

# Future Therapies for HBV

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# What Would HBV Elimination Look Like?

**In the blood:** HBV DNA/HBsAg negative  
anti-HBs positive

**In the liver:** no HBV cccDNA  
no HBV RC/DSL DNA  
HBcAg staining negative  
± HBsAg (occasional)\*

\*[reflecting integrated HBV DNA]

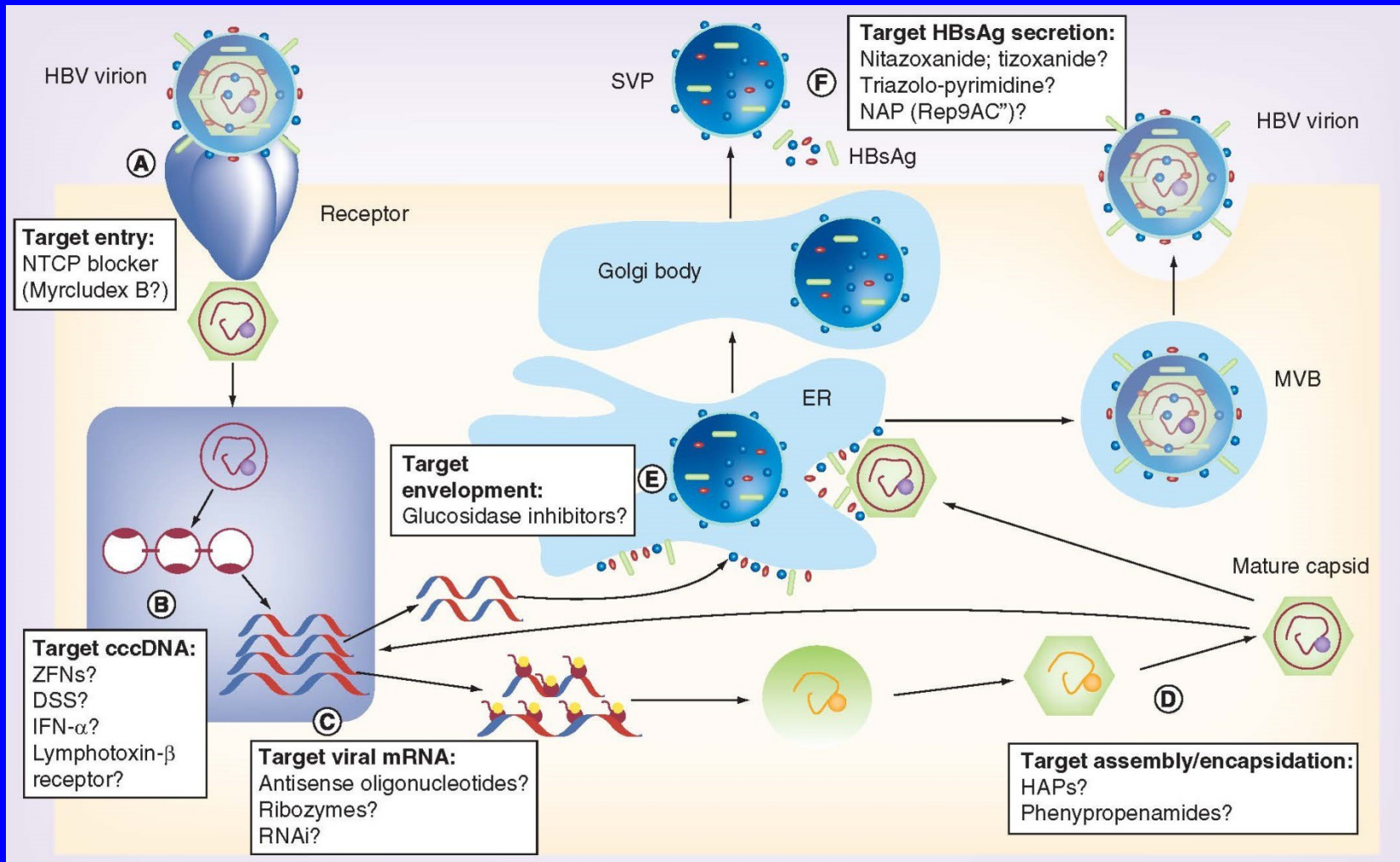
**Functional Cure:** HBsAg loss/Seroconversion: Maintenance of undetectable serum HBV DNA off-treatment

**Absolute or Complete Cure:** No cccDNA or HBV DNA anywhere!

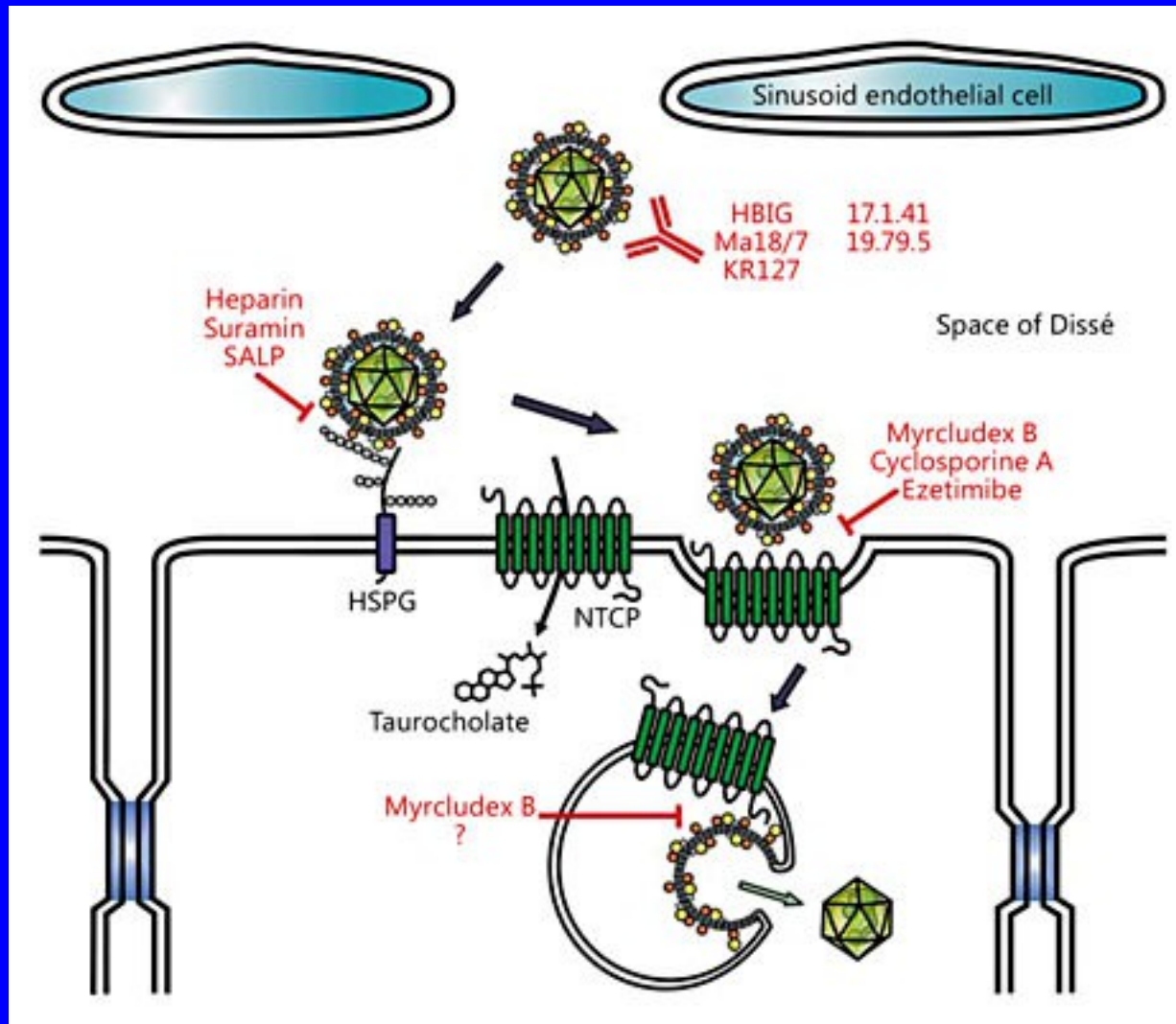
# New Viral Targets

- Attachment and Entry
- cccDNA Generation & Processing (HBcAg and HBx)
- Reverse Transcription
- HBV Nucleocapsid Assembly (HBcAg)
- Packaging Inhibitors
- Molecular Based Therapies (RNAi)
- Combination Therapy

# HBV Lifecycle Showing Novel Approaches for Viral Targets



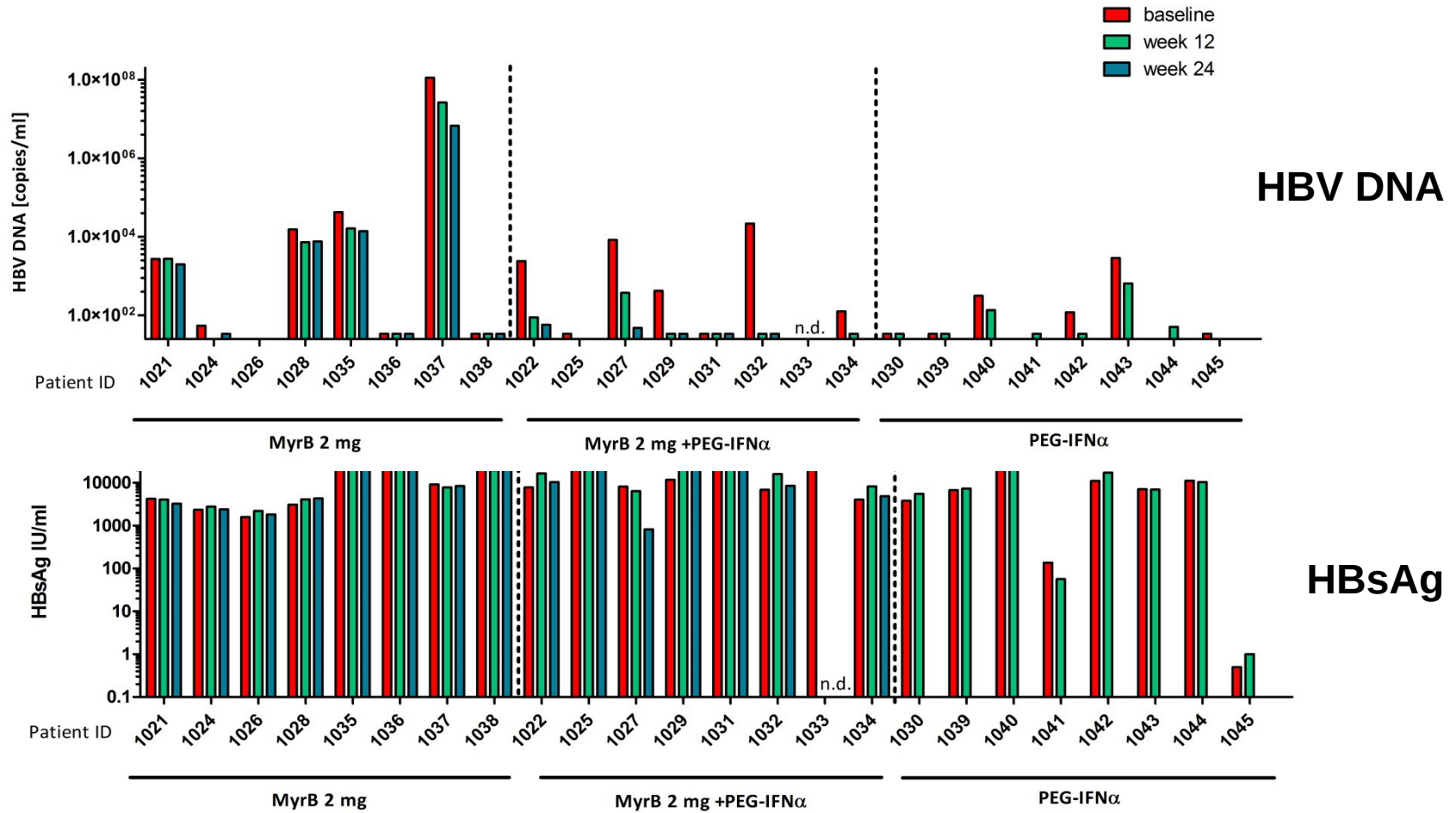
# Inhibitors of HBV Attachment and Entry



Sodium taurocholate cotransporting polypeptide (NTCP) identified as HBV and HDV receptor in 2012

Myrcludex in phase 2 trials in chronic HBV and chronic HDV decrease in HBV DNA and HDV RNA

# (B) HBV Serum DNA- and HBsAg Levels During Myr B and Myr B/IFN $\alpha$ Treatment



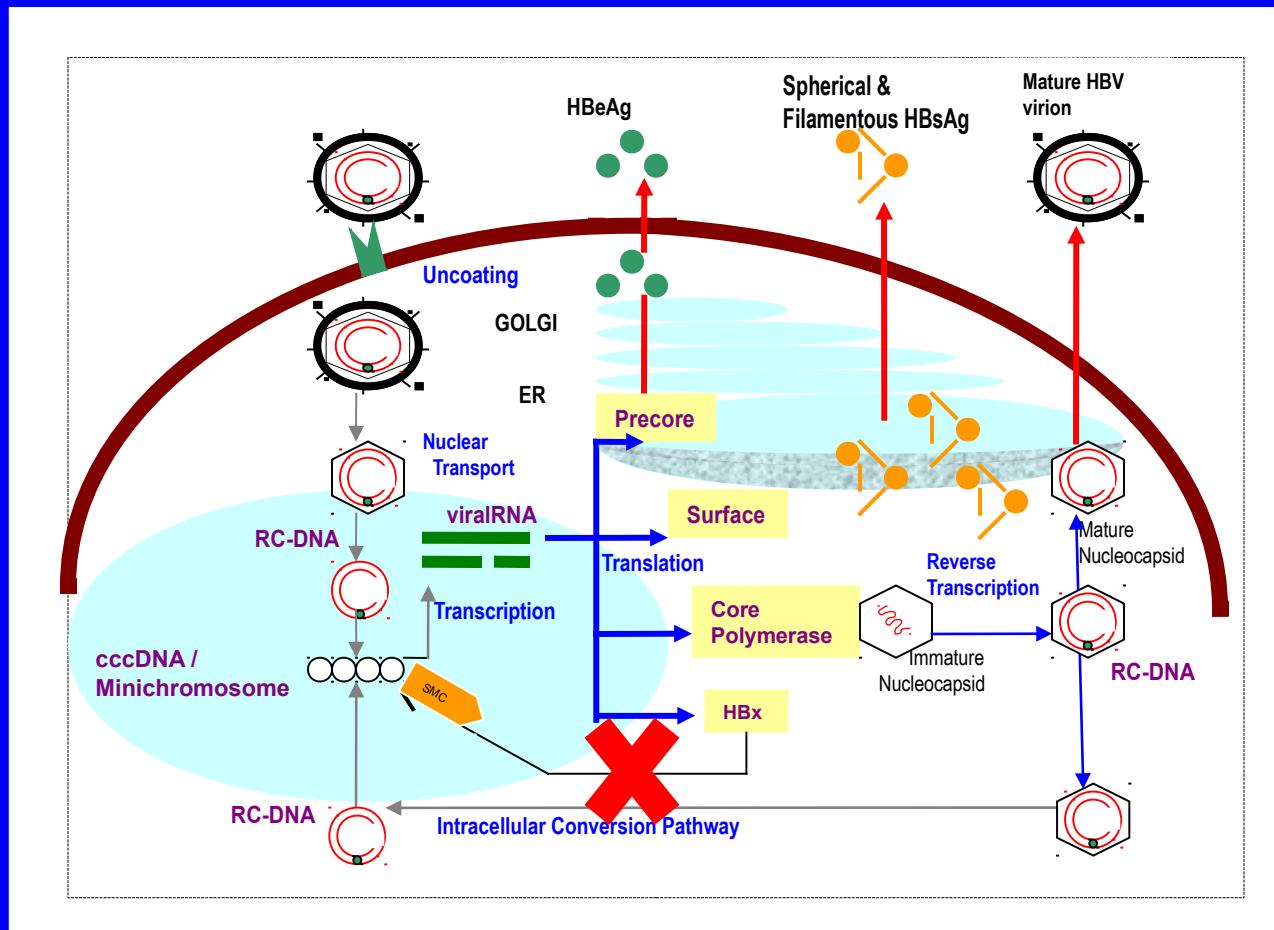
⇒ HBV DNA levels decline during Myrcludex B treatment in 4/8 patients (consistent with HBV trial).

⇒ More pronounced decline of HBV DNA in the Myrcludex B/PEG-IFN $\alpha$  group (5/8 patients).

⇒ No significant changes (except patient 1027) in HBsAg levels .



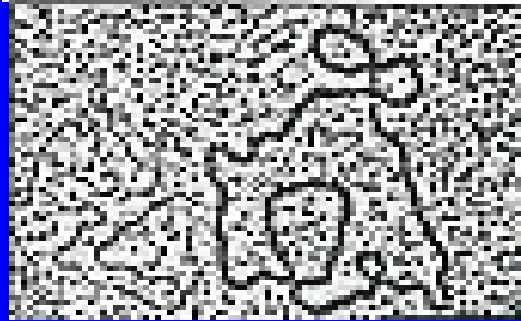
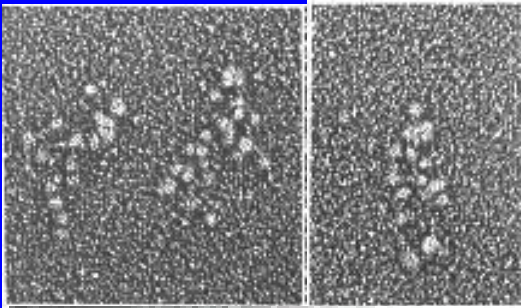
# HBV Replication: cccDNA Pathway



1. RC DNA  $\rightarrow$  cccDNA
  - DNA repair
  - TDP-2

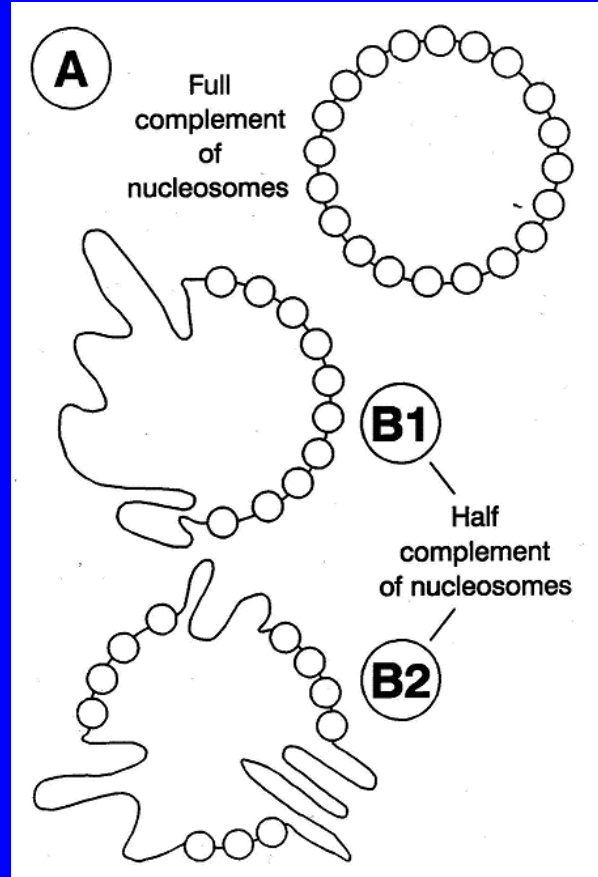
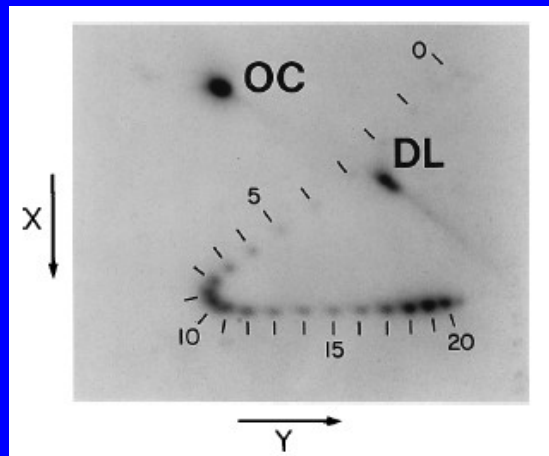
2. HBeAg (early protein)
  - synthesised from precore mRNA

# cccDNA Generation and Processing: cccDNA is a Minichromosome



Bock, T. et al 1994. *Virus Genes*;8:215

Bock, T. et al 2001. *JMB*;307:183

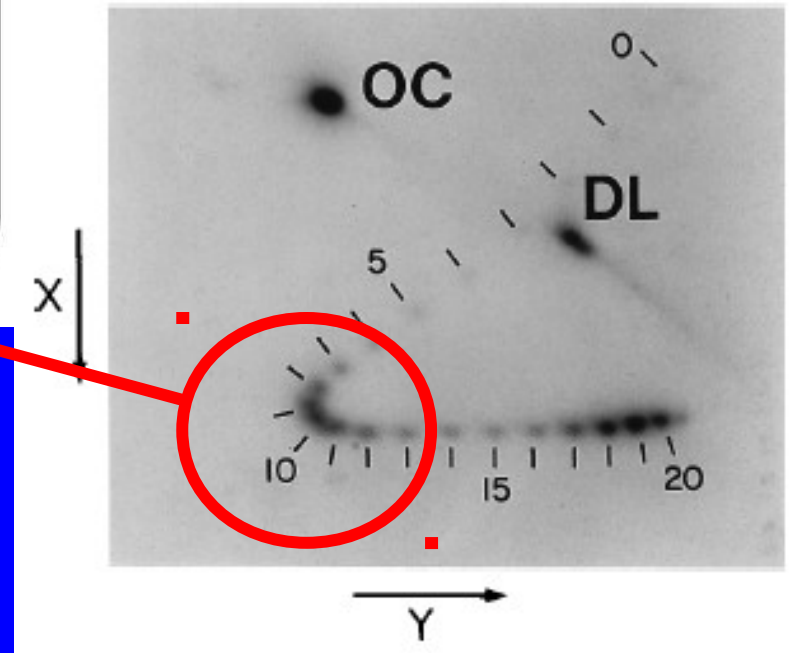
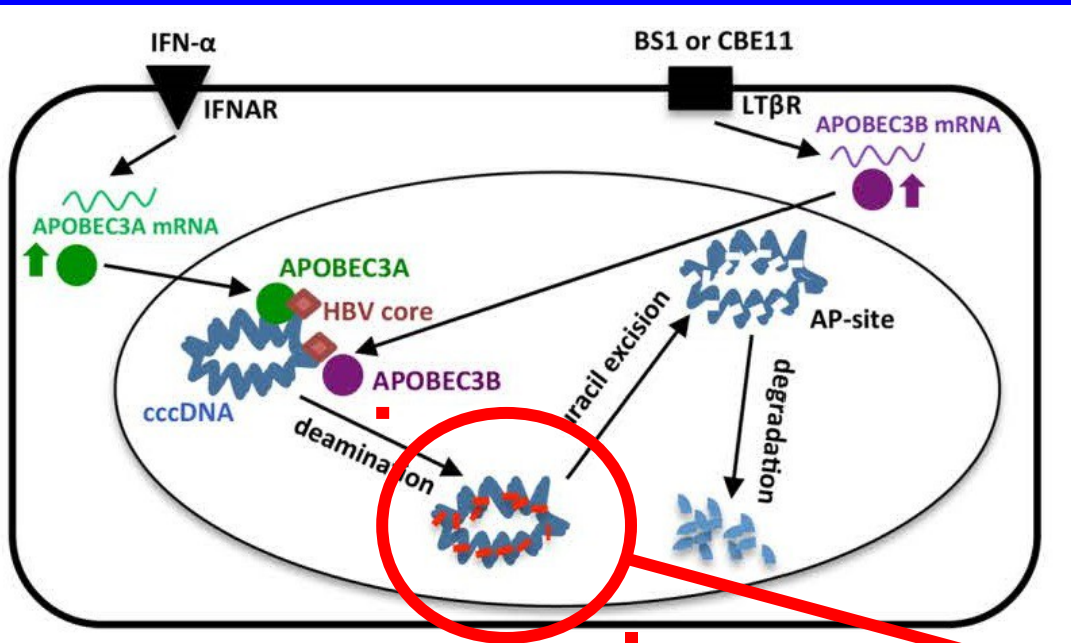


**Low Replication Phenotype**  
Quiescent or active  
Medium to Low Viraemia

**High Replication Phenotype**  
Transcriptionally Active  
High Viraemia

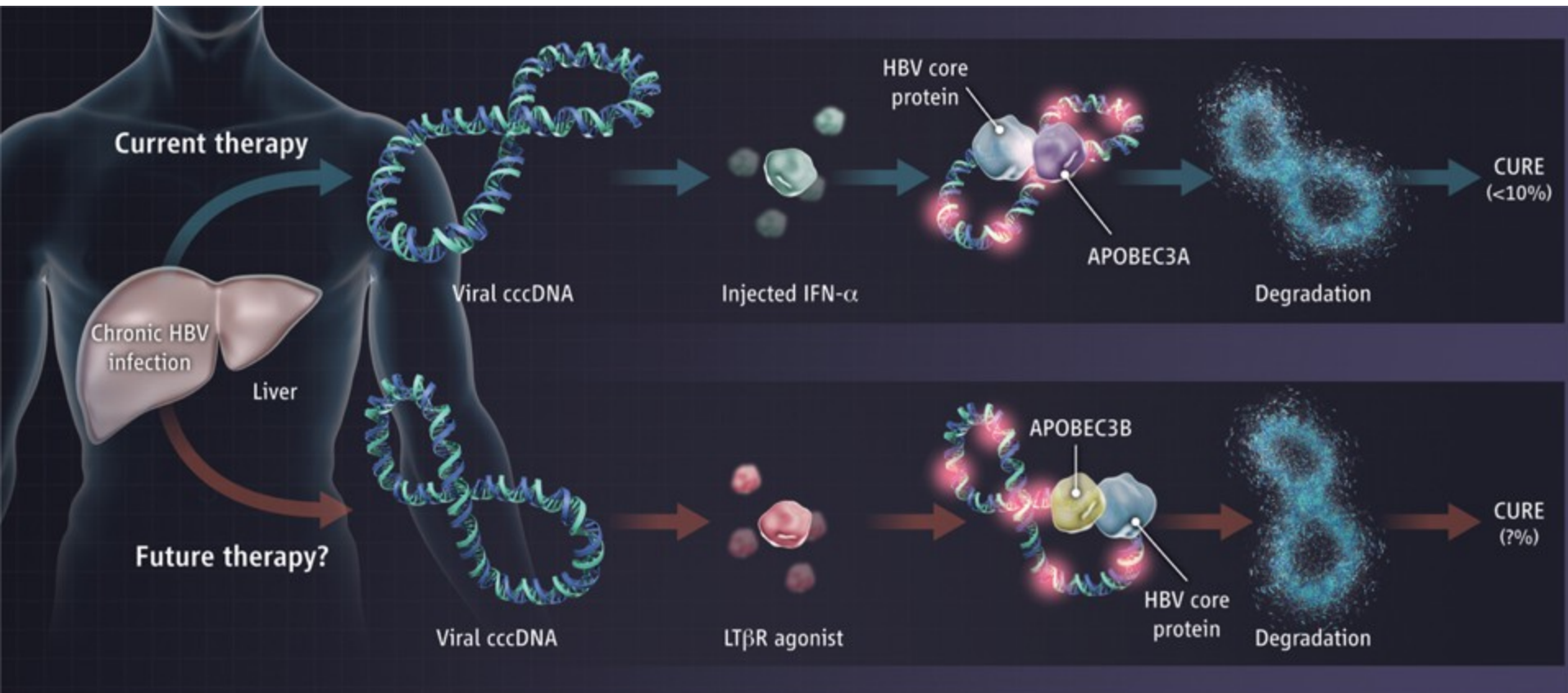


# Interaction of APOBEC 3A/3B, HBV Core Protein (HBc) and cccDNA



# Model for cccDNA Degradation

IFNalpha /Lymphotoxin beta can induce **APOBEC3A/B** dependent degradation of HBV cccDNA



*Lucifora et al, Science 2014; Shlomai & Rice, Science 2014*

**Similar observation with IFN $\gamma$  and TNF $\alpha$  – Xia et al, Gastroenterology 2015**

# HBx Induces Degradation of the Structural Maintenance of Chromosomes (SMC) Complex

## Hepatitis B virus X protein identifies the Smc5/6 complex as a host restriction factor

Adrien Decorsière<sup>1\*</sup>, Henrik Mueller<sup>1†\*</sup>, Pieter C. van Breugel<sup>1†\*</sup>, Fabien Abdul<sup>1\*</sup>, Laetitia Gerossier<sup>2</sup>, Rudolf K. Beran<sup>3</sup>, Christine M. Livingston<sup>3</sup>, Congrong Niu<sup>3</sup>, Simon P. Fletcher<sup>3</sup>, Olivier Hantz<sup>2</sup> & Michel Strubin<sup>1</sup>

*2016. Nature;531:386-389*

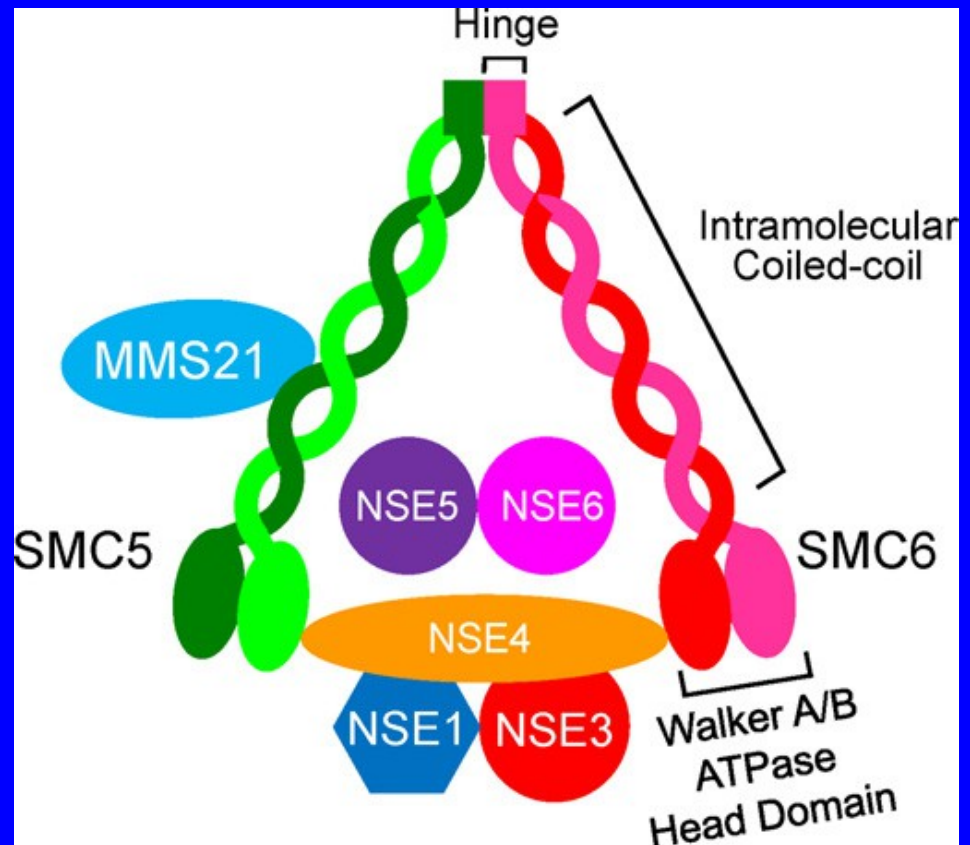
# The Smc5/6 Complex

## Structural Maintenance of Chromosomes (Smc) Complexes

- Condensin
- Cohesin
- Smc5/6 complex

### Smc5/6

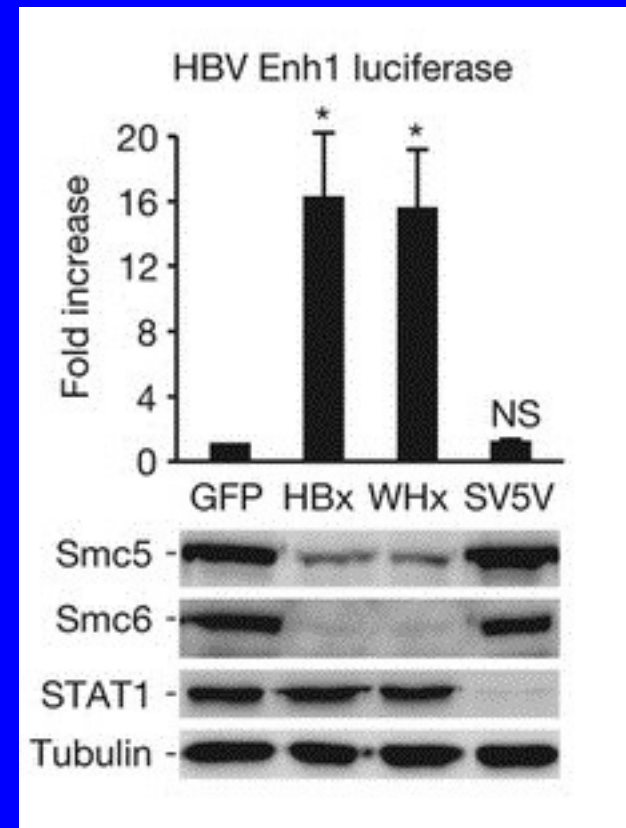
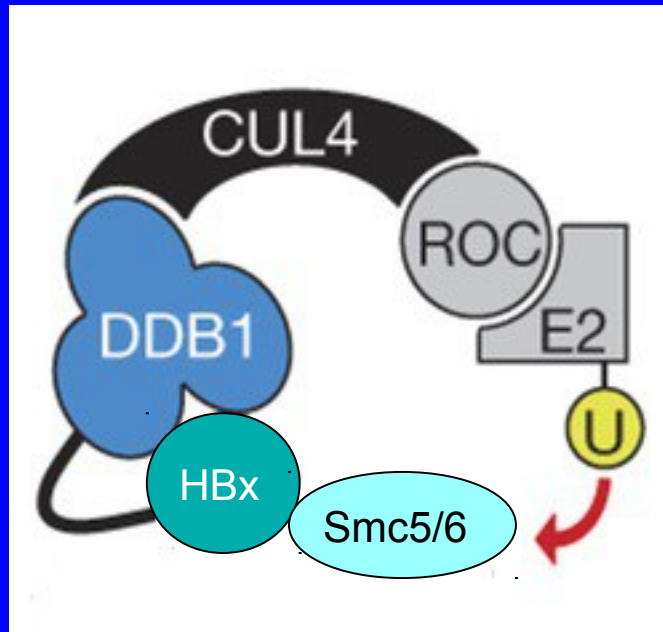
- Nuclear complex
- DNA repair
- Chromosome topology and organization



*Potts, PR. 2009. DNA Repair;8:499*

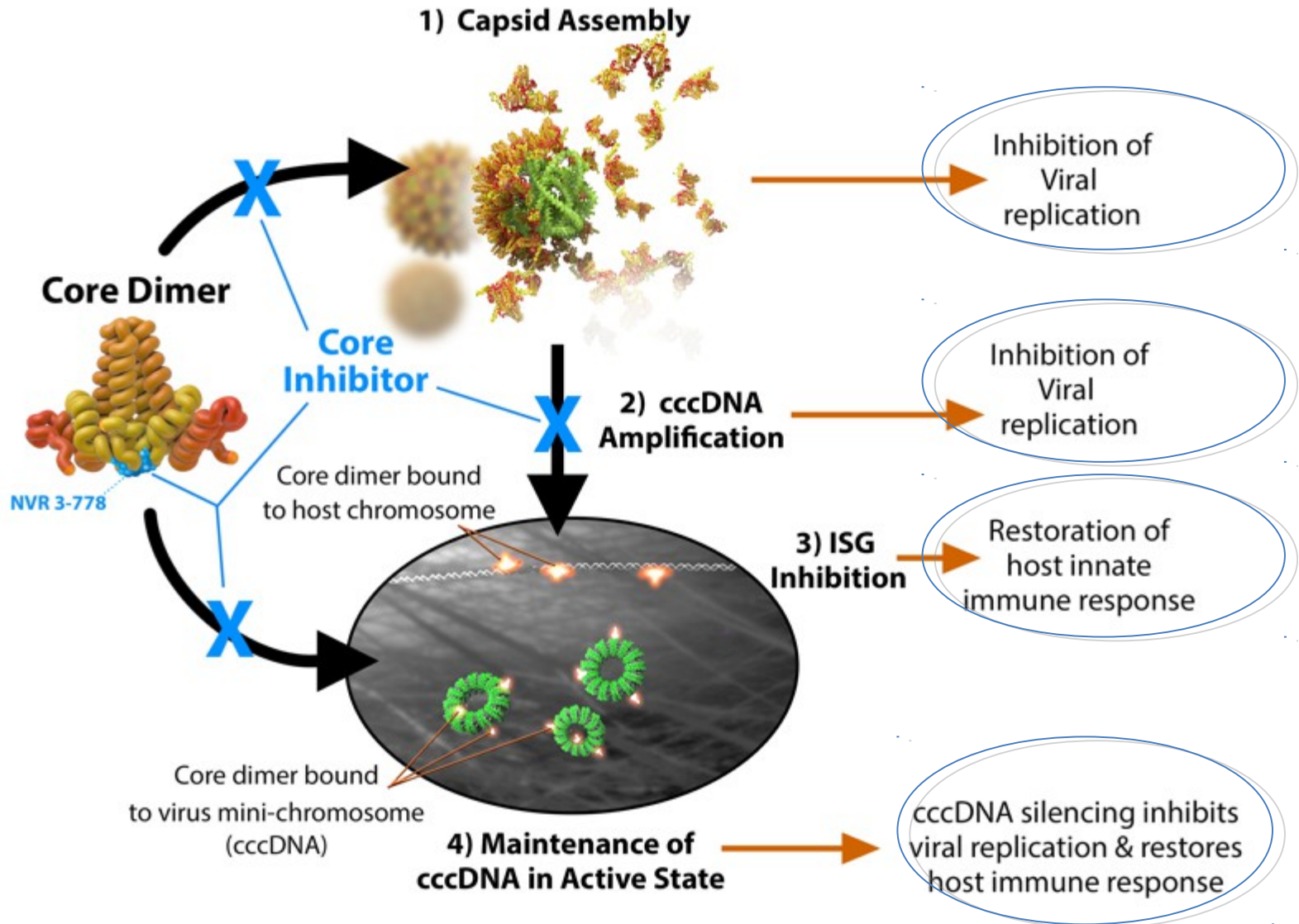
- Depletion of any subunit results in destabilization of the complex (*Taylor, 2008*)

# HBx Induces Degradation of Smc5/6



- The effect is rapid
- Blocked by proteasome inhibitors
- Blocked by E3 ligase inhibitors
- No changes in Smc5/6 mRNA levels

# Targeting the HBV Nucleocapsid



# Targeting HBV Nucleocapsids

## Heteroaryldihydropyrimidines

Destabilization of nucleocapsids

*Deres et al, Science 2003*

*Klumpp et al, PNAS 2015*

## Phenylpropenamide derivatives

Prevent pgRNA encapsidation

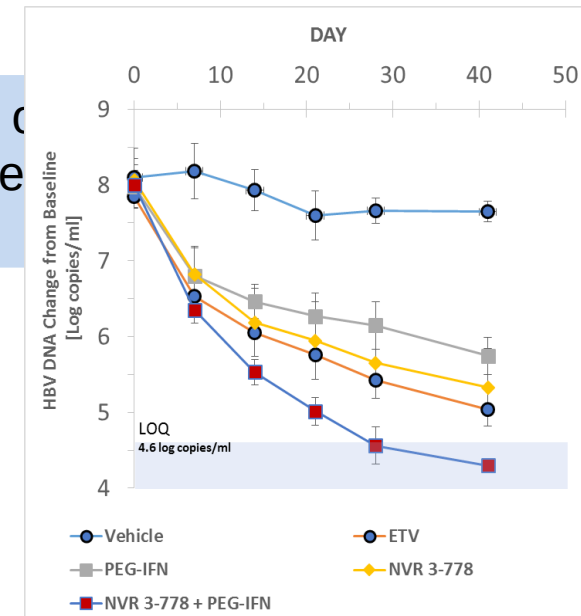
*Feld, J et al 2007. AVR; 76:168-177*

*Antimicrob Agents Chemother. 2002.*

**Novel classes** of capsid inhibitors based on the 3D structure of HBV nucleocapsid  
Novira, Assembly Biosciences, Janssen, Roche, and others  
Phase 1 studies with Novira completed

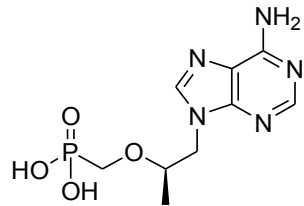
*Lam A, et al. AASLD 2015, San Francisco. #33*

[sulphonamide/sulfamoyl benzimide derivatives]



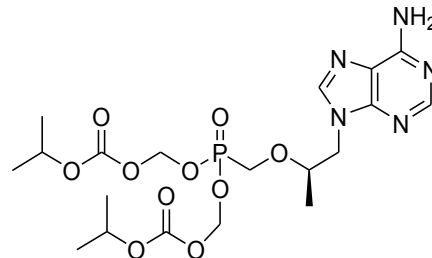
# Reverse Transcription: Improved Potency of NA Tenofovir Alafenamide (TAF)

- TAF = orally bioavailable phosphonoamidate prodrug of tenofovir (TDF)
- In comparison with tenofovir, TAF enables enhanced delivery of the parent nucleotide and its active diphosphate metabolite into lymphoid cells and hepatocytes.
- This is attributed to an improved plasma stability and differential intracellular activation mechanism for TAF relative to TDF

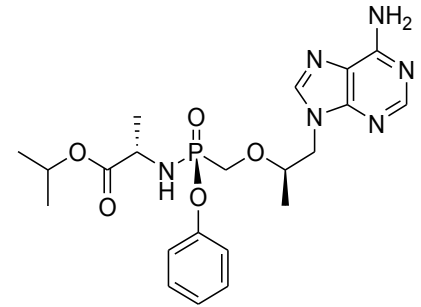


**EC<sub>50</sub> HIV-1  
(PBMCs)**

**Tenofovir  
1.2 μM**



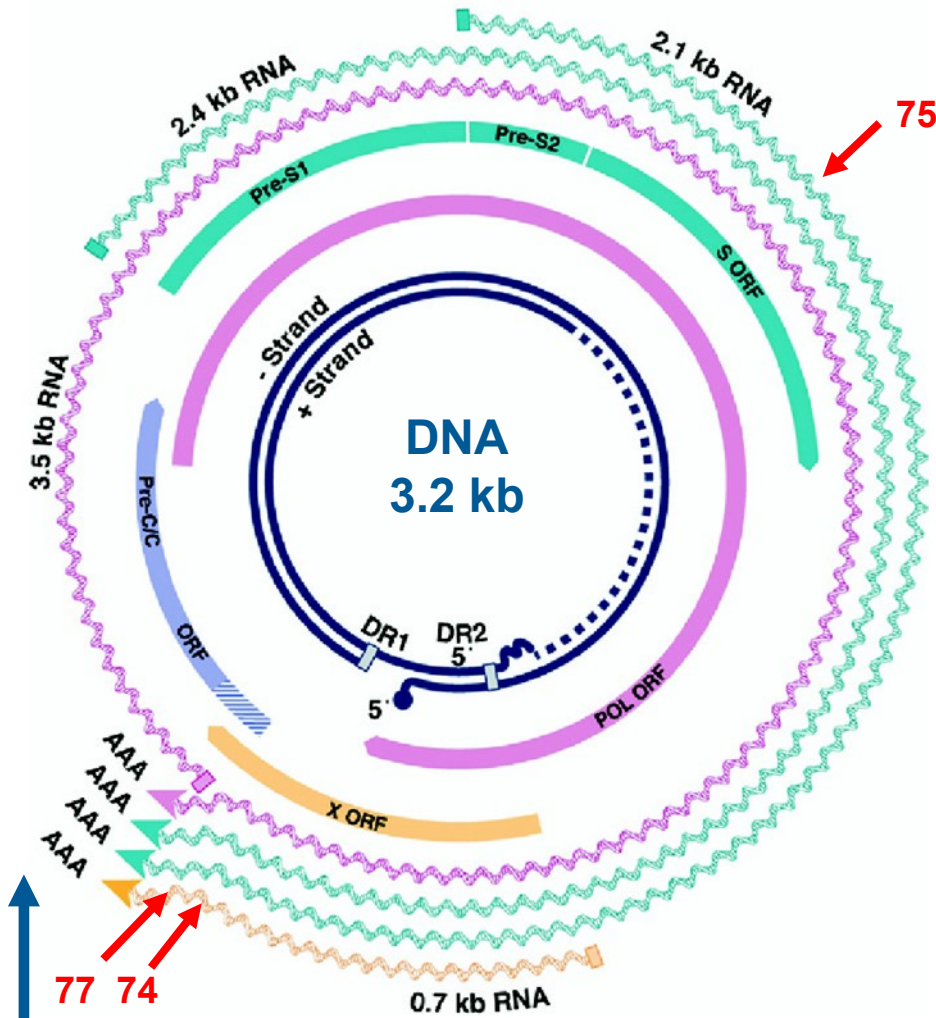
**Tenofovir Disoproxil  
0.015 μM**



**TAF  
0.003 μM**



# HBV genome and siRNA target sites



Same polyadenylation signal for all mRNAs

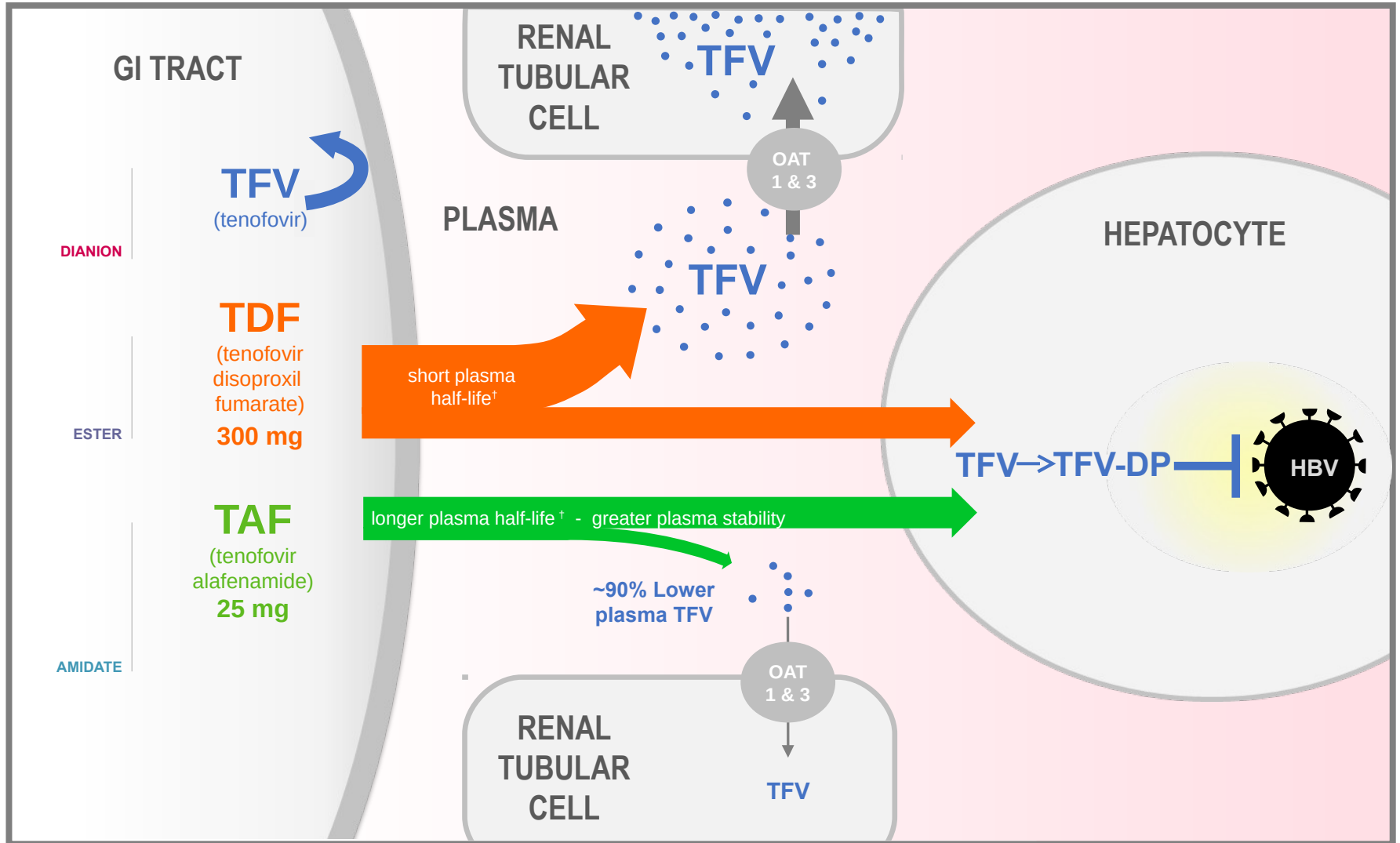
## •HBV mRNA

- 3.5 kb pre-genomic RNA
- 3.5 kb pre-core mRNA
- 2.4 kb pre-S1 mRNA
- 2.1 kb pre-S2/S mRNA
- 0.7 kb X mRNA

## •HBV proteins

- Polymerase (with reverse transcriptase function)
- Core (HBcAg), forms capsid
- E antigen (HBeAg), also called pre-core, a secreted protein
- Large, middle and small surface proteins (HBsAg), form envelope
- X protein (Transactivator)

# Tenofovir alafenamide (TAF) – A Novel Prodrug of Tenofovir

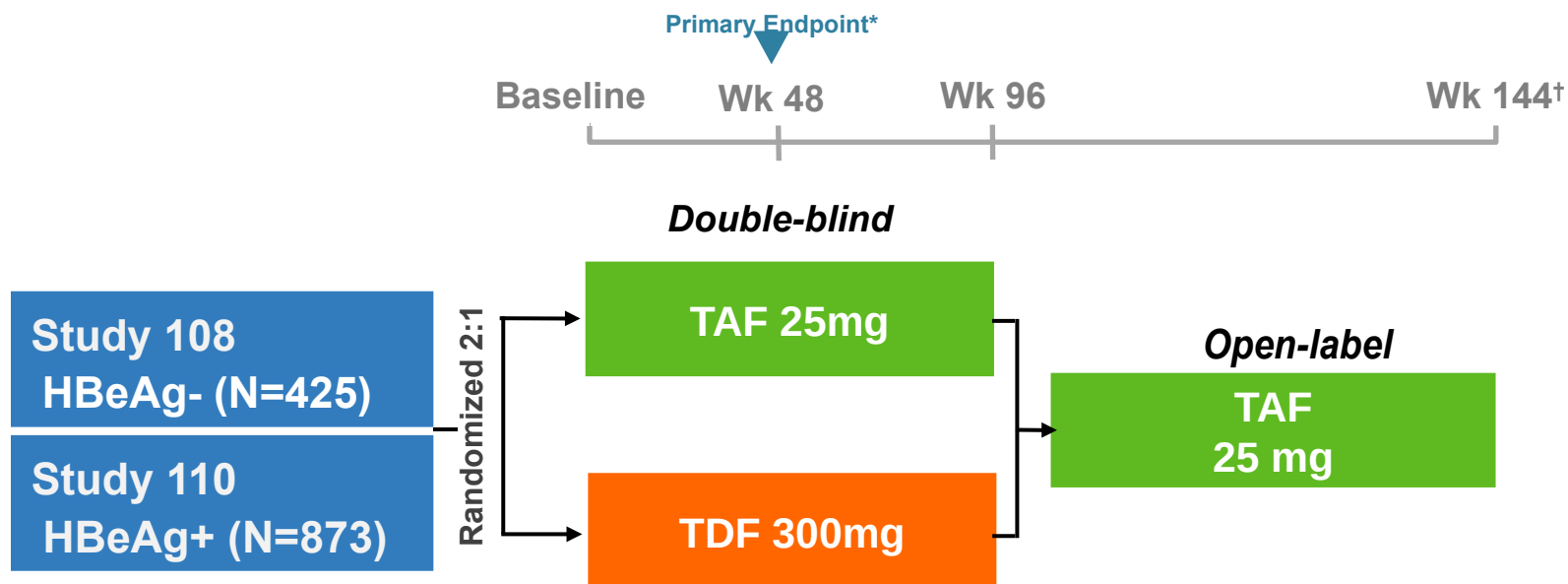


<sup>†</sup>T<sub>1/2</sub> based on *in vitro* plasma data - TDF = 0.4 minutes, TAF = 90 minutes.



Study 108 and 110: Phase 3 CHB Studies: TAF vs TDF

# TAF HBV Phase 3 Program



- Two phase 3, randomised, double-blind studies
- Inclusion criteria
  - HBV DNA  $\geq 20,000$  IU/mL; ALT  $>60$  U/L (males),  $>38$  U/L (females)
- Primary endpoint (non inferiority margin of 10%):
  - HBV DNA  $<29$  IU/mL at Week 48
- Key secondary safety endpoints
  - Bone mineral density and renal parameters at Week 48

<sup>†</sup>Amendment to extend double-blind to Week 144 and open-label phase to Week 384 (Year 8) is currently underway

\*Non-inferiority margin of 10%

Buti EASL 2016, Oral GS06; Chan, EASL 2016, Oral GS12

## HBV DNA Response at 48 Weeks

Study 108  
(HBeAg-subjects)

HBV DNA <29 IU/mL (%)

Log<sub>10</sub> HBV DNA Change



● TAF  
● TDF

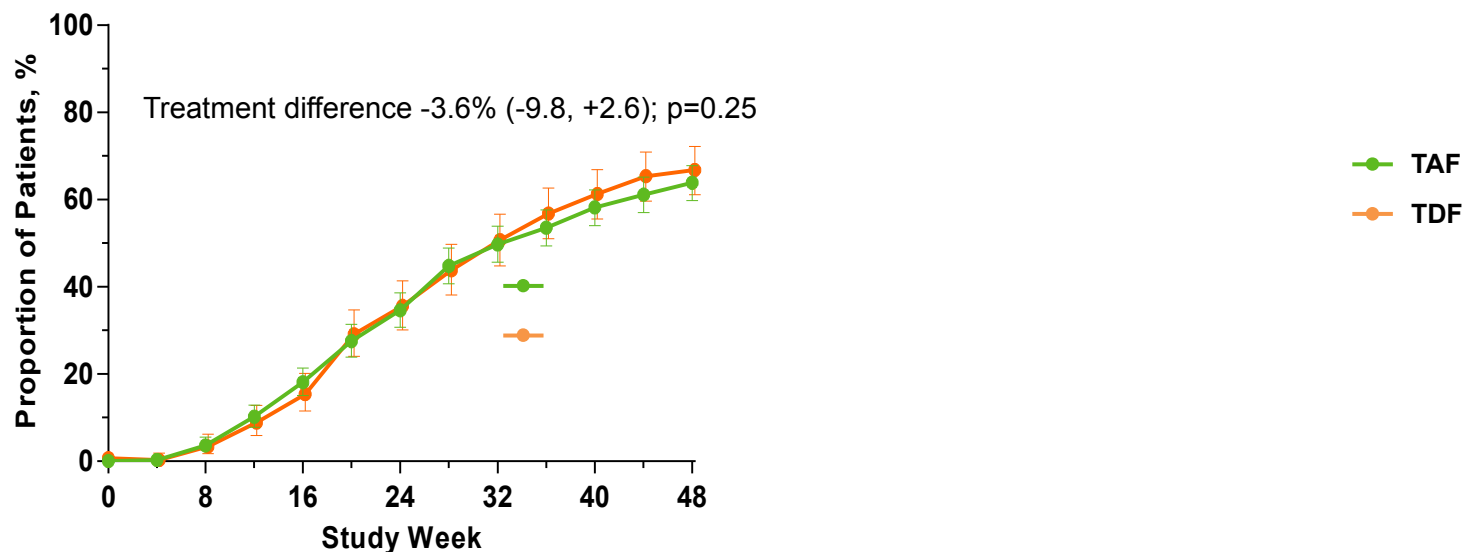
- Similar and non-inferior rates of virologic suppression with TAF and TDF at Week 48
- No resistance detected in either treatment group

## HBV DNA Response at 48 Weeks

HBV DNA <29 IU/mL (%)

Log<sub>10</sub> HBV DNA Change

Study 110  
(HBeAg+ subjects)



- Similar and non-inferior rates of virologic suppression with TAF and TDF at Week 48
- No resistance detected in either treatment group



# Results: Renal Safety

Mean ( $\pm$ SD) change in eGFR<sub>CG</sub> (mL/min)

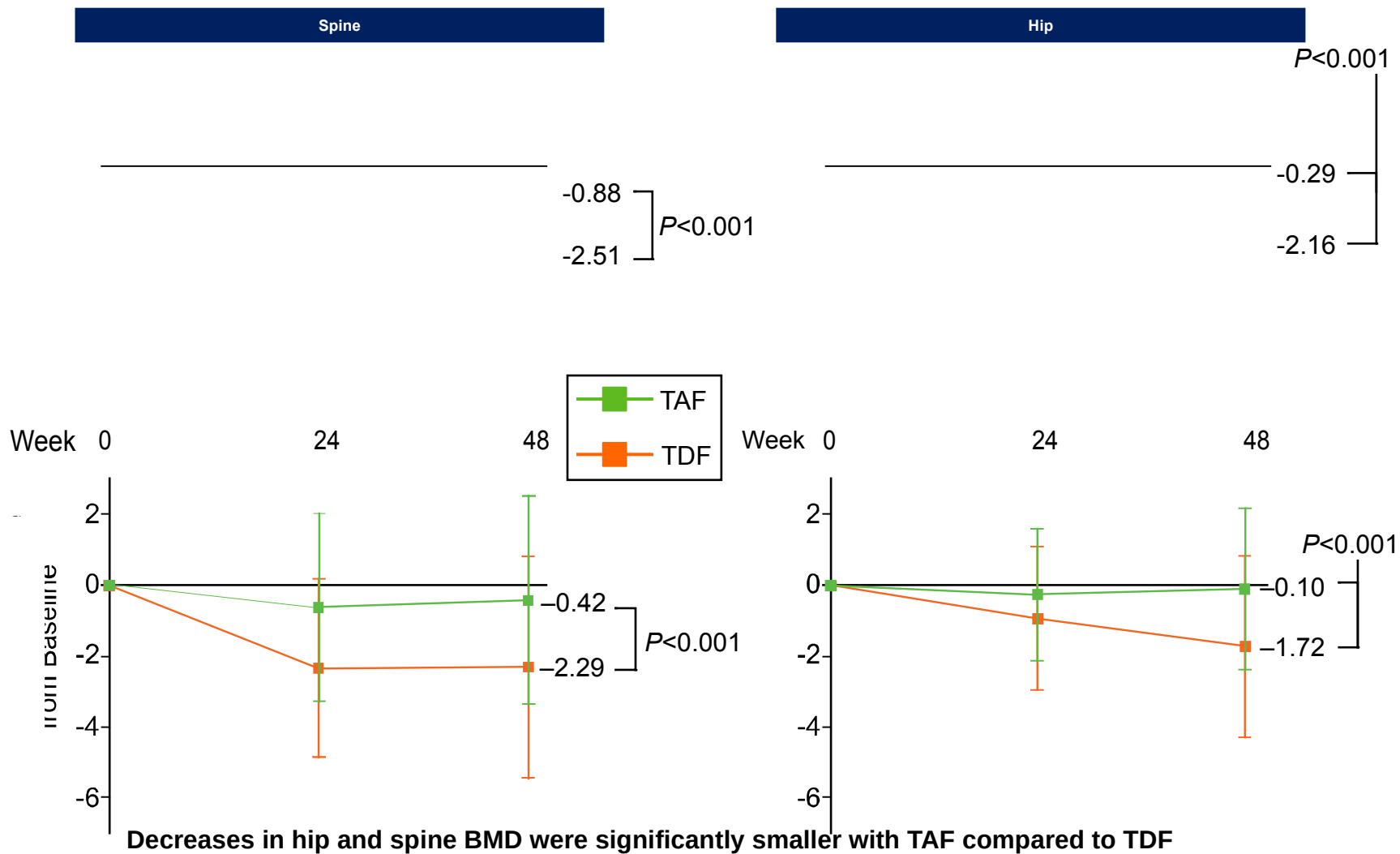
-0.6  
-4.7 } p < 0.001

	TAF n=866	TDF n=432	P-value
Change in sCr, mg/dL	0.010 (0.11)	0.024 (0.10)	0.012

# Changes in Spine and Hip BMD Through Week 48

Study 108

Study 110

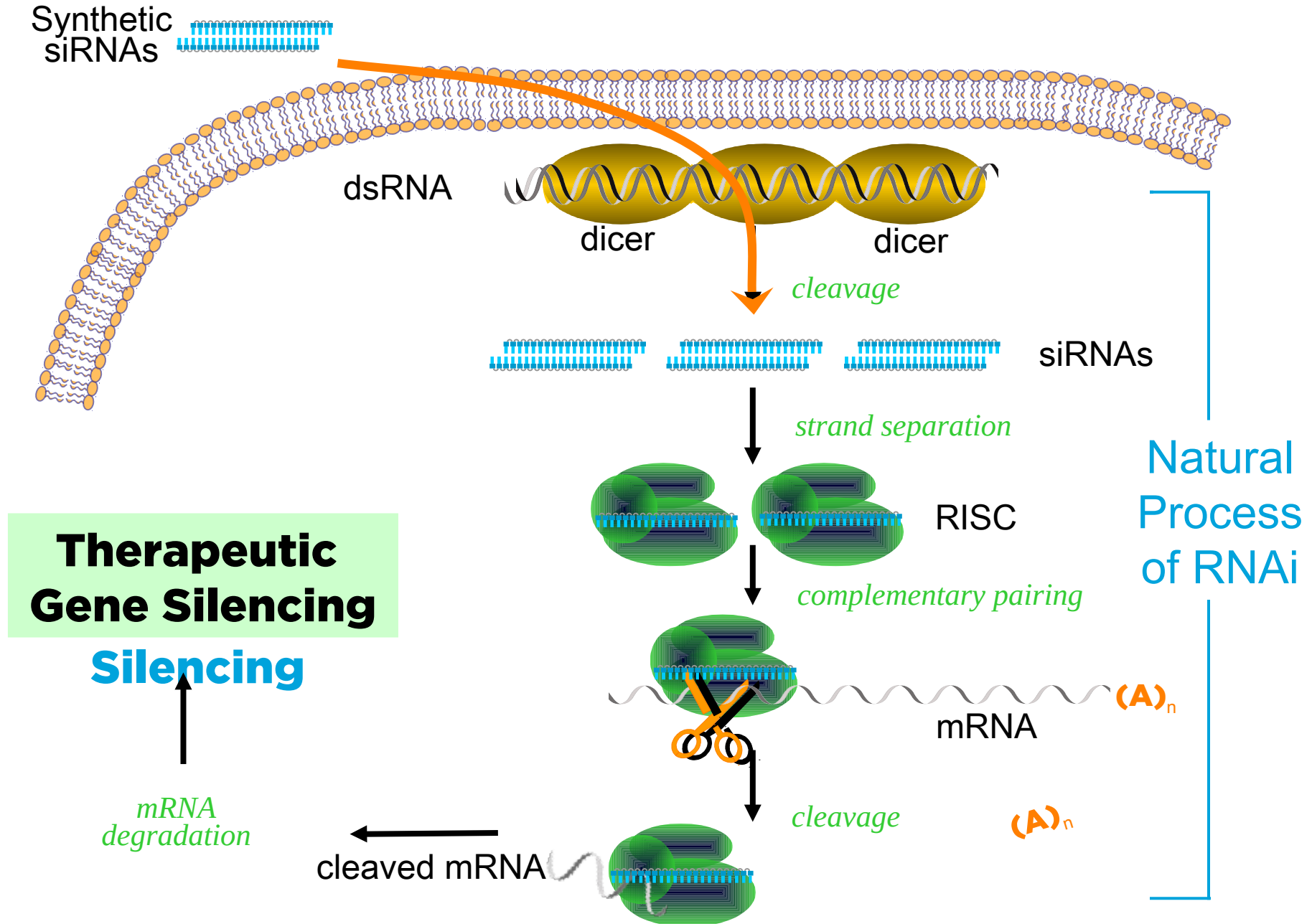


# Authors' Conclusions

- Treatment with TAF for 48 weeks demonstrated:
  - Non-inferior efficacy to TDF for the proportion with HBV DNA <29 IU/mL
  - Improved rates of ALT normalization
  - No resistance development in either treatment group
  - Rates of HBeAg loss and seroconversion similar to TDF in Study 110
- TAF was well tolerated in HBeAg-negative and -positive patients
  - Treatment-emergent AEs similar to TDF
  - Significantly smaller increases in SCr (integrated safety analysis) and decreases in eGFR<sub>CG</sub> compared to TDF, with improved markers of renal tubular function
  - Significantly less declines in hip and spine BMD compared to TDF, with improved bone biomarkers



# Mechanism of RNA Interference (RNAi)



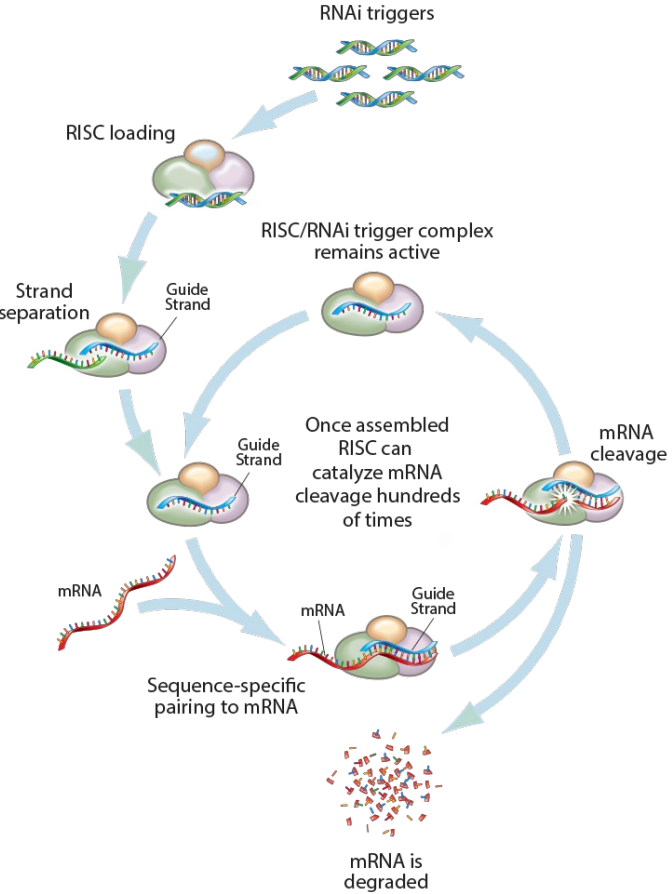
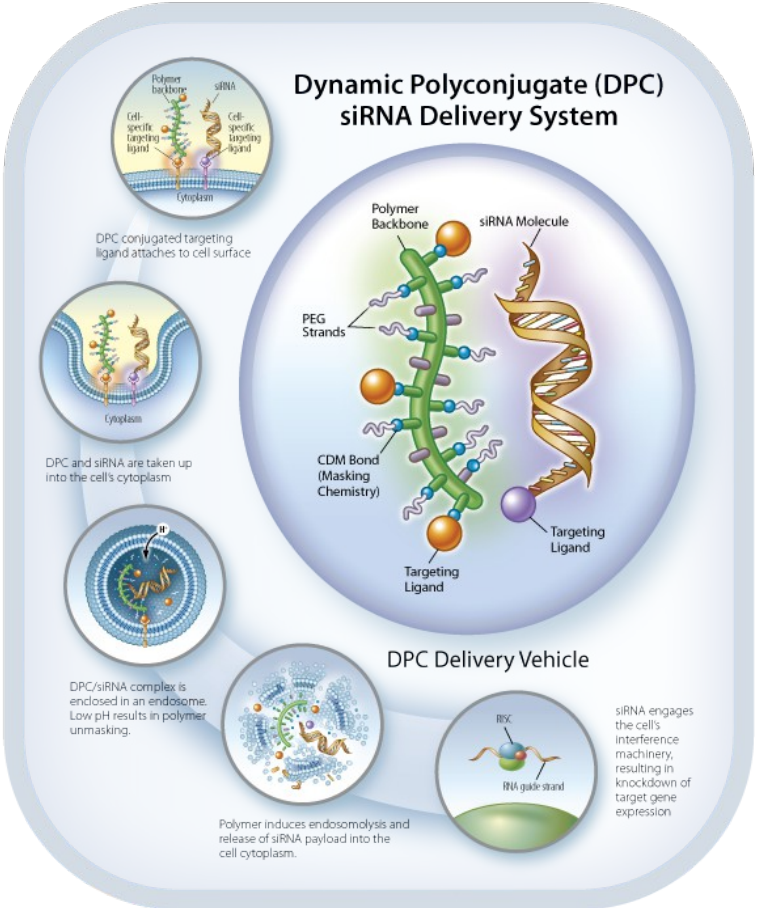
# Groups Involved in RNAi Therapy and HBV

- Arrowhead Pharmaceuticals
  - ARC-520 (phase 2)
  - ARC-521 (phase 1/2)
- Arbutus Biopharma
  - ARB-1467 (phase 1/2)
- Alnylam Pharmaceuticals
  - ALN-HBV (phase 1)

# ARC-520 RNAi delivery technology

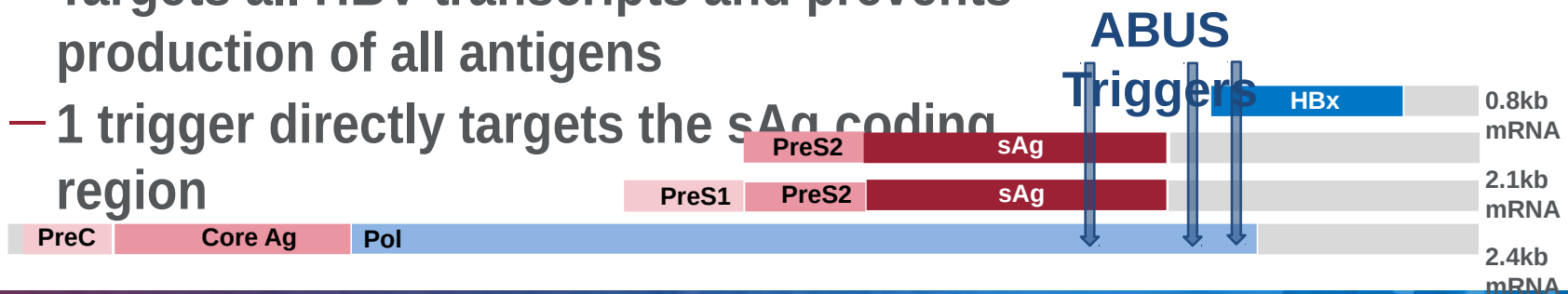
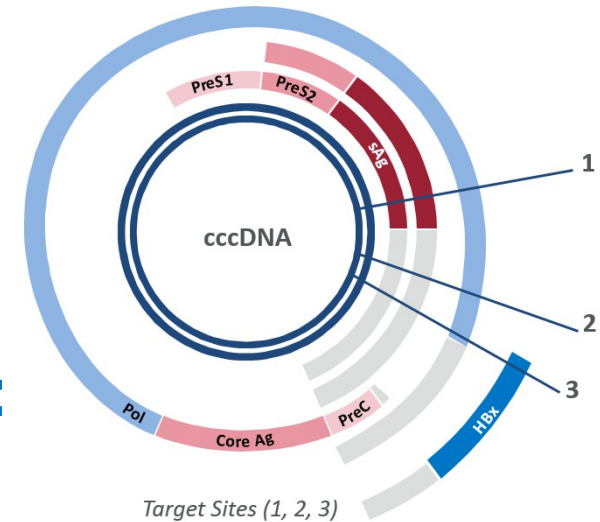
## DPC polymer composition: amphipathic peptide with reversibly "masked" amines

- Ligand-driven cellular uptake of (N-acetyl galactosamine for hepatocytes)
- Liver tropic siRNA attachment by lipophilic ligand (e.g. cholesterol)
- ↓ pH in endosomes unmasks peptide to disrupt endosomal membrane
- siRNA released to cytoplasm



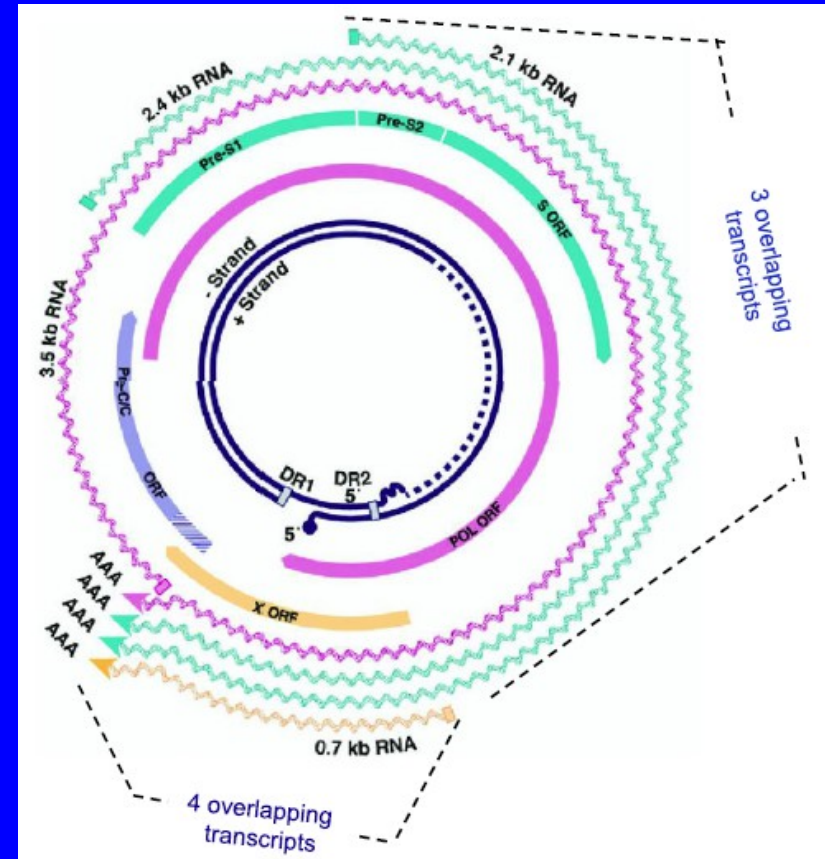
# ARB-1467 Targets Multiple HBV Genomic Sites

- Primary viral target is HBsAg
- Target sites are regions of high conservation in HBV viral genomes
- Advantages of the **3-trigger combo**:
  - Increased potency
  - Coverage extension to 99.8% of HBV genotypes
  - Targets all HBV transcripts and prevents production of all antigens
  - 1 trigger directly targets the sAg coding region



# Alnylam RNAi (Preclinical)

- Delivery: **Multi-component lipid nanoparticles** for delivery to the liver via LDL receptor
- Triantennary Gal/Nac conjugated to 3' end of sense strand of siRNA
- Two target regions:
  - 0.7 kb region overlapping across all 4 HBV transcripts.
  - 1.4 kb region overlapping across 3 transcripts
- Inhibits replication, assembly and secretion of virus as well as subviral antigens that overlaps across 3 HBV transcripts



# ARC-520 Produces Deep and Durable Knockdown of Viral Antigens and DNA in a Phase II Study in Patients with Chronic Hepatitis B

HBV antigen reduction in ETV  
experienced HBeAg-positive  
patients with a single 4 mg dose  
(cohort 5)

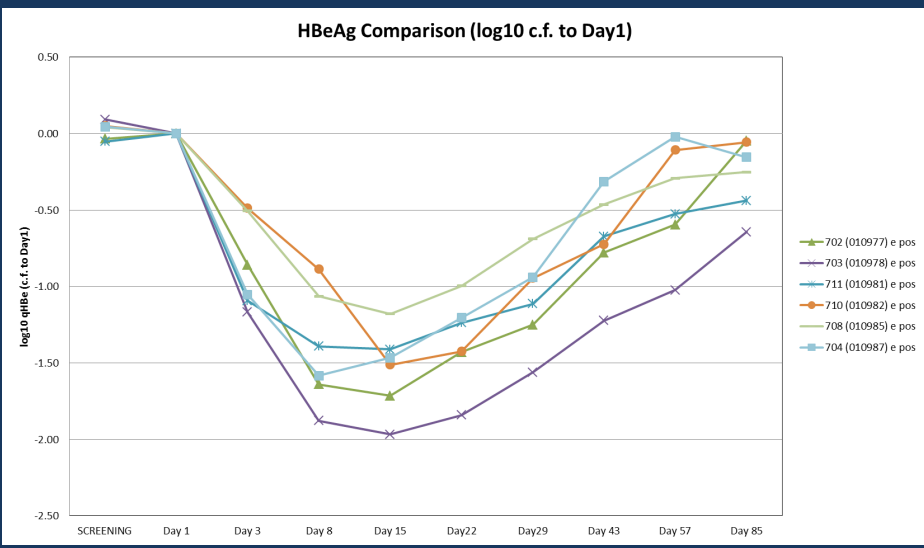
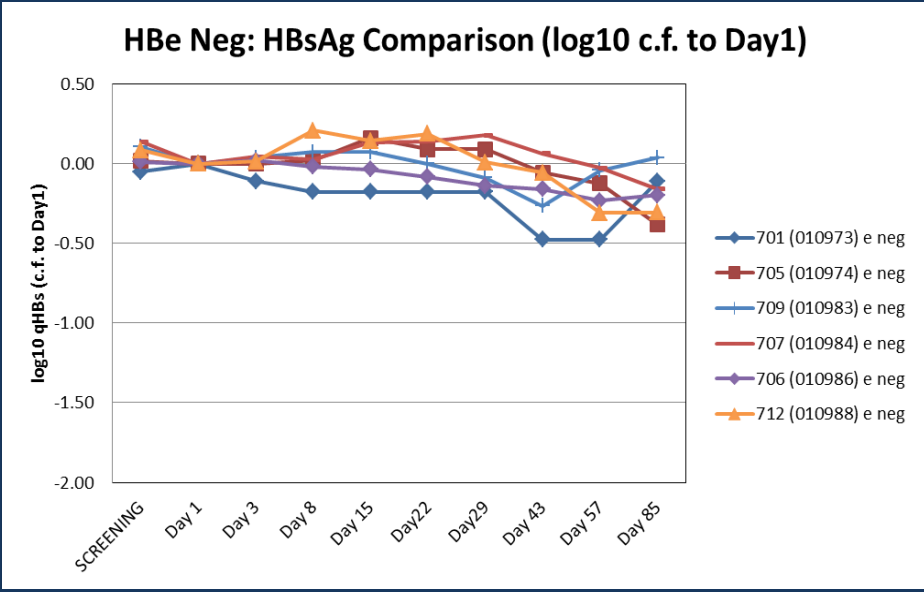
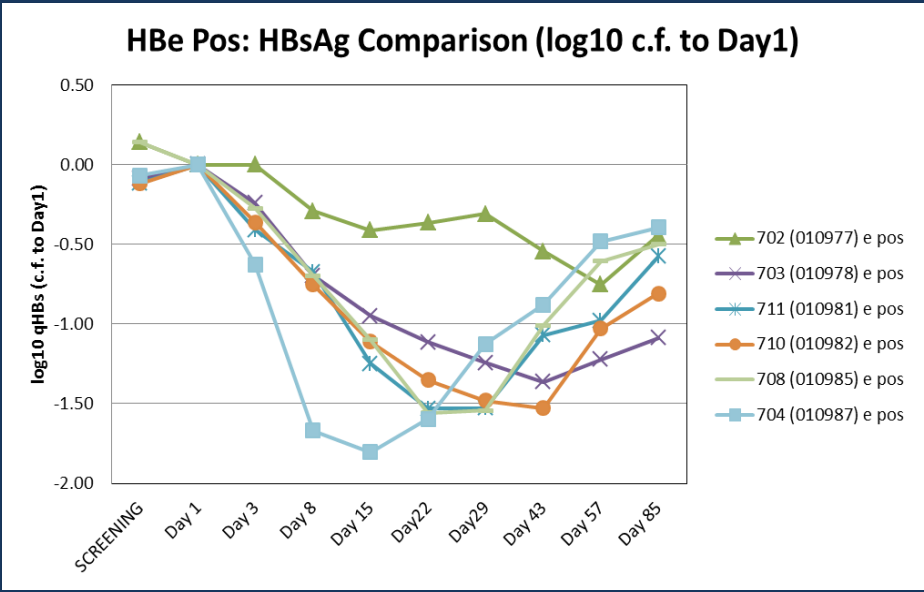
HBsAg reduction in ETV naïve  
patients with a single 4 mg dose  
(cohort 7)

Direct antiviral effect lasted up to 57 days  
after a single dose of ARC-520, delayed  
response duration >85 days

- Small dose-related reduction in HBsAg
- Maximum effective dose not reached
- HBV DNA results pending in ETV naïve patients

# ARC-520 RNAi: clinical responses

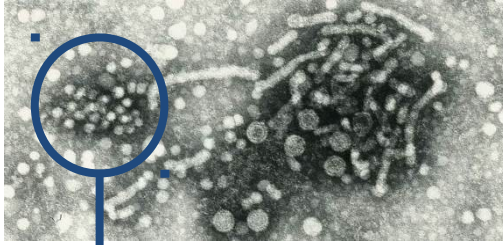
**NUC naïve cohort (n=12):** 50% HBeAg positive, 1x 4mg dose



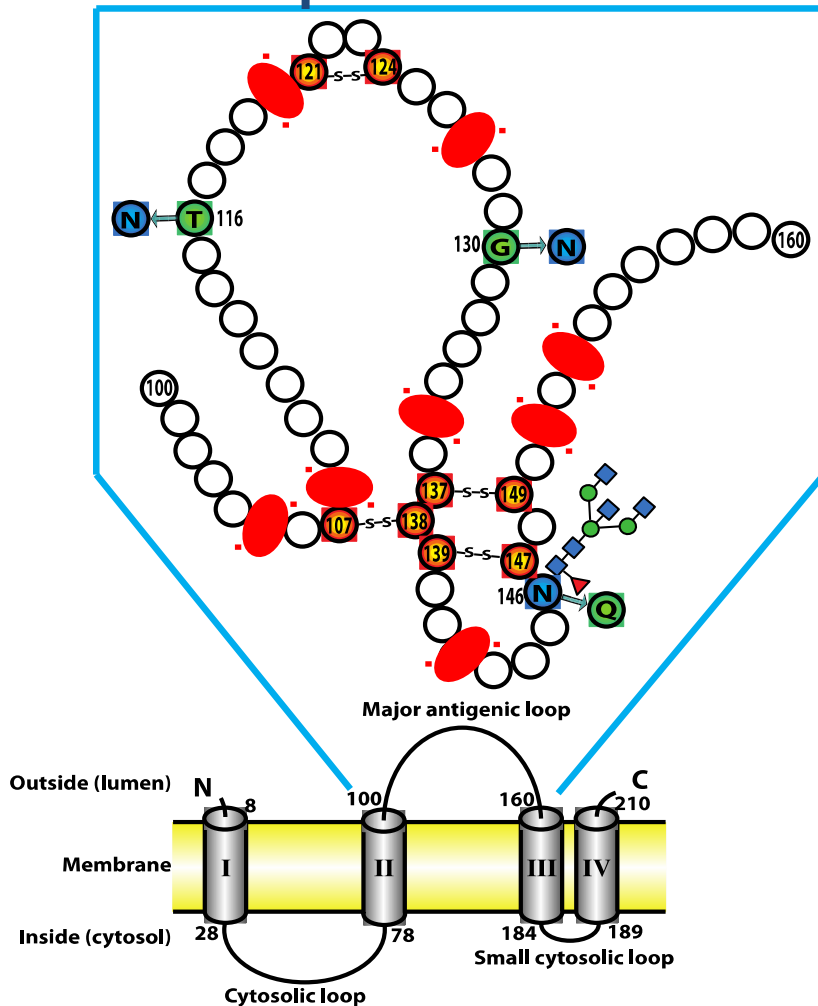
**HBsAg:**  
 •> 1log drop in HBsAg achieved by all HBe pos subjects (excluding 702; transitional HBe <0.1 PE IU/mL at BL)

**HBeAg:**  
 •> 1log decline in HBeAg achieved

# Immune Regulation by HBsAg



- HBsAg secreted in vast excess over virions ( $>10^3$  fold)
- Circulate in blood 100-400  $\mu\text{g/ml}$  (1% of total serum protein)
- Unique conformational structure (8 cysteines ● and 8 prolines ●)
- Associated with increased risk of HCC (Yuen, MF. et al 2008. *Gastro*;135:1192–1199)
- Plays a key role in HBV persistence
- Suppress both innate (TLR-2, TLR-9 and IFN- $\alpha$ ) as well as adaptive (mDC) responses to infection



Wang, S et al 2013. *J Immunol*;190:5142.; Xu, Y et al 2009. *Mol Immunol*;46:2640.; Op den Brouw, ML et al 2009. *Immunol*;126:280.



# Importance of HBsAg Clearance/Seroconversion

- ▽ ↓ Hepatic decompensation
- ▽ ↓ HCC
- ▽ ↑ Survival
- ▽ ↓ Levels of cccDNA

- **As close to cure as we can expect to achieve in chronic hepatitis B**

# Egress

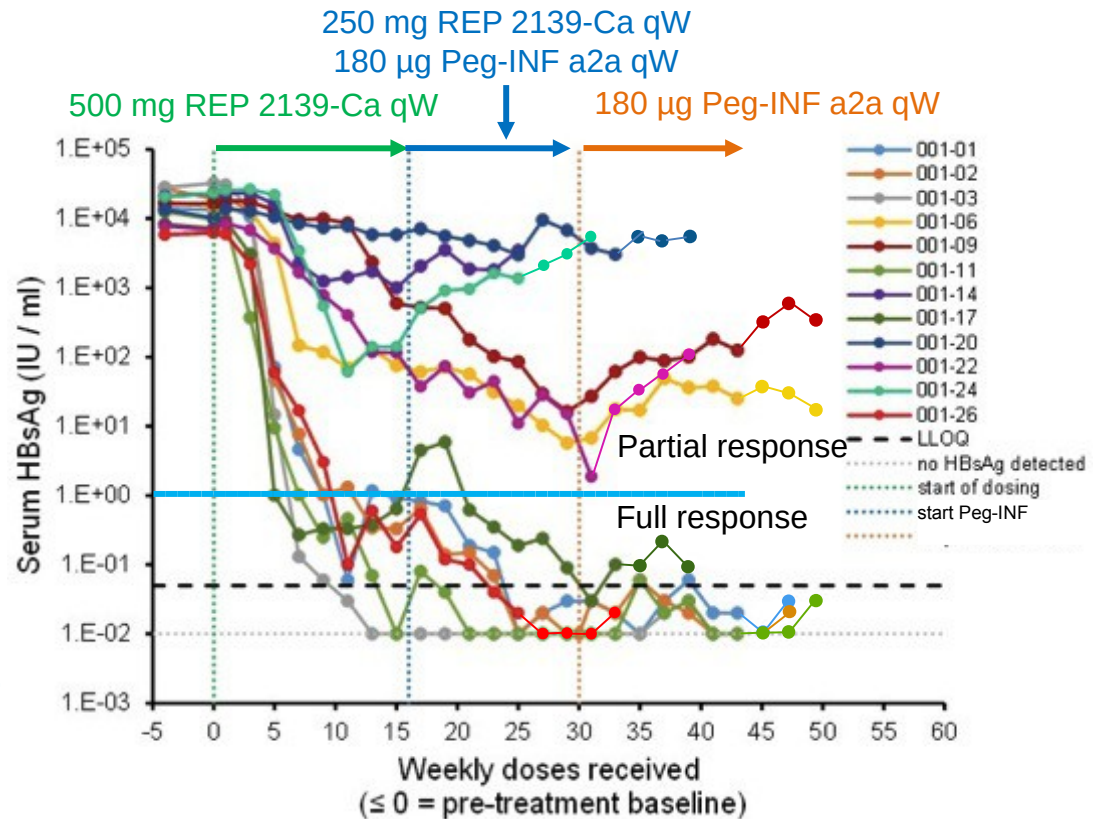
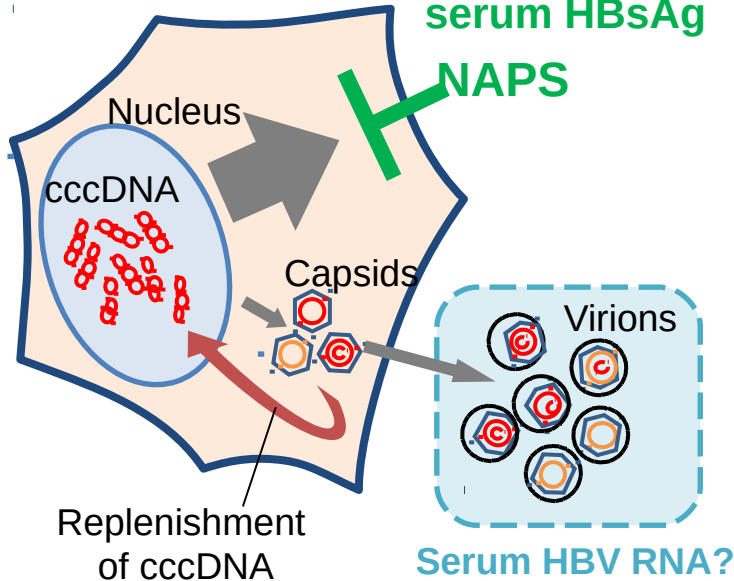
## Update on the Safety and Efficacy of REP 2139 Monotherapy and Subsequent Combination Therapy with Pegylated Interferon Alpha-2a in Chronic HBV/HDV Co-Infection in Caucasian Patients

### MOA of Nucleic Acid Polymers (NAP)

•NAPs have entry and post-entry antiviral effects in HBV infection *in vitro*<sup>1</sup>

Restoration of immune response?<sup>2,3</sup>

Elimination of serum HBsAg



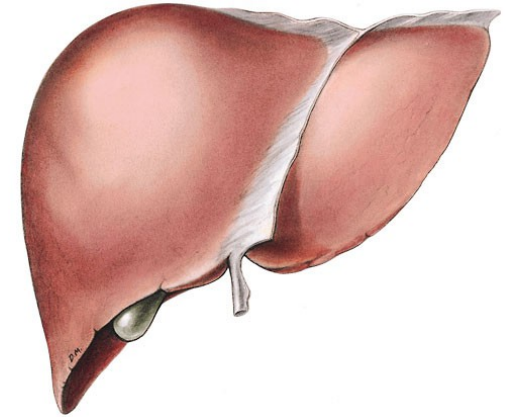
Bazinet M, et al. AASLD 2015, San Francisco. #31.

1. Noordeen, F et al. AAC. 2013; 2. Wu et al. Hepatology. 2009; 3. Boni et al. Gastroenterology. 2012

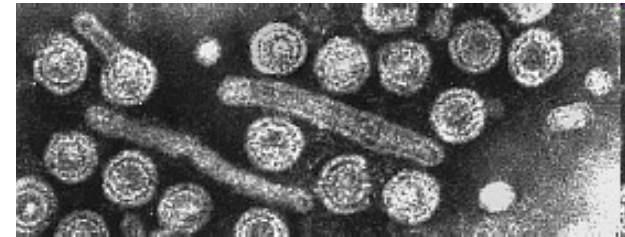
# Hepatitis B Virus Immune Evasion/Exhaustion

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## ➤ Hepatotropic virus



## ➤ High Antigen burden



## ➤ Immune regulation at the innate and adaptive level

# Rationale for Immunotherapy in HBV

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- **Spontaneous sustained immune control in adults following acute infection**
- **Bone marrow transplant > Immune reconstitution > clearance of chronic HBV**
- **Patients that spontaneously clear chronic HBV display robust T cell responses**

# Potential Immunological Targets for HBV Therapy

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## Innate

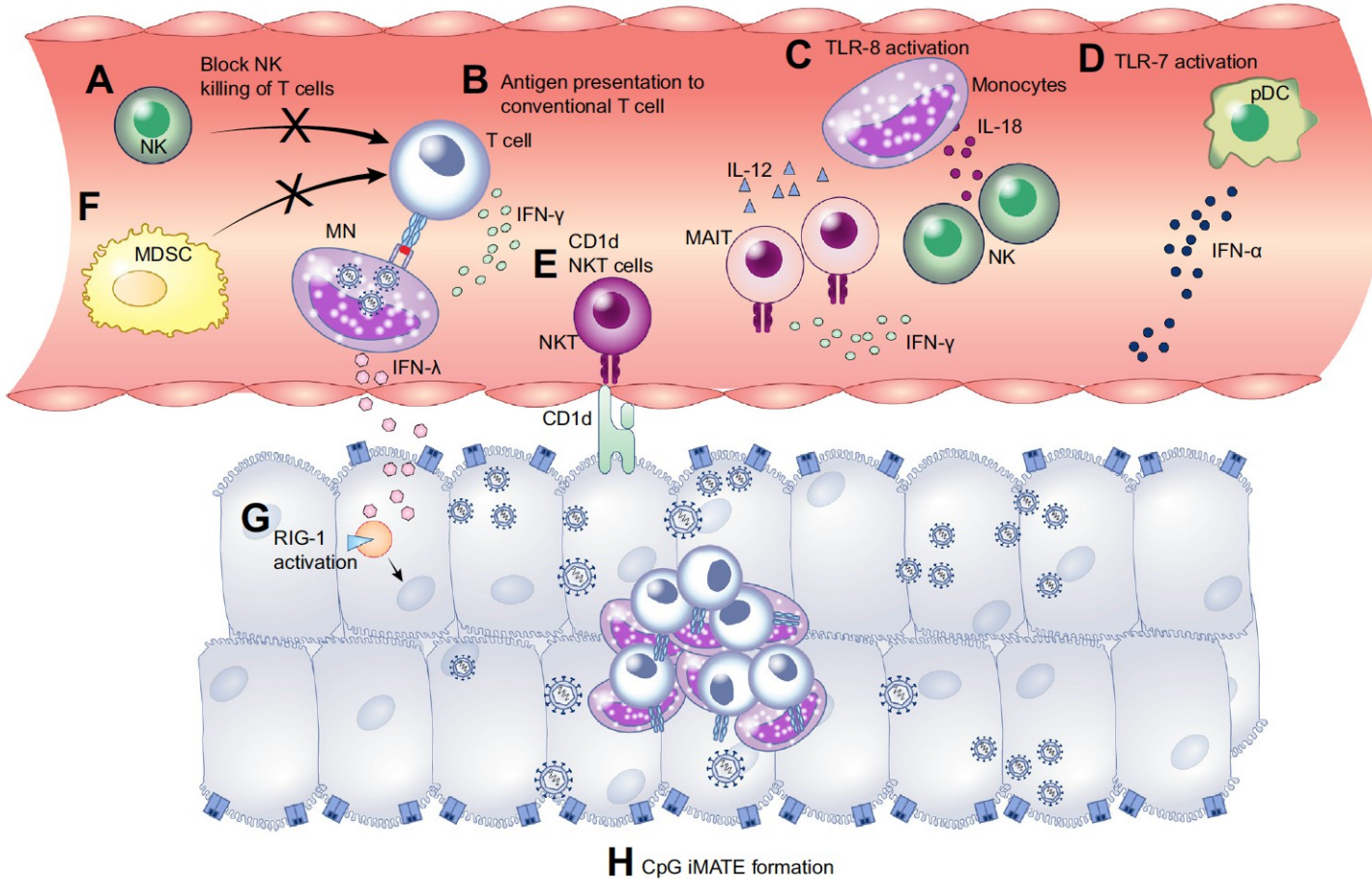


## Adaptive

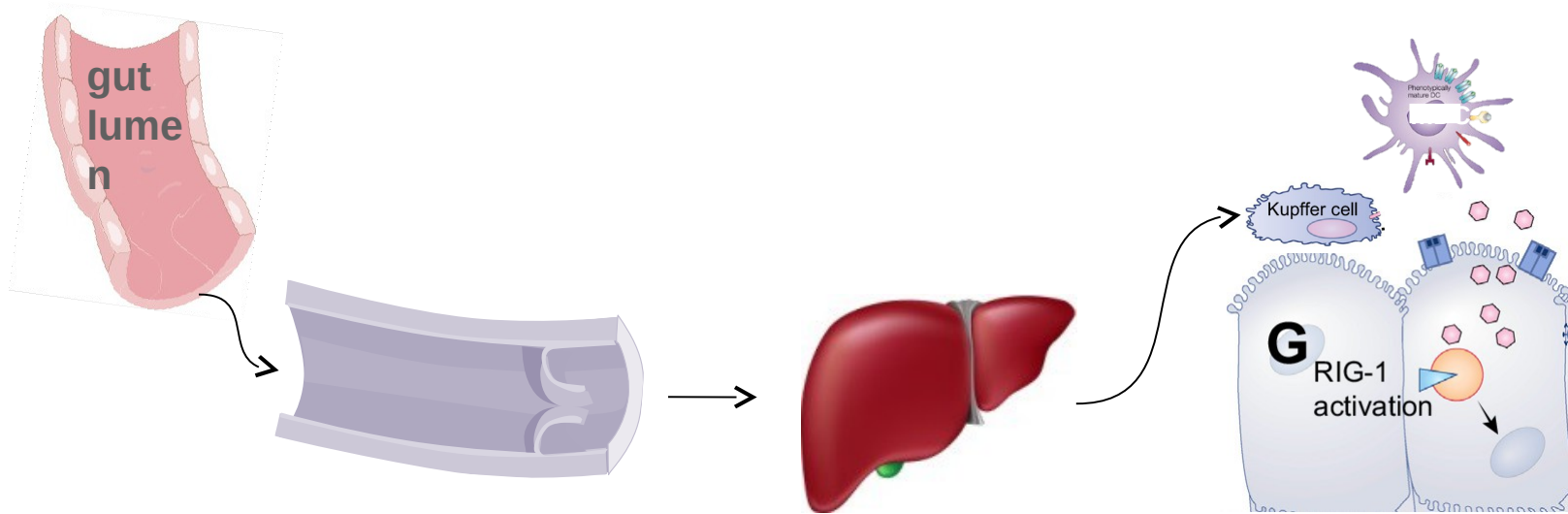


# Opportunities for Innate-targeted Immunotherapy

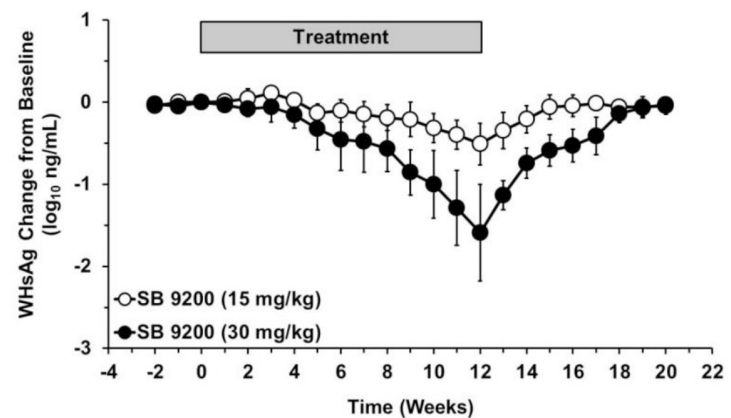
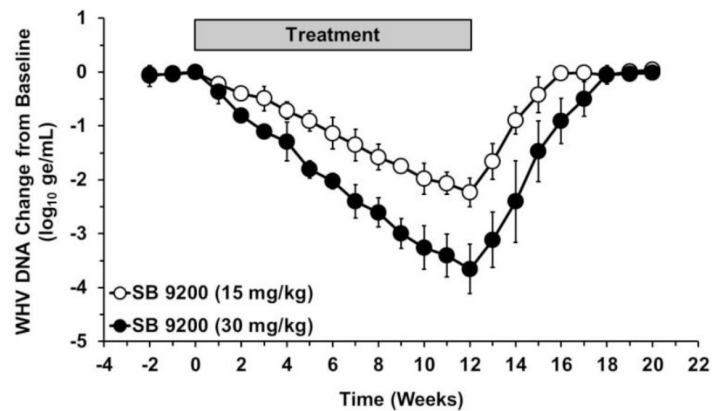
## - Stimulating antiviral cytokine production --



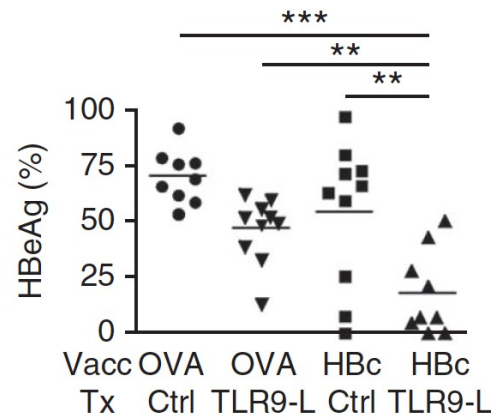
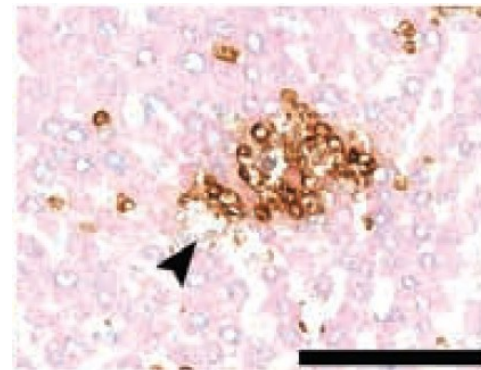
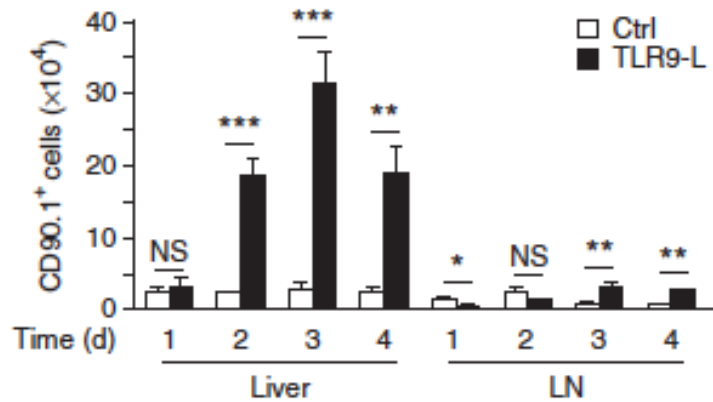
# Direct triggering of RIG-I in infected hepatocytes



## Chronic Woodchuck Hepatitis Virus Infection model



# CpG induction of iMATEs





# Potential Immunological Targets for HBV Therapy

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## Innate



## Adaptive



# Therapeutic Vaccination in Chronic HBV Infection

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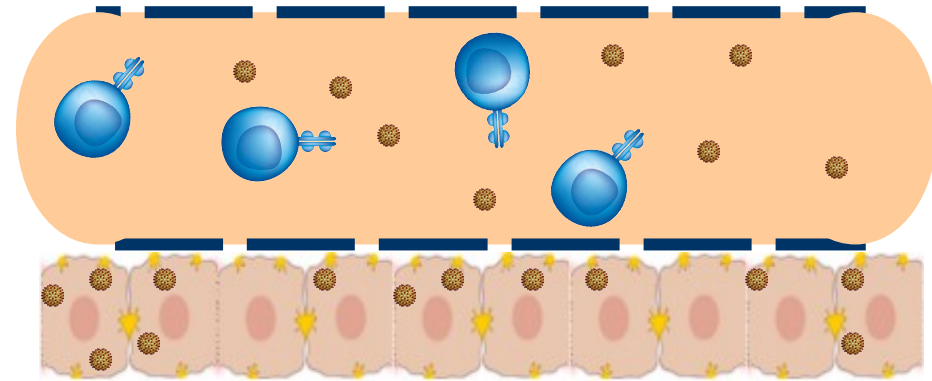


# Inhibition of HBV-specific T cell Function

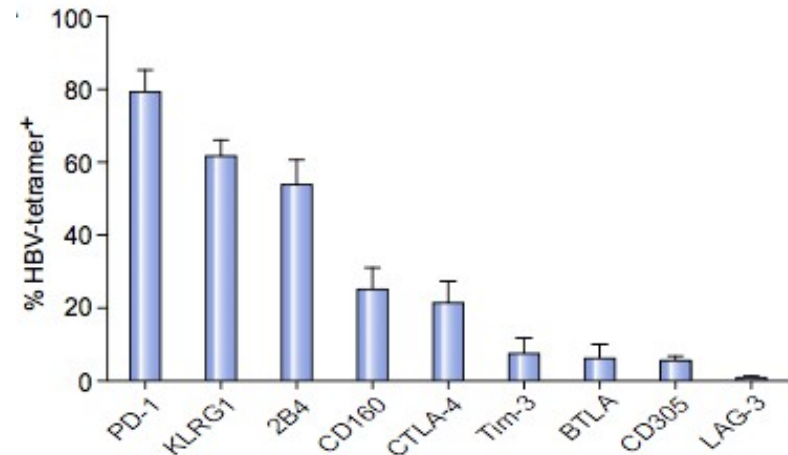
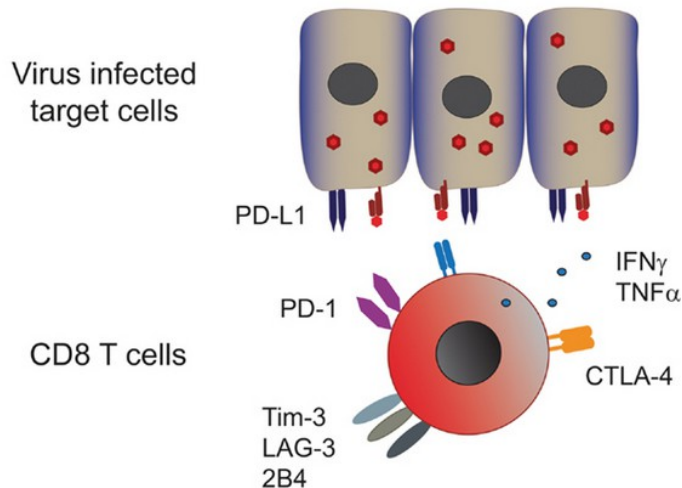
## Chronic

Weak T cell response + no effective antibodies

- 1.
- a.
- 1.
- a.
- b.
- c.



### Exhaustion

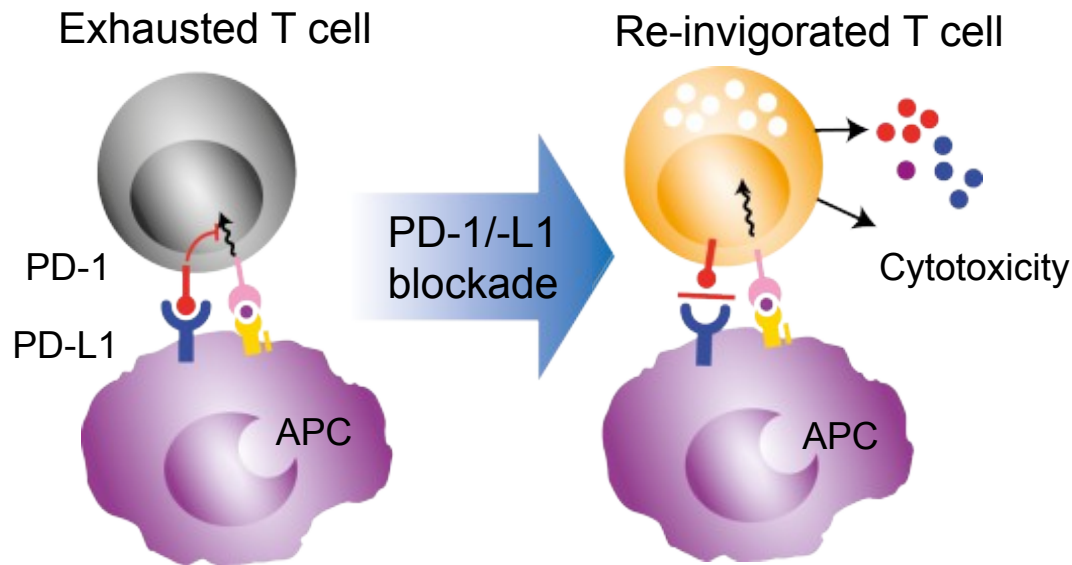


# Therapeutic vaccination with checkpoint modulation

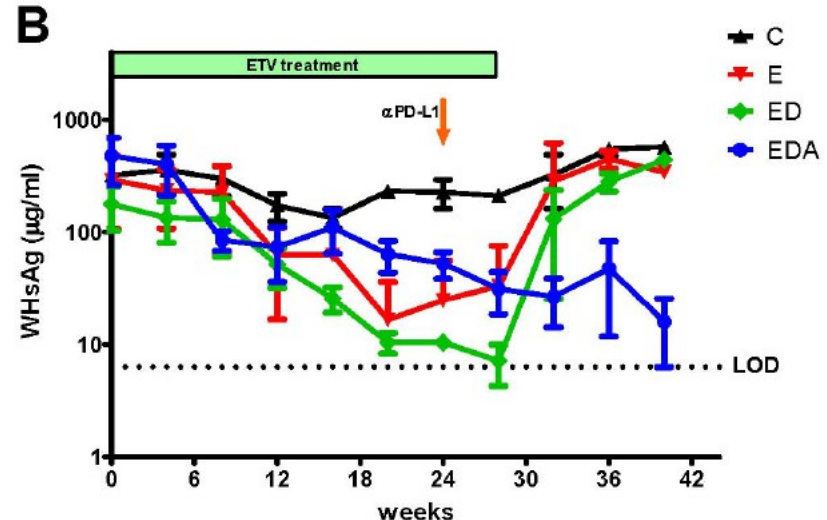
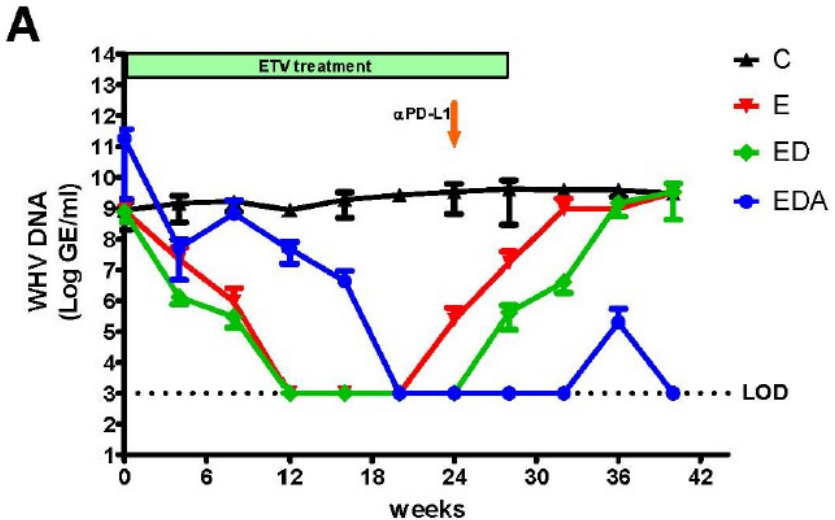
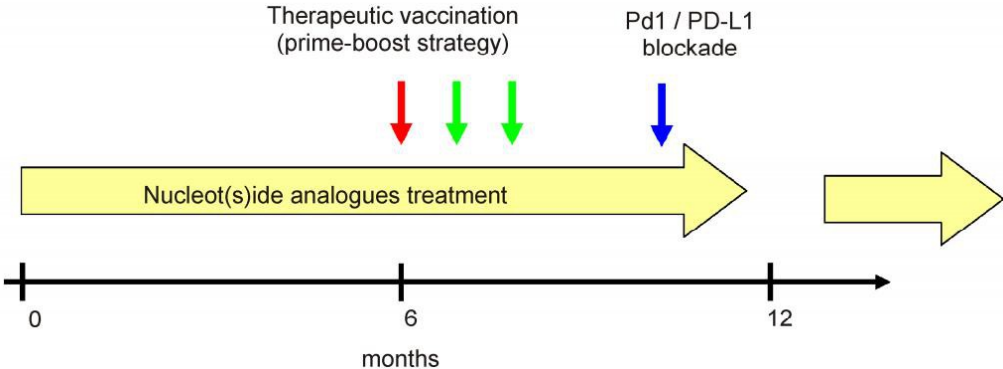
*The NEW ENGLAND JOURNAL of MEDICINE*

Safety and Activity of Anti-PD-L1 Antibody  
in Patients with Advanced Cancer

N ENGL J MED 366;26 NEJM.ORG JUNE 28, 2012



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# Stopping Treatment

## APASL Recommendation to Stop Antiviral Treatment

(Liaw, Y-F et al 2008. *Hepatol Int*;2:263)

In HBeAg-positive patients: when HBeAg seroconversion has developed > 6 months

In HBeAg-negative patients: when HBV DNA remaining undetectable for three separate occasions 6 months apart

### •Outcomes

- 25-50% develop viral relapse with hepatitis
- up to 40% remain treatment free (SVR)
- half of these lose HBsAg

### •Factors

- HBV DNA undetectable at stop
- HBsAg < 100 IU/ml [low]
- duration of AV therapy (4-5 years)

*Hadziyannis, S et al 2012. Gastro;143:629.*

*Liang, Y et al 2011. Aliment Pharmacol Ther;34:344.*

*Patwardham, N et al 2014. Aliment Pharmacol Ther;40:804.*

*He, D et al 2013. BMC Infec Dis;13:458.*

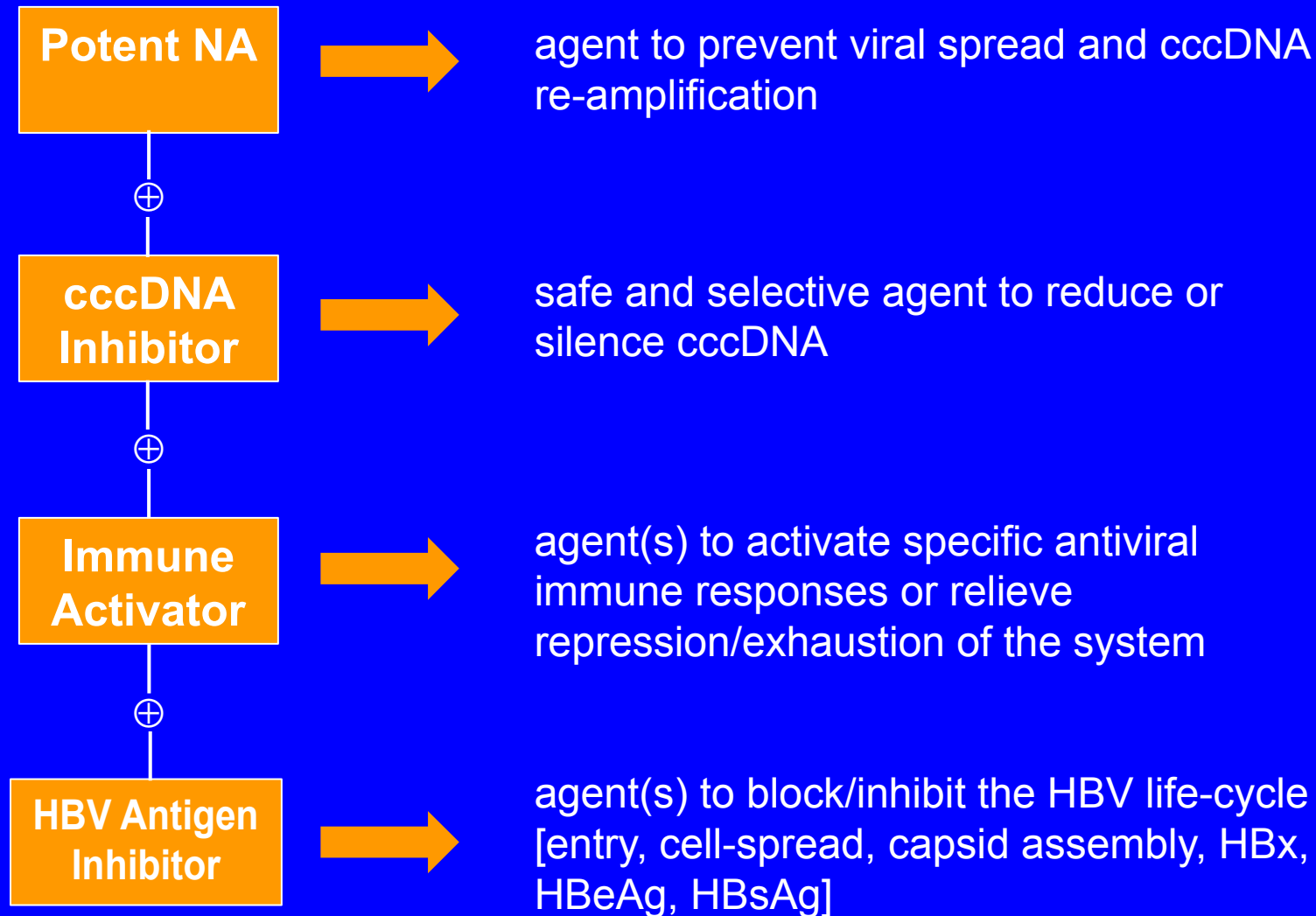
*Jeng, W-J et al 2013. Hepatol;58:1888.*

# Future Directions and Challenges

- The goalposts are shifting
- The medium-term aim for the field is to achieve functional “cure”
  - HBsAg seroconversion; HBV DNA undetectability
  - An immunomodulator is likely to be required
- New and Novel agents for CHB are starting to emerge
  - Identification of the HBV-R (NTCP) a major breakthrough
  - Improved delivery to the liver for molecular therapeutics
  - Molecular therapies blocking HBV protein/production antigen (HBsAg; HBeAg; HBx)

**PALPABLE OPTIMISM**

# What Might a HBV Curative Regimen Look Like?





# The Concept of Combination Therapy

