Durable regulation of gene expression without cutting or nicking the DNA



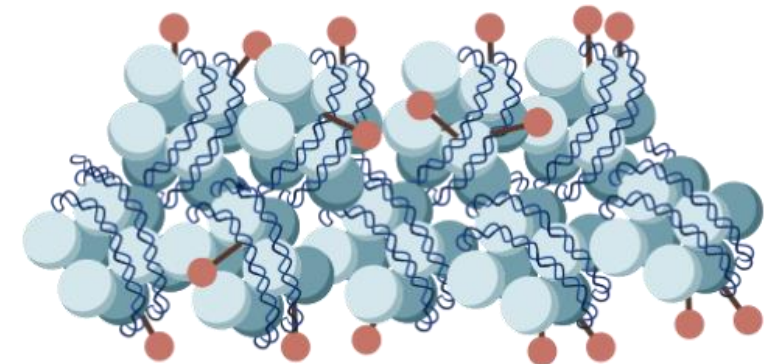
Gene is Active  
DNA is Open and Accessible

Epigenetic Silencer  
Methylates Targets



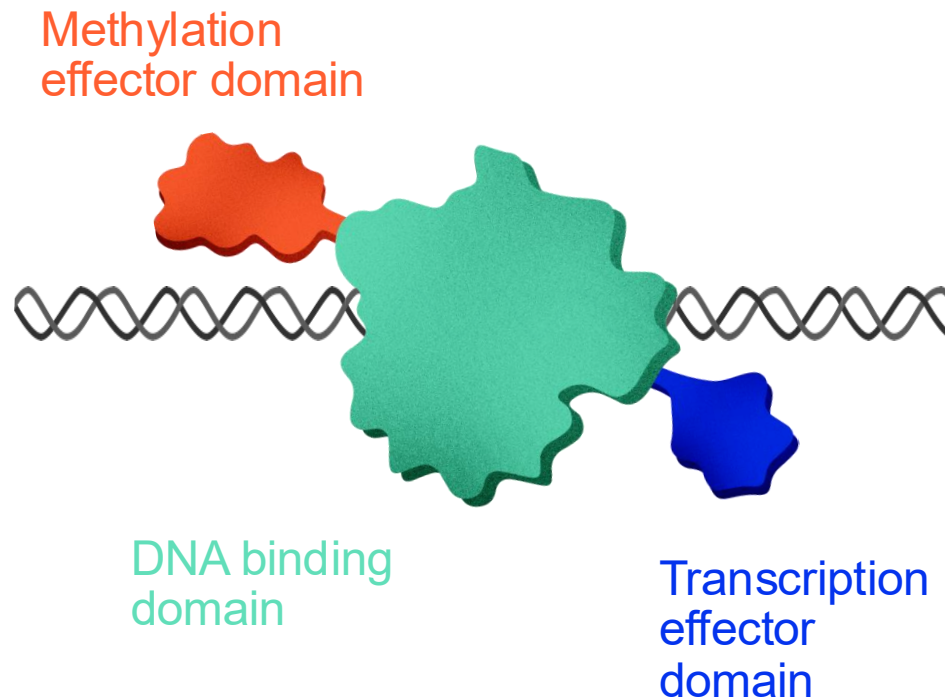
Transient Application

Gene is Inactive  
DNA is Closed and Inaccessible



# Epigenetic silencers are modular constructs that can be delivered to the liver by LNPs

## Epigenetic Silencer Protein

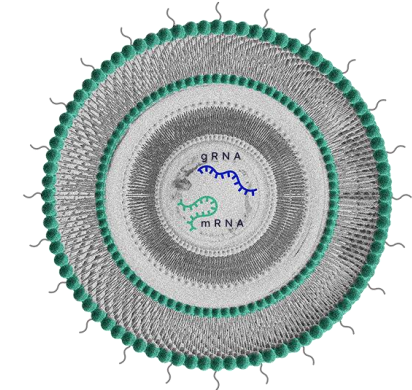


**DNA binding domain** precisely localizes effector domains to target sequence

**Transcription effector domain** transiently suppresses the target gene

**Methylation effector domain** durably represses the target gene

## LNP delivery



mRNA and gRNA delivered via LNP

# Novel therapies needed for chronic HBV to drive functional cure

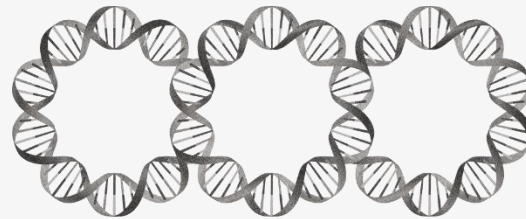
## Key Mechanisms Driving HBV Pathophysiology



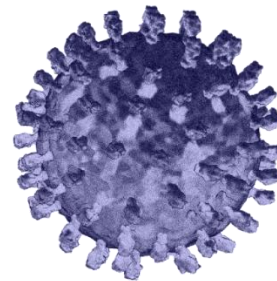
### Current Therapies are Limited

- 254M chronic hepatitis B patients globally
- 1.1M hepatitis B related deaths annually
- Current therapies don't robustly target both mechanisms – very low functional cure rates
- NUC therapies generally require lifelong treatment

## 1 Viral Replication



cccDNA



HBV DNA

HBsAg

## 2 Immunosuppressive Environment



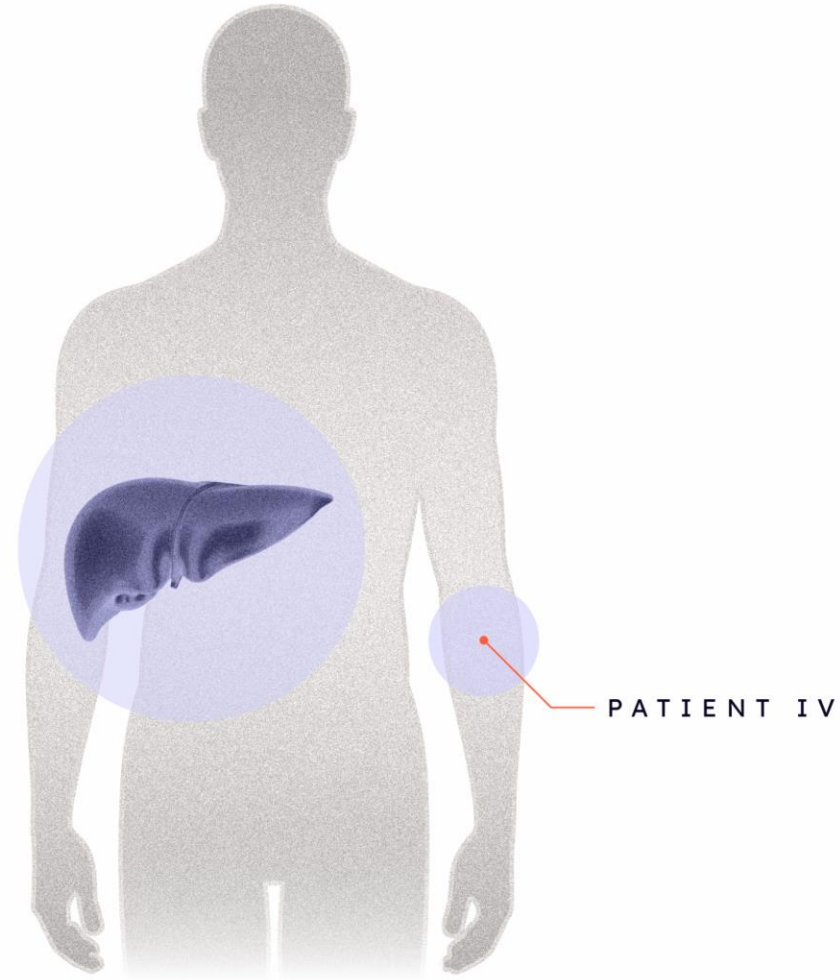
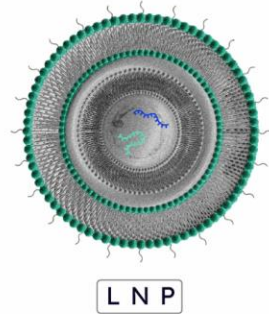
intDNA



HBsAg

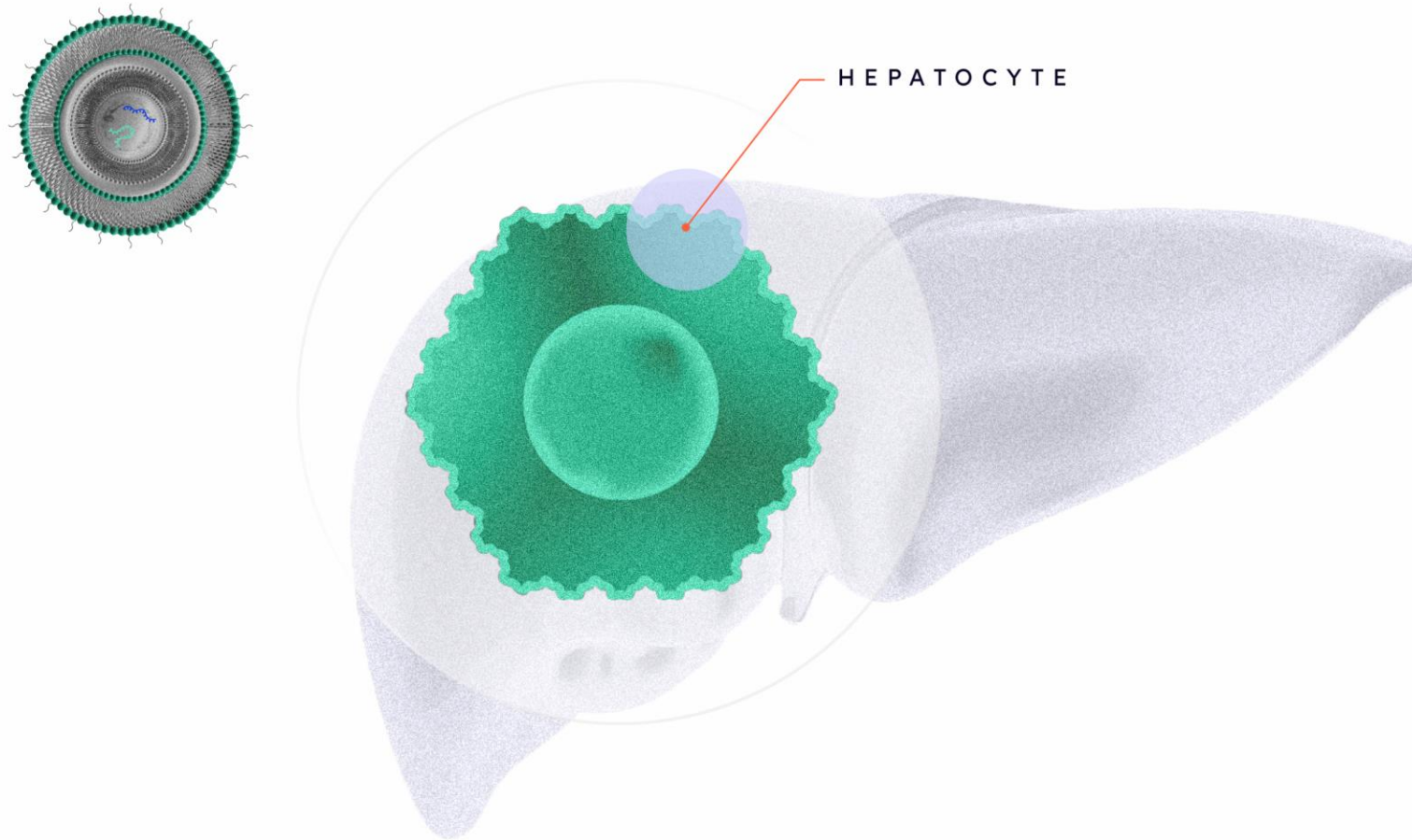
# CRMA-1001 silences both HBV integrated and cccDNA

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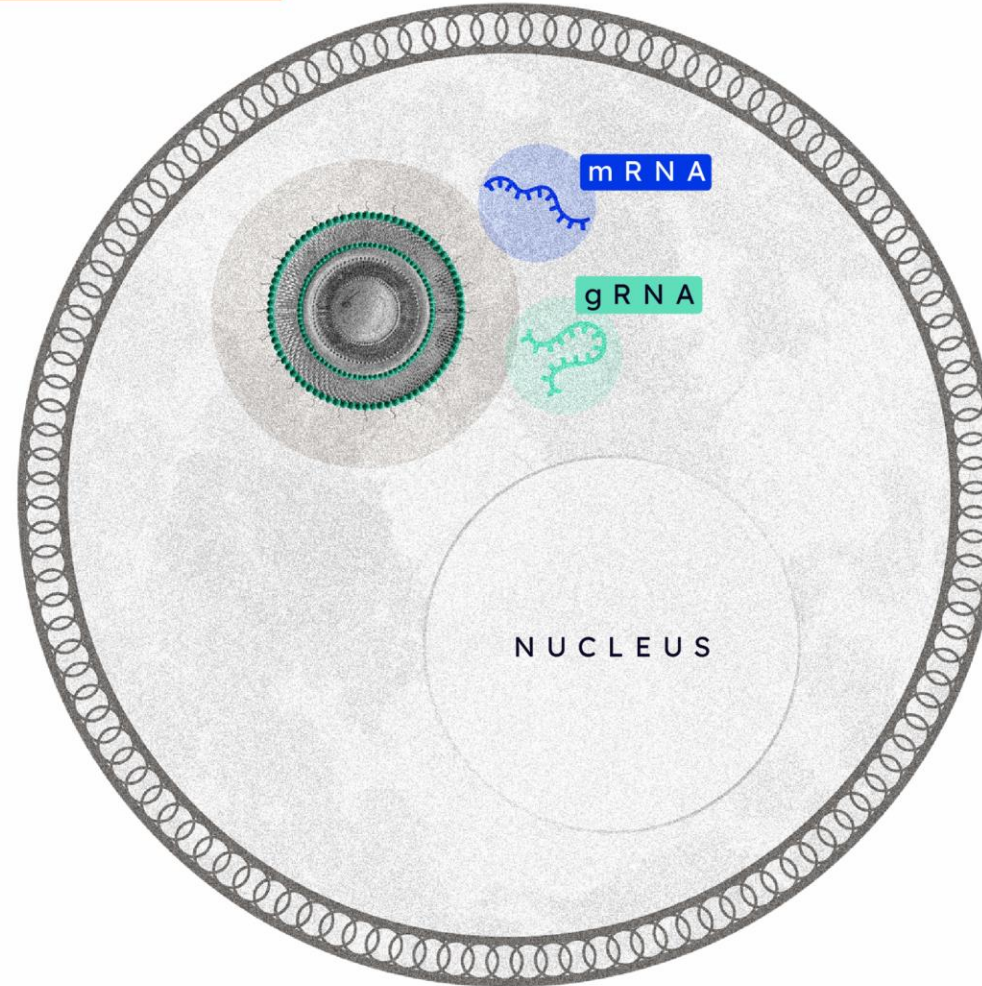
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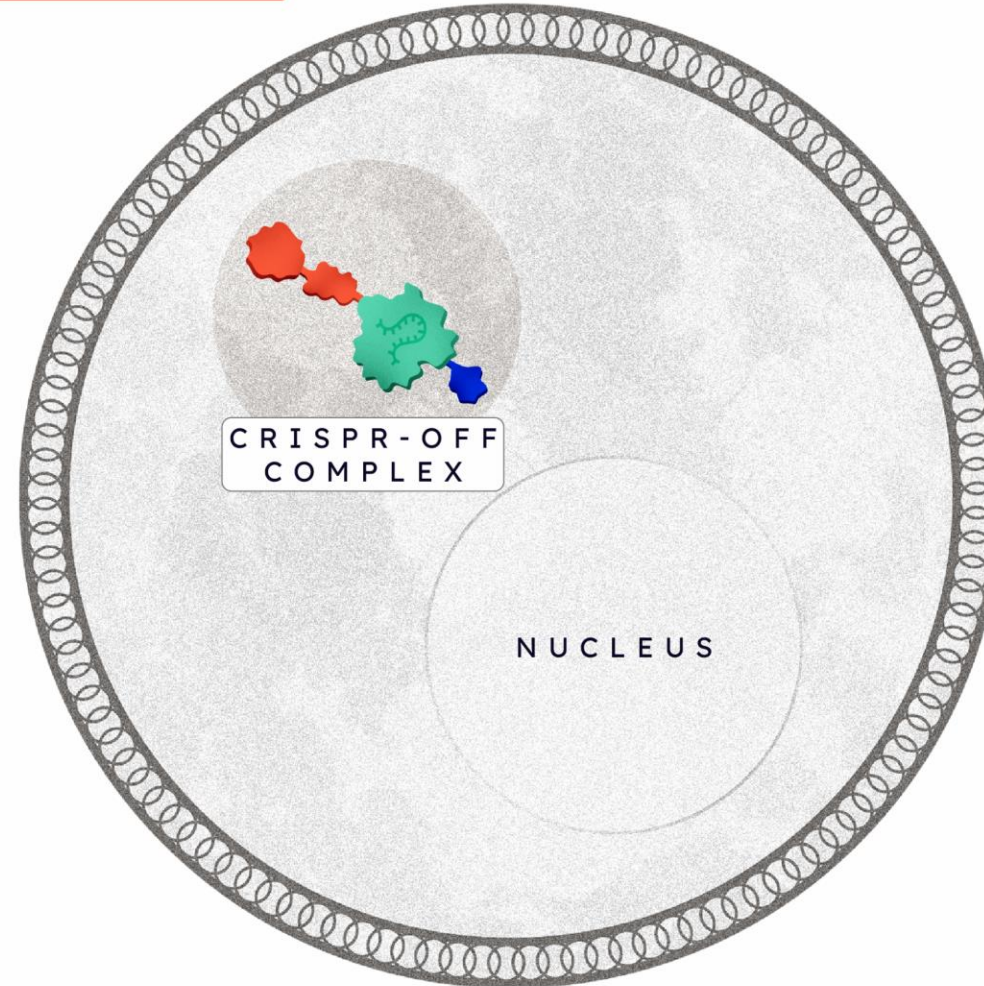
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## RNP COMPLEX FORMATION

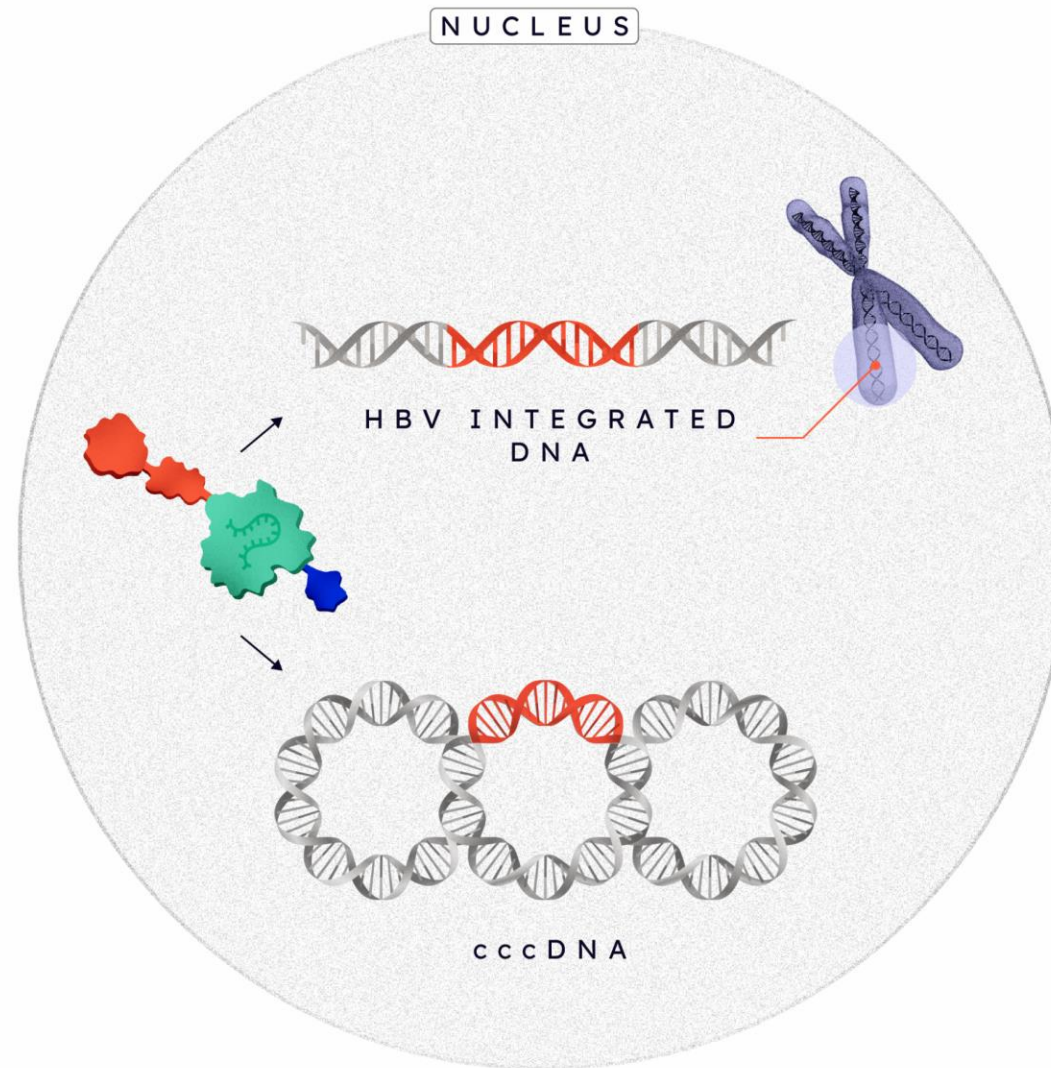


# CRMA-1001 silences both HBV integrated and cccDNA

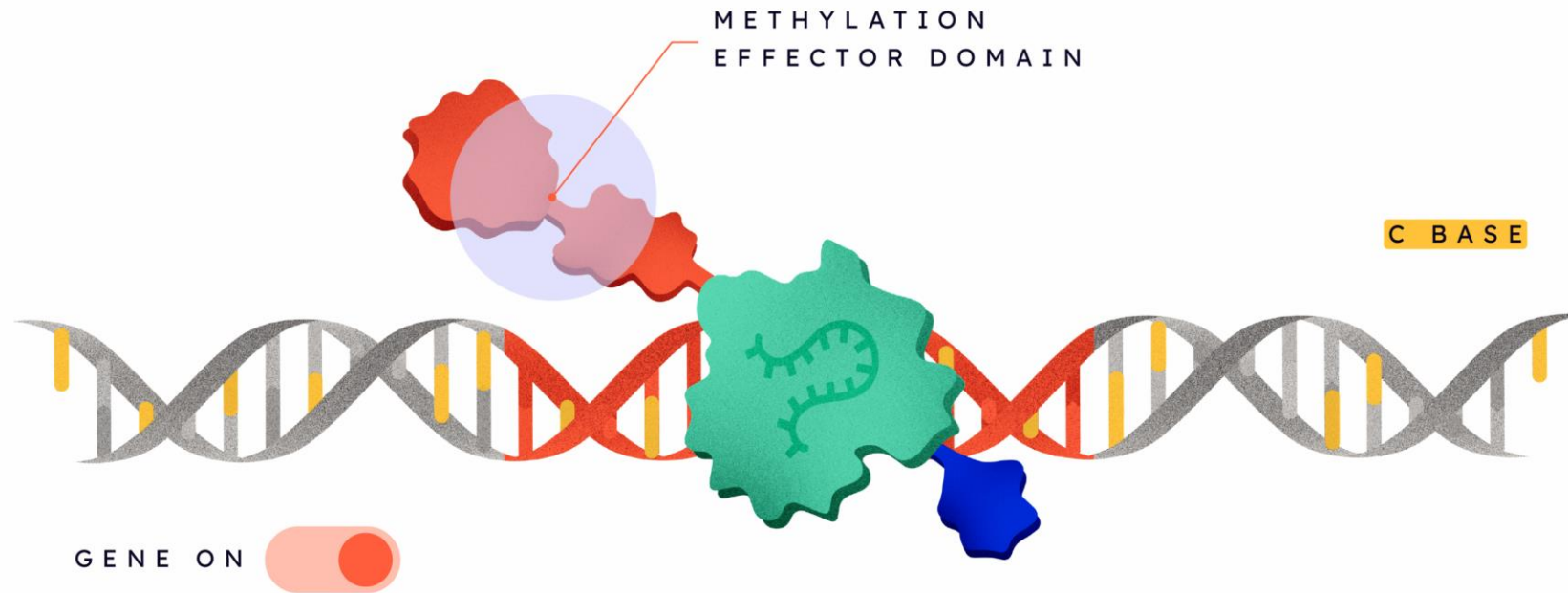
## RNP COMPLEX FORMATION



# CRMA-1001 silences both HBV integrated and cccDNA



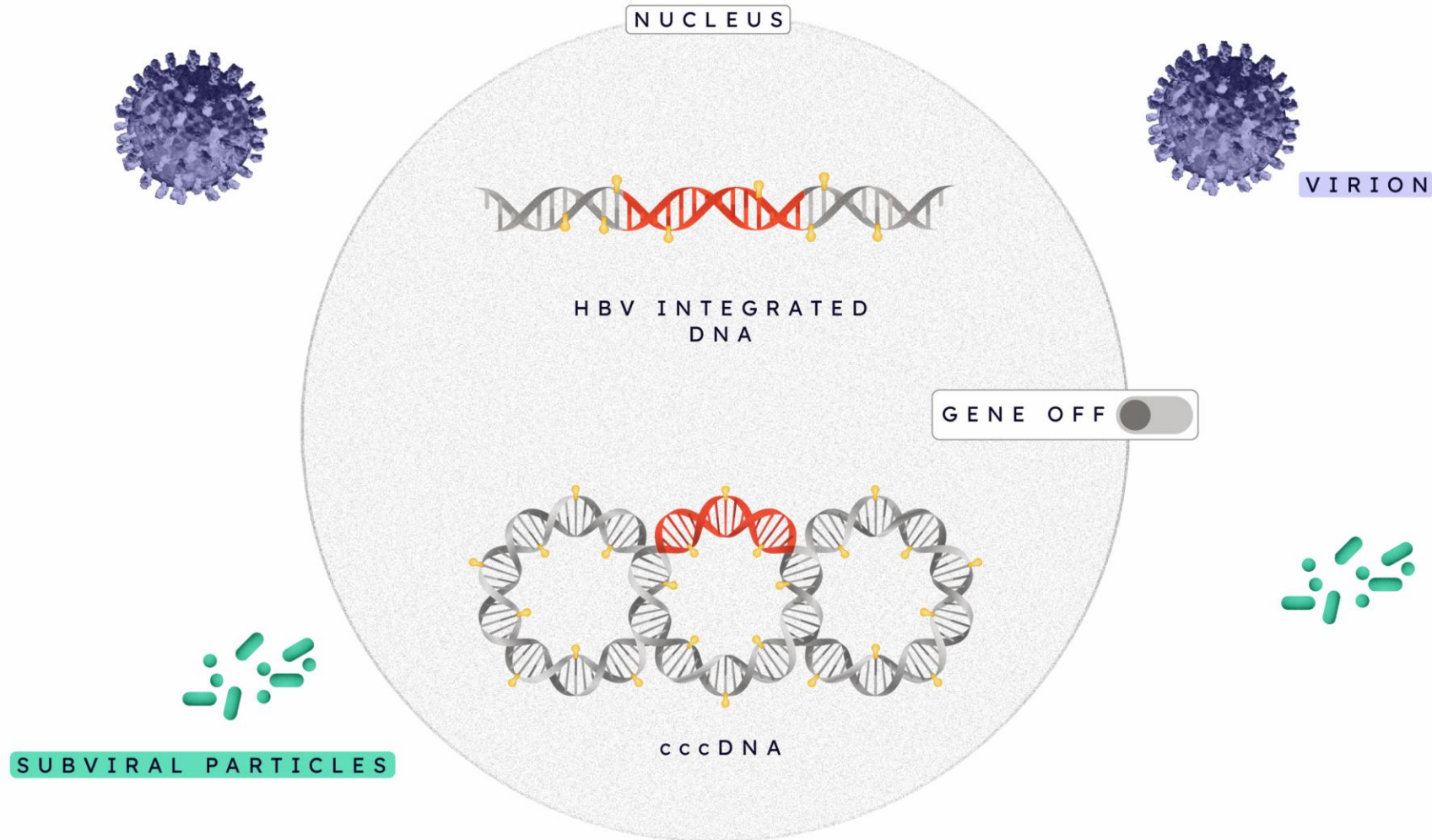
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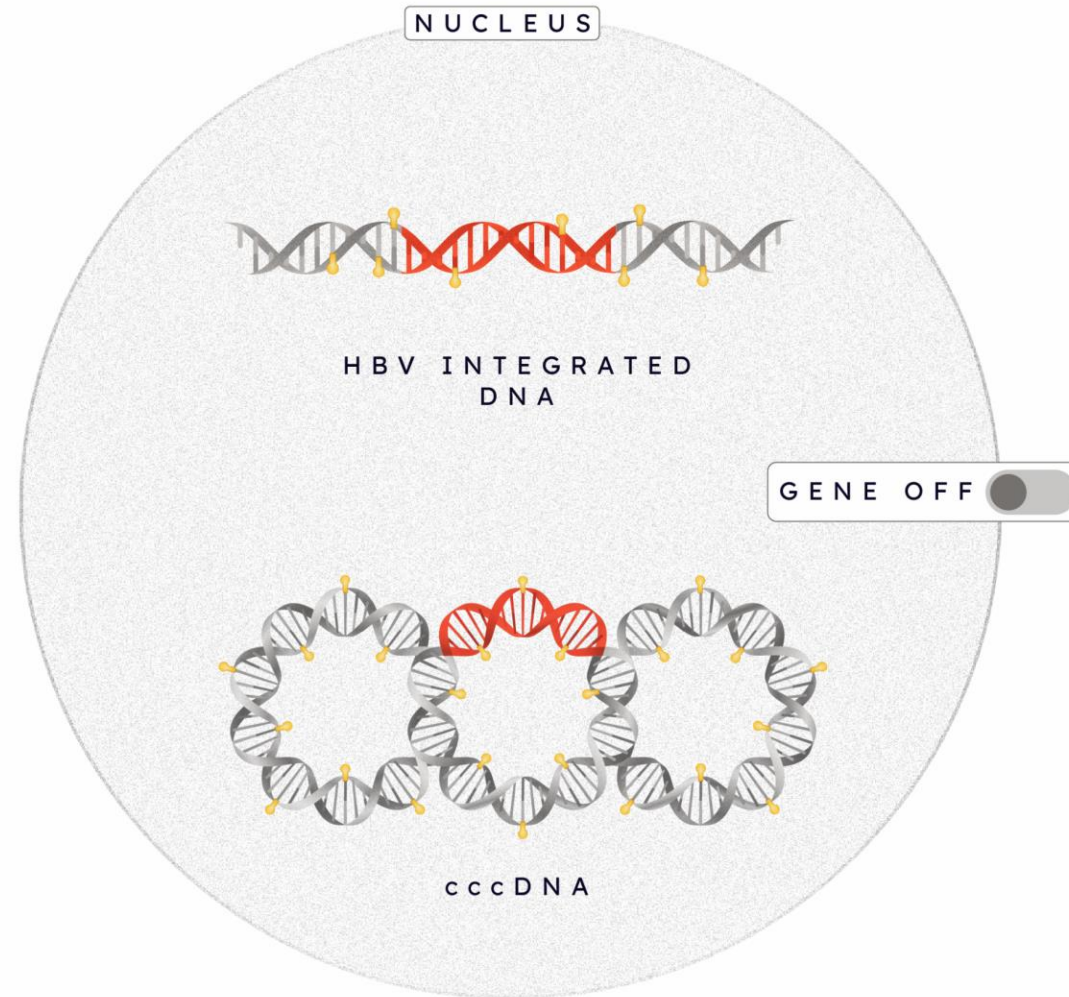
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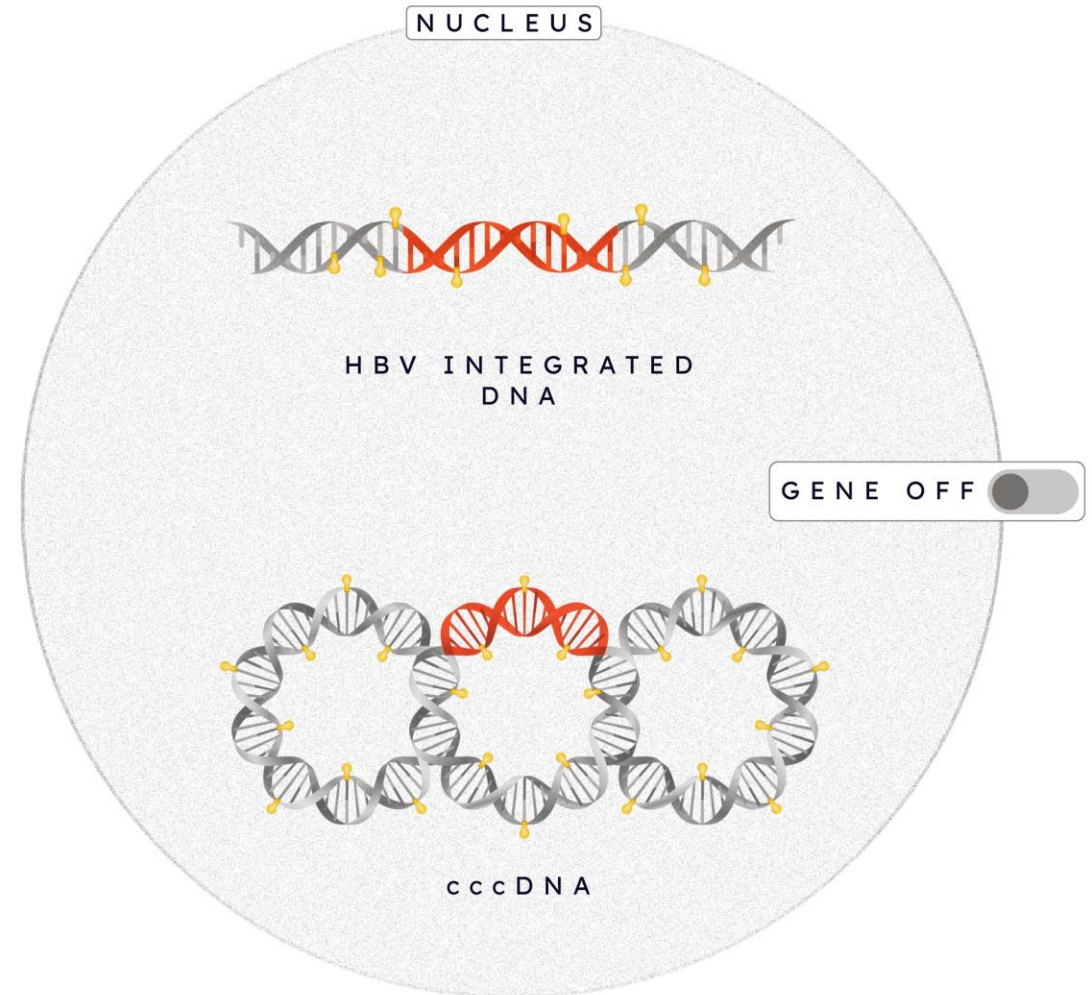


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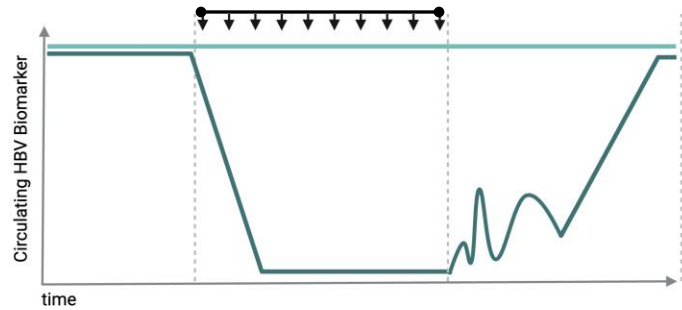
# Epigenetic silencing is uniquely suited to address chronic HBV

- ✓ **Durably silences both forms of the genome**, HBV cccDNA and intDNA, at the level of transcription
- ✓ **Conserved** target site across HBV genotypes (>94% across genotypes A-H)
- ✓ **Highly potent with 99.9% silencing** of viral antigen after a single dose in vivo, supporting delivery to virtually all hepatocytes
- ✓ **Potential for clinically meaningful rates of functional cure with a single course of treatment**
- ✓ **Mechanism avoids unintended cutting or nicking DNA** as compared to other gene editing approaches

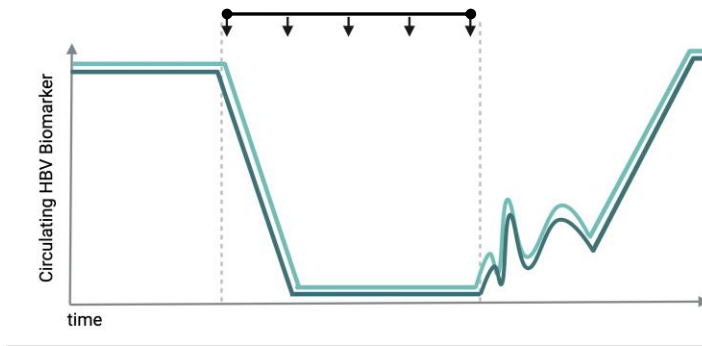


# CRMA-1001 will potentially enable functional cures in chronic HBV where other technologies may be limited

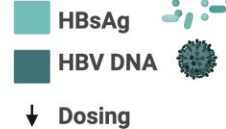
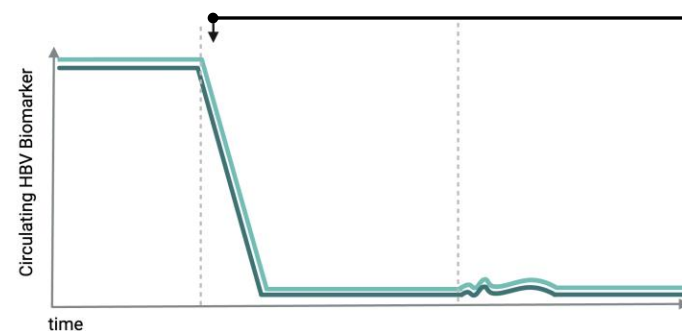
## Nucleos(t)ide Analog



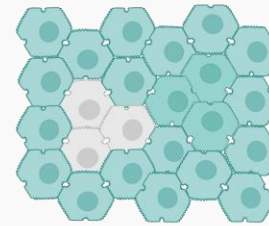
## ASO/siRNA



## CRMA-1001

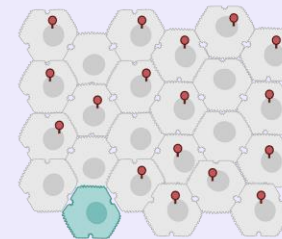


## Nucleos(t)ide Analog and ASO/siRNA



At cessation of treatment, infected cells are **functionally active** producing virions and antigens. Significant disease burden **overwhelms the immune system**, limiting clearance of infected cells and rates of functional cure.

## CRMA-1001

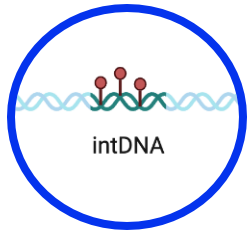


CRMA-1001 treatment is intended to **functionally inactivate** HBV, resulting in **minimal residual disease burden** for immune system to clear, enabling functional cure.

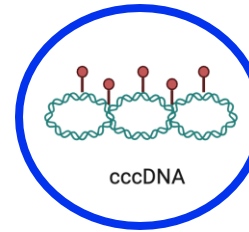
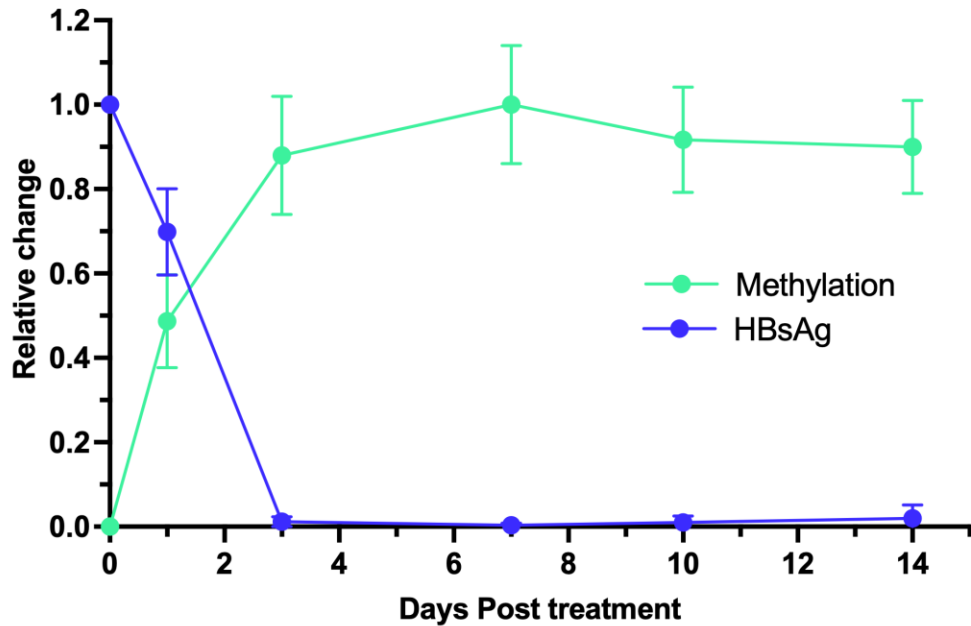


Plots are illustrative

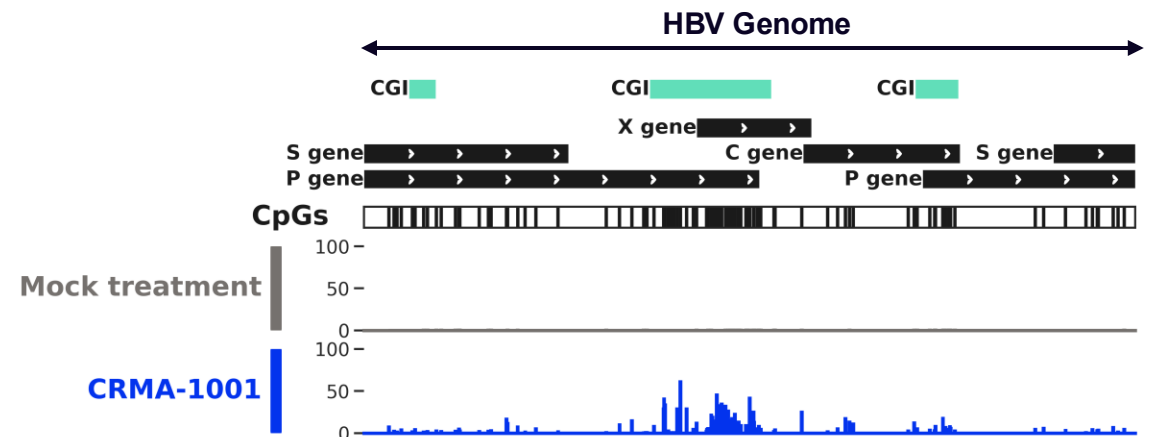
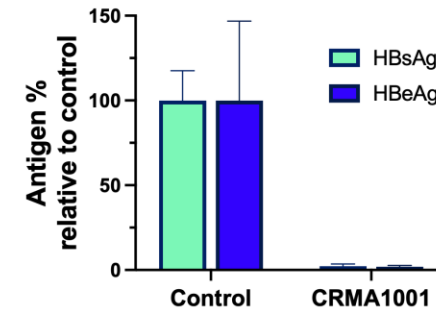
# Increase of HBV CpG methylation in both intDNA and cccDNA correlates with reduction of viral biomarkers



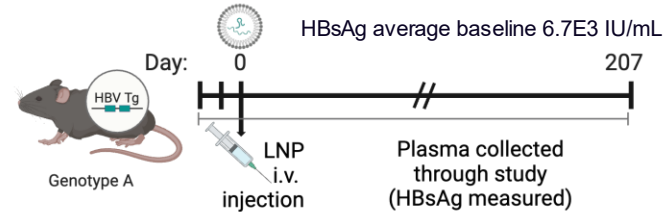
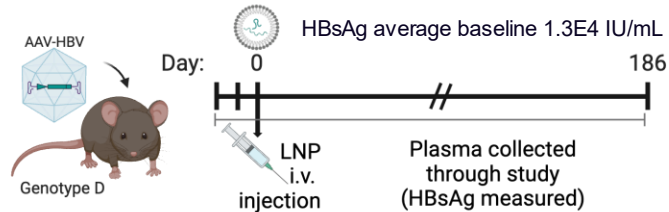
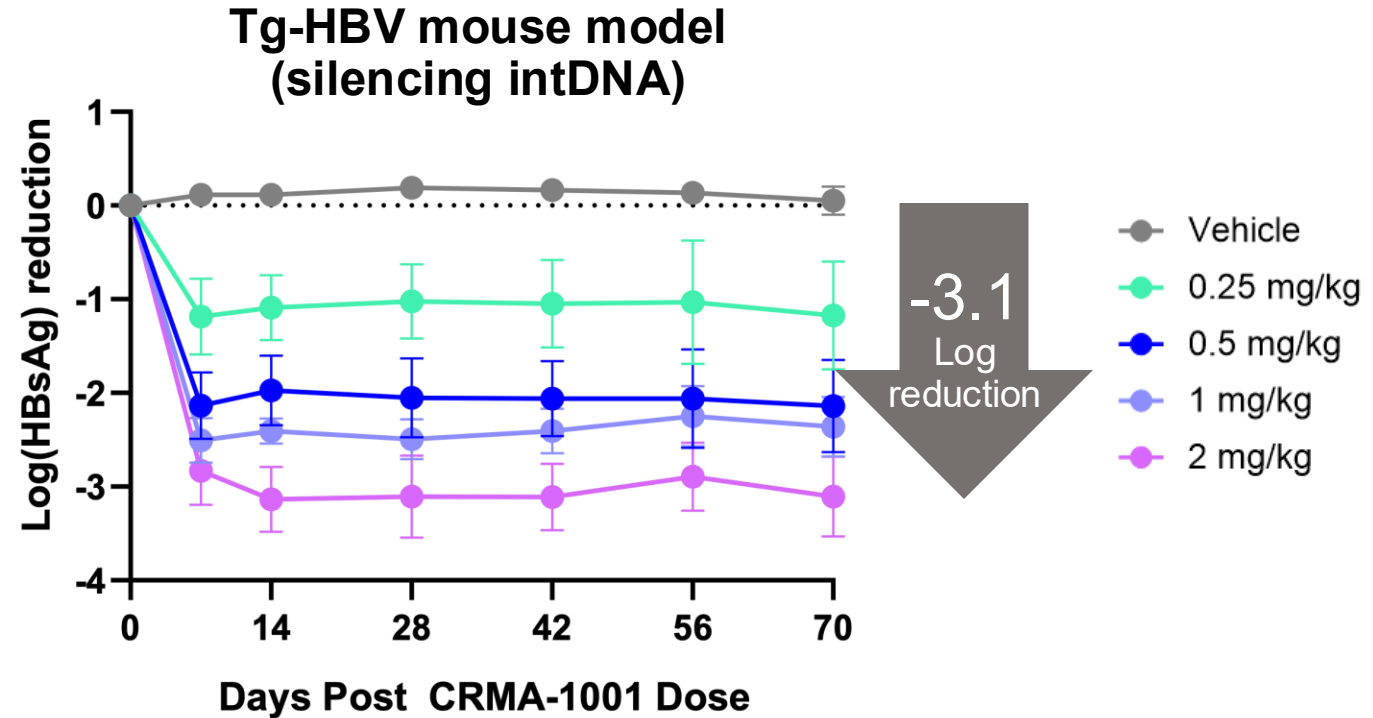
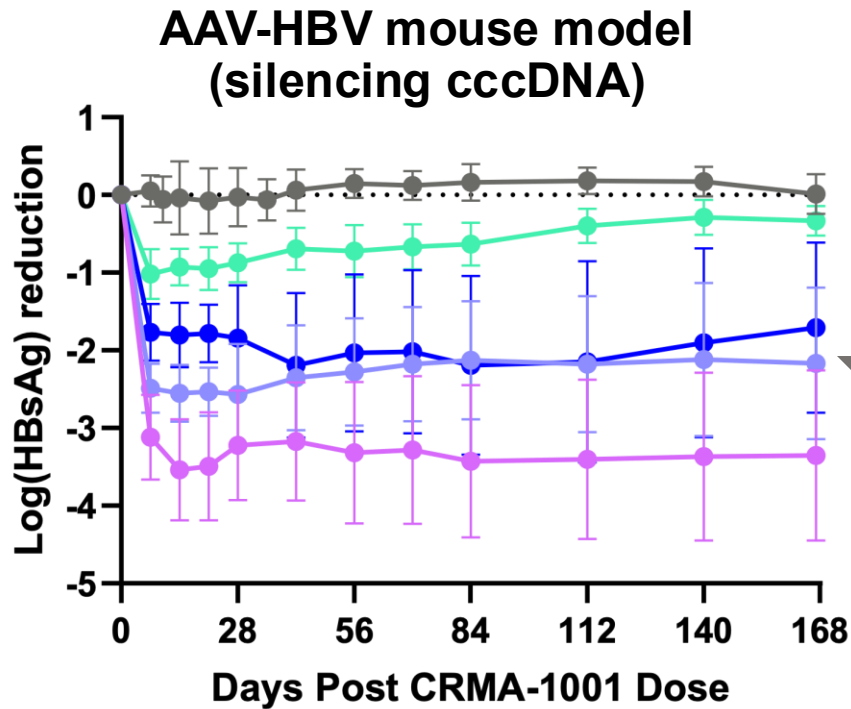
HepG2.2.15 cells 14 days after treatment with CRMA-1001



Infected PHH cells 14 days after treatment with CRMA-1001

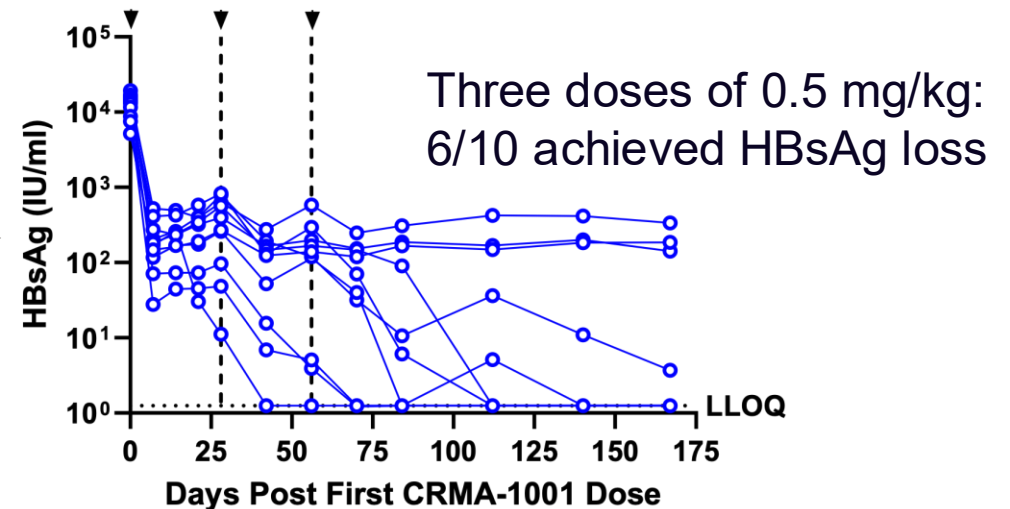
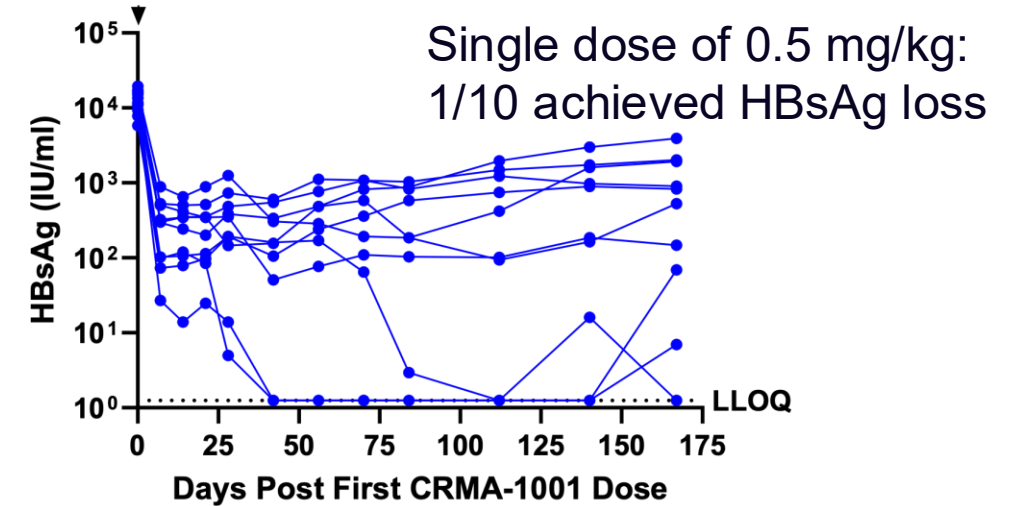
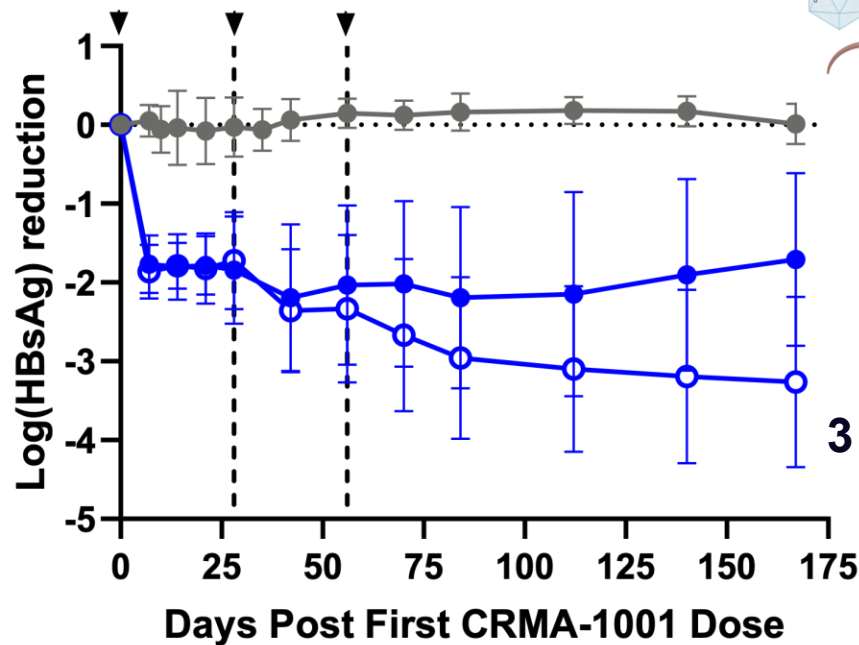


# CRMA-1001 durably reduces HBsAg in HBV mouse models for at least 6 months following a single dose



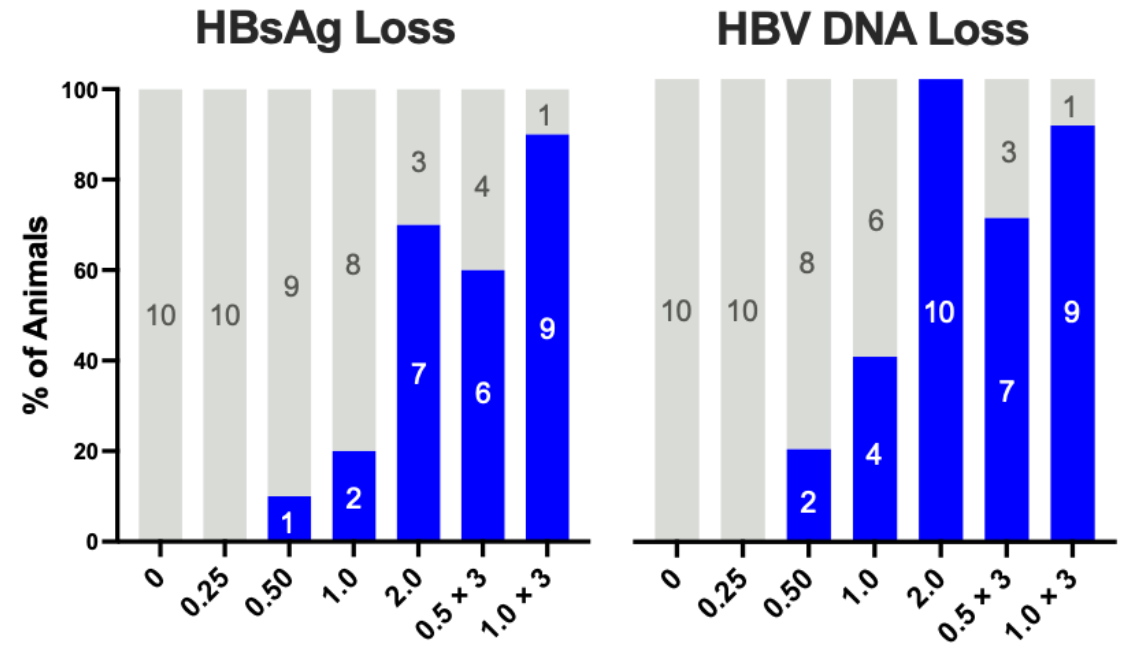
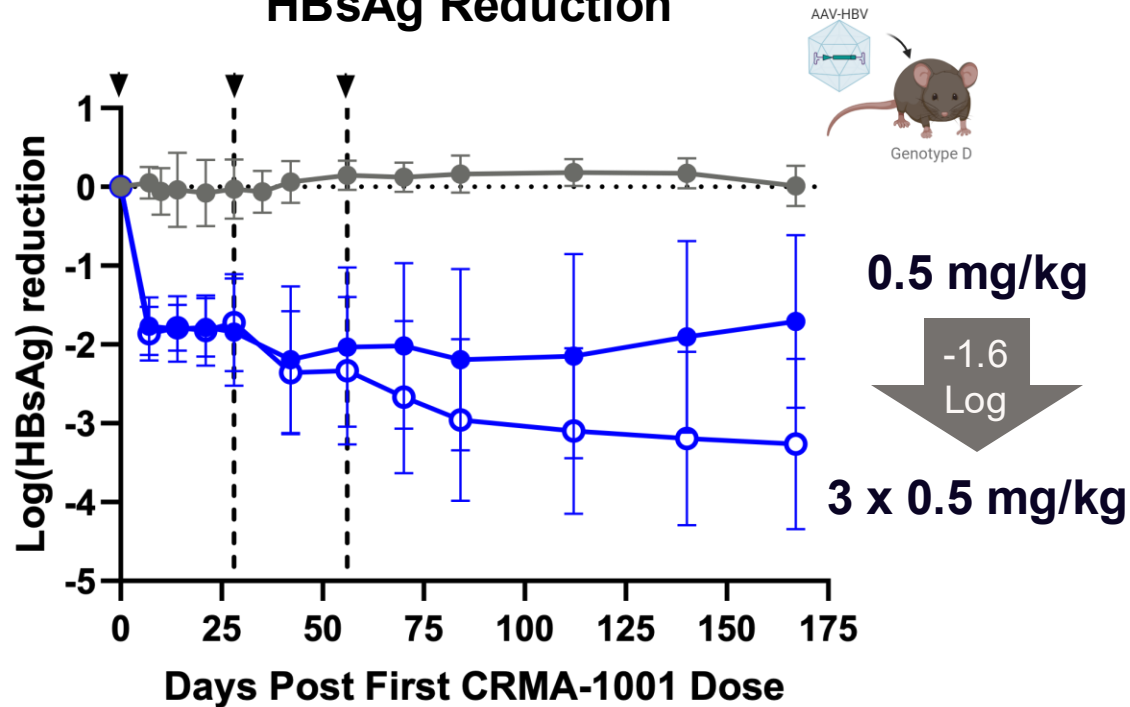
# CRMA-1001 repeat dosing regimen drives additive pharmacology and results in more animals with HBsAg loss

**CRMA-1001 single vs three doses  
HBsAg Reduction**



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**CRMA-1001 single vs three doses  
HBsAg Reduction**

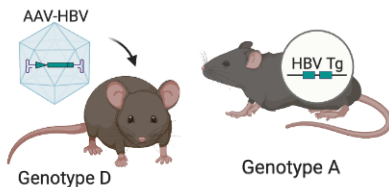
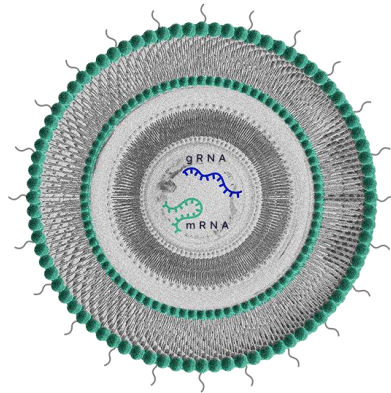


**CRMA-1001 Dosage [mg/kg]**

Number of animals with:  
 ■ Detectable Biomarker  
 ■ Biomarker Loss

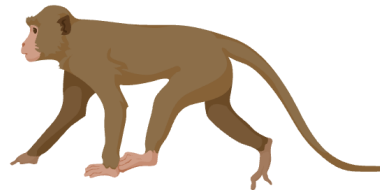
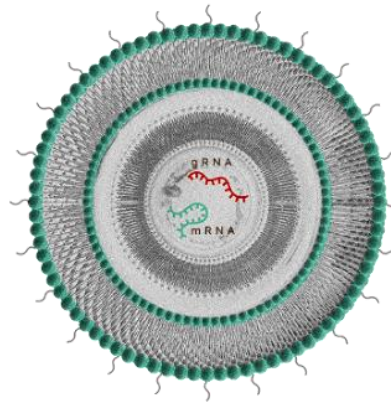
# Epigenetic silencing of PCSK9 in NHPs informs efficiency, durability, and tolerability of silencing HBV with CRMA-1001

## CRMA-1001



**No large animal models available for HBV**

## PCSK9-EE

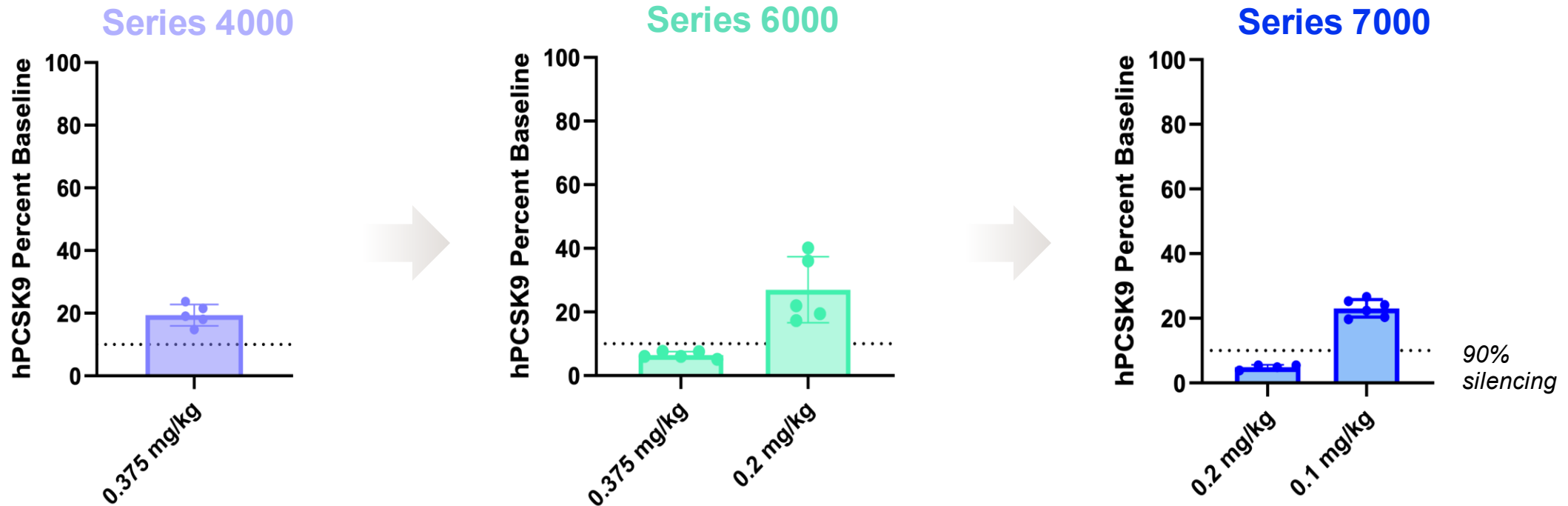
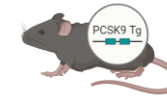


**PCSK9 silencing can be evaluated in NHPs**

- CRMA-1001 and PCSK9-EE use the same mRNA epigenetic effector construct and LNP delivery vehicle to silence the target, only the gRNA is different
- As HBV does not have a large animal model, nChroma Bio is leveraging *PCSK9* to evaluate the efficiency, durability, and tolerability of epigenetically silencing HBV

# Construct optimization efforts have led to a significant increase in potency in animal models

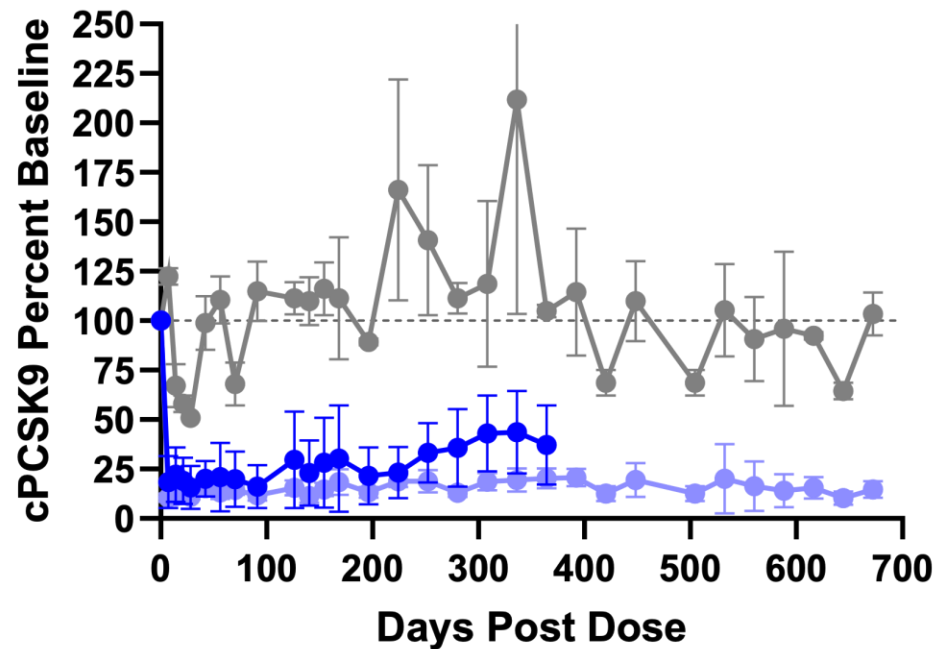
## Optimization in hPCSK9 Tg Mice



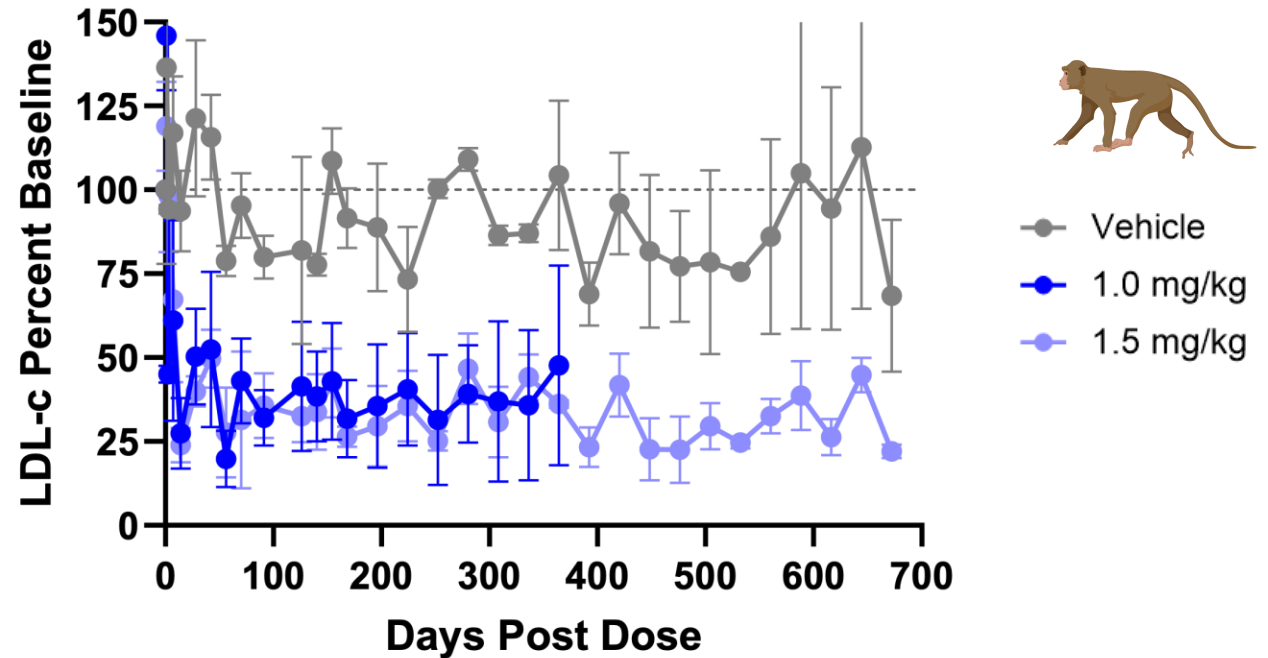
- Construct optimization has yielded a significant increase in epigenetic silencer potency in hPCSK9 Tg-mice
- NHP study with 7000 construct (used in CRMA-1001) demonstrated potency gains and >22 months durability

# PCSK9-EE (Series 7000) achieved saturating pharmacology at $\geq 1$ mg/kg in NHPs with durability beyond 22 months

### Durability of cPCSK9 silencing

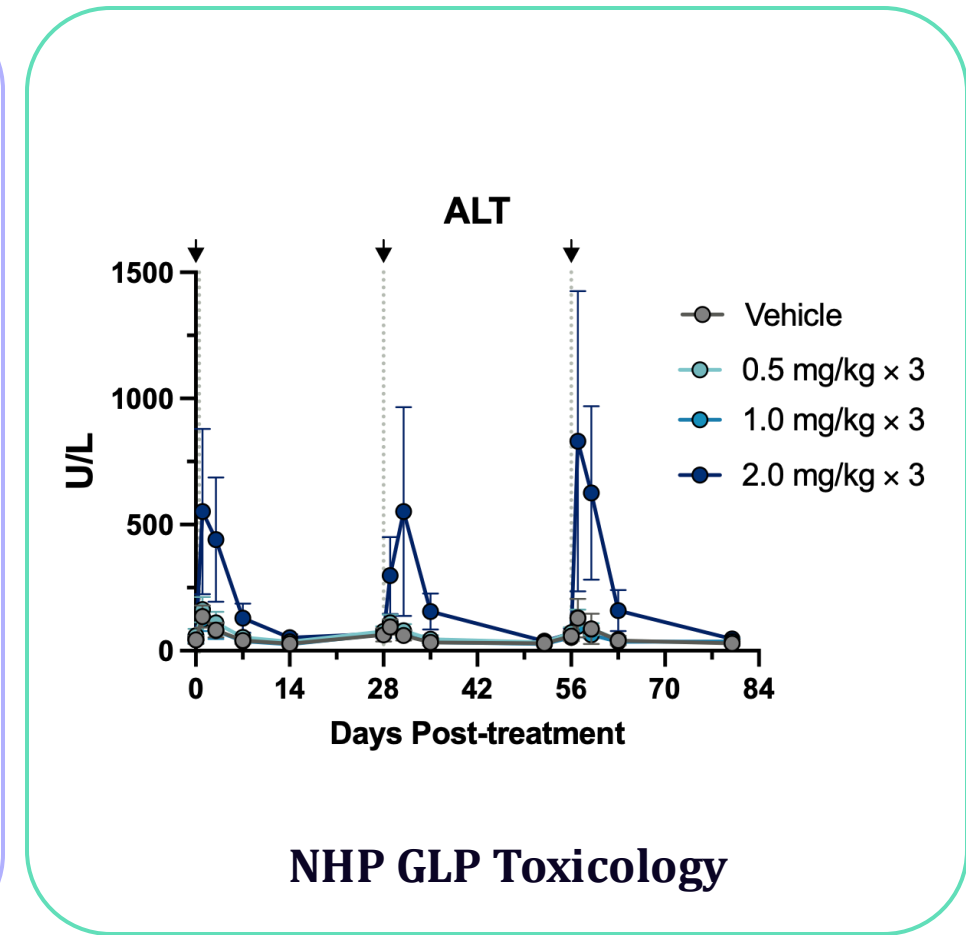
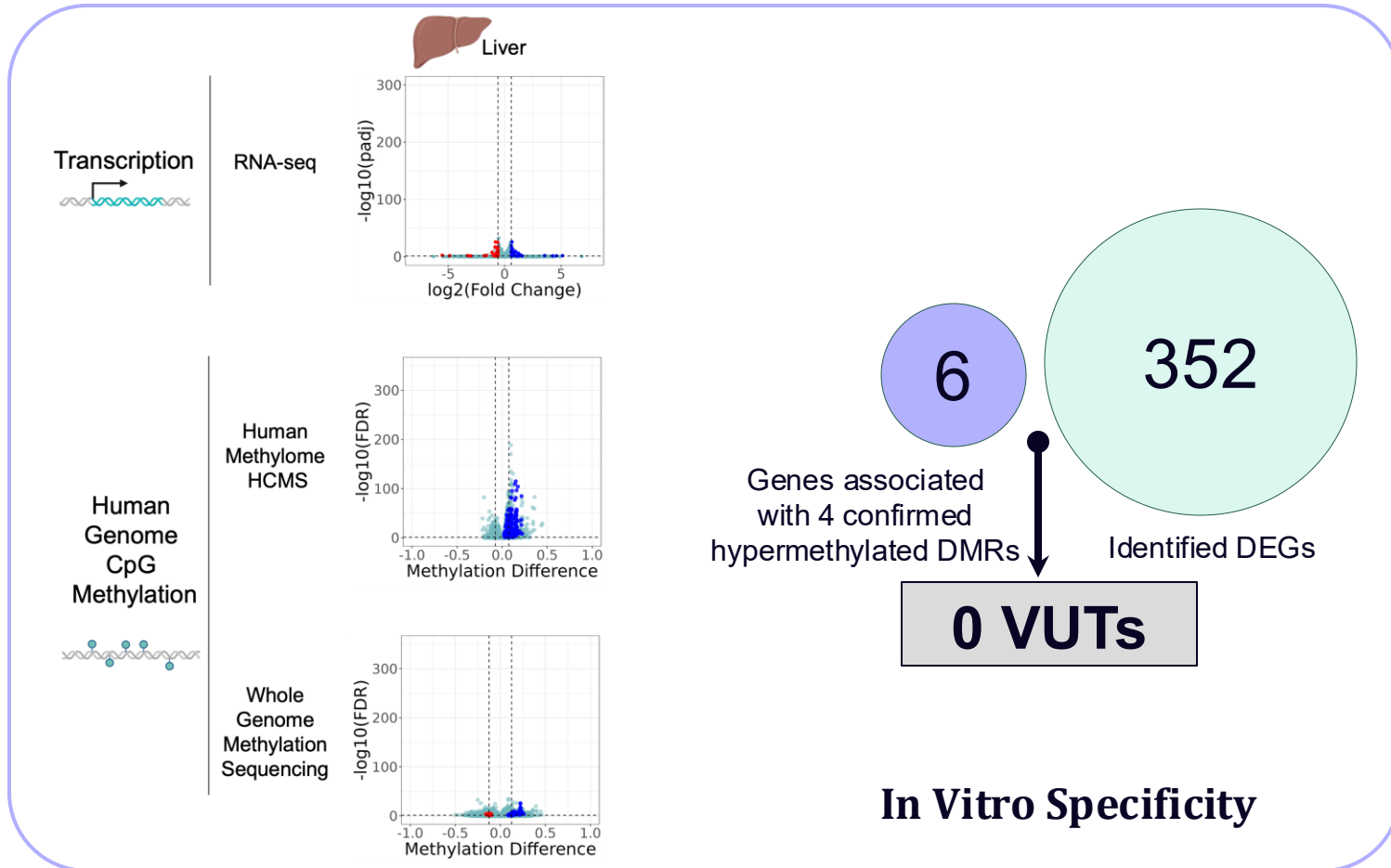


### LDL-c Reduction

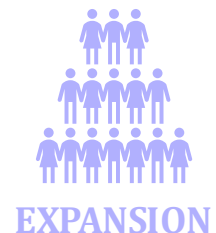


# No verified unintended targets (VUTs) have been identified in the human genome

A comprehensive evaluation of CRMA-1001 specificity, and safety in NHPs supported clinical trial initiation



# CRMA-1001-101: A Phase 1/2, open-label, SAD/MAD study of CRMA-1001 in adults with chronic hepatitis B



## Key Eligibility Criteria

### Inclusion Criteria:

- Adults with chronic hepatitis B
- On oral antiviral therapy
- ALT and AST  $\leq 1.5 \times$  ULN
- Total bilirubin  $\leq$  ULN

### Exclusion Criteria:

- Significant hepatic fibrosis or cirrhosis
- Current or prior liver disease other than HBV

## Select Endpoints

### Primary

- Safety and tolerability of single and multiple doses of CRMA-1001

### Secondary

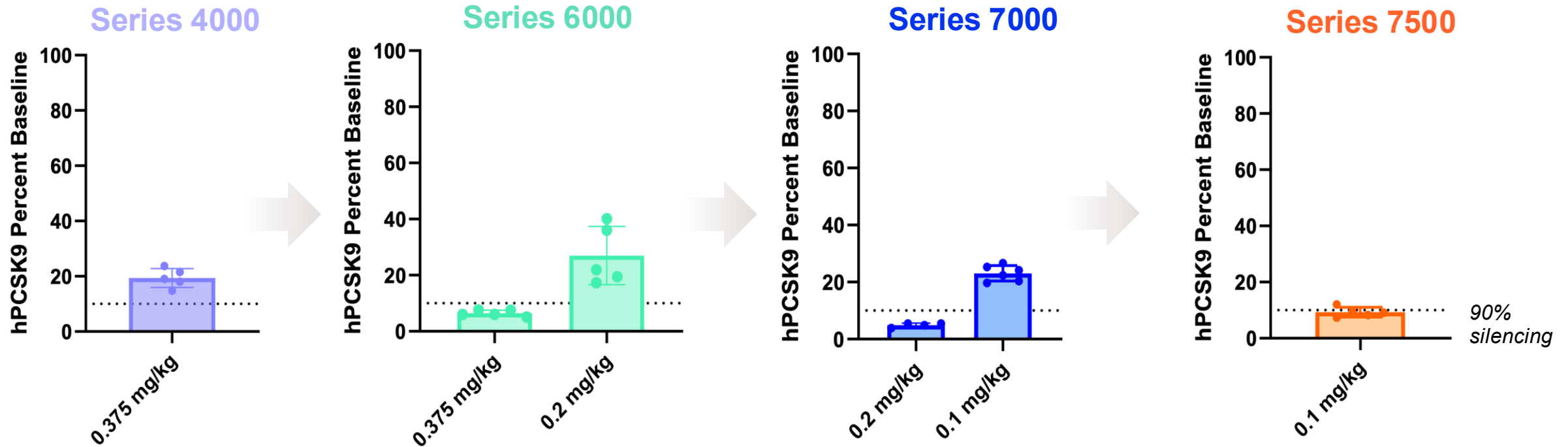
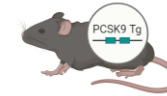
- Pharmacokinetics of CRMA-1001
- Immunogenicity of CRMA-1001
- Change in HBV Biomarkers

The study is actively recruiting and dosing participants in Hong Kong and New Zealand. Additional sites are planned with CTA approvals in the United Kingdom and France.

ClinicalTrials.gov ID: [NCT07200193](https://clinicaltrials.gov/ct2/show/study/NCT07200193) EUCT number: [2025-523619-12-00](https://eudra-ct.europa.eu/eudra-ct/index.jsp?view=detail&tab=summary&id_trial=2025-523619-12-00)

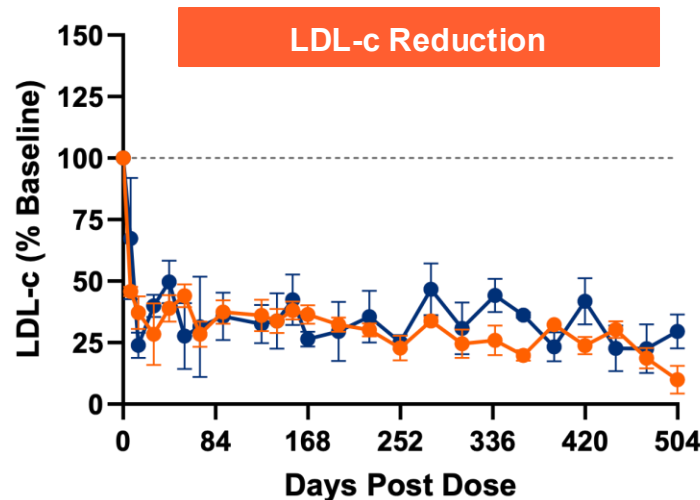
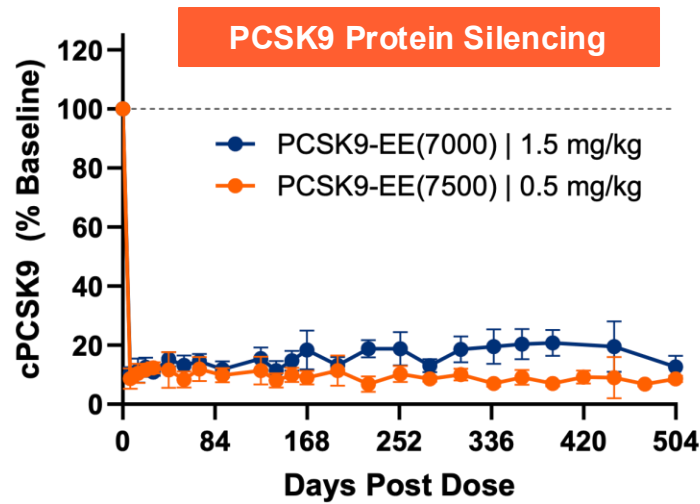
# Ongoing construct optimization efforts demonstrate significant increases in potency in animal models

## Optimization in hPCSK9 Tg Mice



- Further construct optimization yields increased potency in epigenetic silencer potency in hPCSK9 Tg-mice
- NHP study with Series 7500 construct has demonstrated additional potency gains and >18 months durability

# PCSK9-EE (Series 7500) achieved >3X improved potency over 7000 series editor in NHPs with durability beyond 18 months



# CRMA-1001: Designed to durably silence cccDNA and intDNA with the potential to achieve high rates of functional cure

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- ✓ **Potency:** >3 log reduction of HBsAg at a single high dose or with repeat lower doses
- ✓ **Durability:** Suppressed HBV viral markers for over 6 months (observation period of study)
- ✓ **Translation:** Effector saturated *PCSK9* silencing at  $\geq 1$  mg/kg in NHPs for up to 22 months
- ✓ **Safety:** Confirmed methylation at target site with no detectable unintended targets in the human genome, PK/Biodistribution & Toxicology profile supported further development
- ✓ **Pipeline:** Ongoing construct optimization has demonstrated further gains in potency

**CRMA-1001-101 is actively recruiting and dosing participants in multiple jurisdictions**

# Thank you to nChroma Bio team, collaborators, partners, and study participants



# Thank You

