

Cautionary Note Regarding Forward-looking Statements

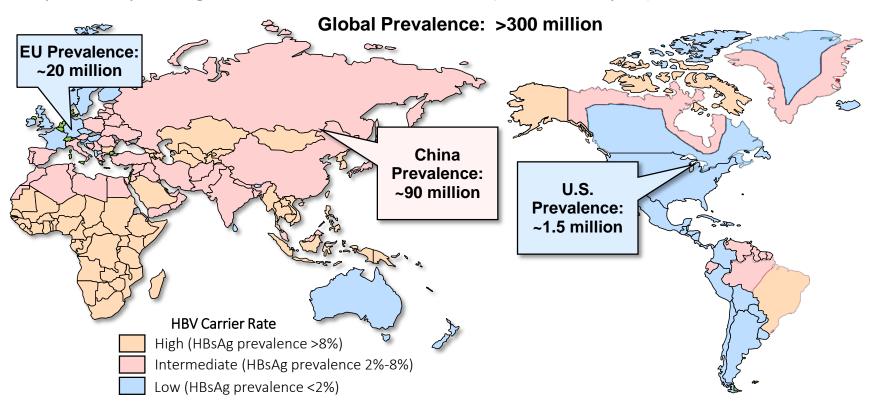


Some of the information in this presentation contains forward-looking statements regarding future events, including statements about the clinical and therapeutic potential of Assembly's development programs and drug candidates. Certain forward-looking statements may be identified by reference to a future period or periods or by use of forward-looking terminology such as "may," "expected," "should" or "predictive." Assembly intends such forward-looking statements to be covered by the safe harbor provisions contained in Section 27A of the Securities Act of 1933, as amended, and Section 21E of the Securities Exchange Act of 1934, as amended. Actual results or developments may differ materially from those projected or implied in these forward-looking statements. More information about the risks and uncertainties faced by Assembly are more fully detailed under the heading "Risk Factors" in Assembly's Annual Report on Form 10-K for the year ended December 31, 2016, and Quarterly Report on Form 10-Q for the quarter ending June 30, 2017 filed with the Securities and Exchange Commission. Except as required by law, Assembly assumes no obligation to update publicly any forward-looking statements, whether as a result of new information, future events or otherwise.

Significant Need Remains for Curative Therapies



- ❖ Number of chronically-infected HBV patients exceeds the number of patients infected with HCV (~170M) plus HIV (~37M) combined
- ❖ The majority are undiagnosed often asymptomatic for years
- Chronic HBV infection results in chronic inflammation and progressive liver damage, potentially leading to liver cirrhosis, HCC and death (~1M deaths/year)



Current HBV Therapies



Currently Approved

- Nucleoside Analogs: Entecavir, Lamivudine, Telbivudine
- Nucleotide Analogs: Tenofovir, Adefovir, Tenofovir Alafenamide
- Interferons (IFN and peg-IFN)

Entecavir and Tenofovir

- Safe, highly effective therapies and the current drugs of choice
- Target the viral polymerase, inhibiting reverse transcription of negativestrand DNA from pgRNA and positive-strand HBV DNA synthesis to generate rcDNA
- Highly effective reduction and maintenance of HBV DNA at undetectable levels in virtually all treatment-naïve patients
- HBV DNA undetectability maintained for prolonged periods (years)
- One pill, once-a-day dosing
- Very well tolerated, with no meaningful resistance emergence over prolonged treatment periods
- Unfortunately, cure rates are very low despite prolonged therapy

Deficiencies of Current Approved Therapies



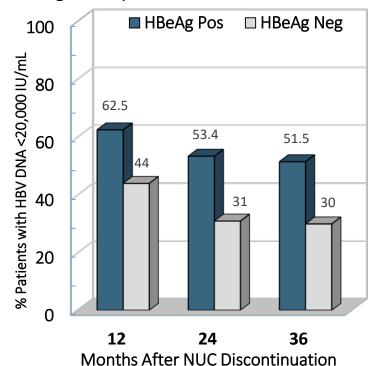
Results at 48 Weeks

HBV Patients	ETV ^{1,2}	TFV ³	Peg-IFN ^{4,5}
HBeAg Positive	N = 354	N = 176	N = 271
HBV DNA Undetectable	67%	76%	25% ^a
HBeAg Seroconversion	21%	21%	27%
ALT Normalization	68%	68%	39%
HBsAg Loss	2.0%	3.2%	2.9%
HBeAg Negative	N = 325	N = 250	N = 177
HBV DNA Undetectable	90%	93%	63%ª
ALT Normalization	78%	76%	38%
HBsAg Loss	0.3%	0%	0.6% ^b

^a HBV DNA <400 copies/mL; ^b72 weeks

Table courtesy of Geoff Dusheiko

Virologic Relapse After Nuc Discontinuation



HBeAg Positive Patients

14 studies, 733 initially HBeAg positive Pooled HBsAg loss: 1%

HBeAg Negative Patients
17 studies, 967 HBeAg negative
Pooled HBsAg loss: 1.7%

Papatheodoridis G. et al, Hepatology 2016

¹ Chang T-T, et al. N Engl J Med 2006:354:1001-10

² Lai C-L, et al. N Engl J Med 2006:354:1011-20

³ Marcellin P, et al. N Engl J Med 2008:359:2442-55

⁴ Lau GKK, et al. N Engl J Med 2005:352:2682-95

⁵ Marcellin P, et al. N Engl J Med 2004:351:1206-1

Thought We Had A Chance for Cure 12 Years Ago



Woodchuck Hepatitis B Virus Model



- WHBV infection at 3 days of age results in a carrier state with life-long viremia
- Chronically-infected animals mimic the HBV carrier state in man (viral pathogenesis & development of HCC)
- Infected woodchucks have a >90% chance of dying of HCC within 4 years
- Predictive model for toxicity and effectiveness of antivirals in man
- ETV potency against WHBV equivalent to HBV

Long-term treatment study conducted to determine if prolonged ETV therapy could cure woodchucks?

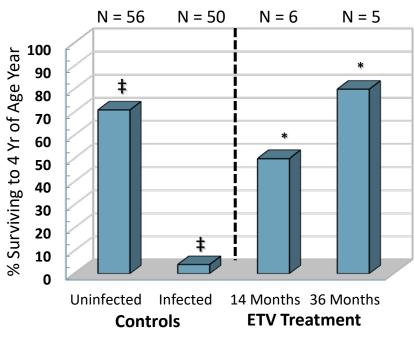
Long-Term Woodchuck Study: Cure and Survival



ETV Treatment (0.5 mg/kg)

- 2 months of daily treatment
- 12 or 34 months of weekly treatment
- Sustained suppression (≥ 8 logs) of viral DNA levels for 1-3 years – no rebounds or evidence of resistance
- cccDNA levels reduced >4 logs
- ❖ WHBsAg levels reduced 91% at Week 96
- Survival in ETV-treated animals significantly improved over historical controls
- Clear evidence that ETV can cure woodchucks based on multiple parameters

Woodchuck Survival



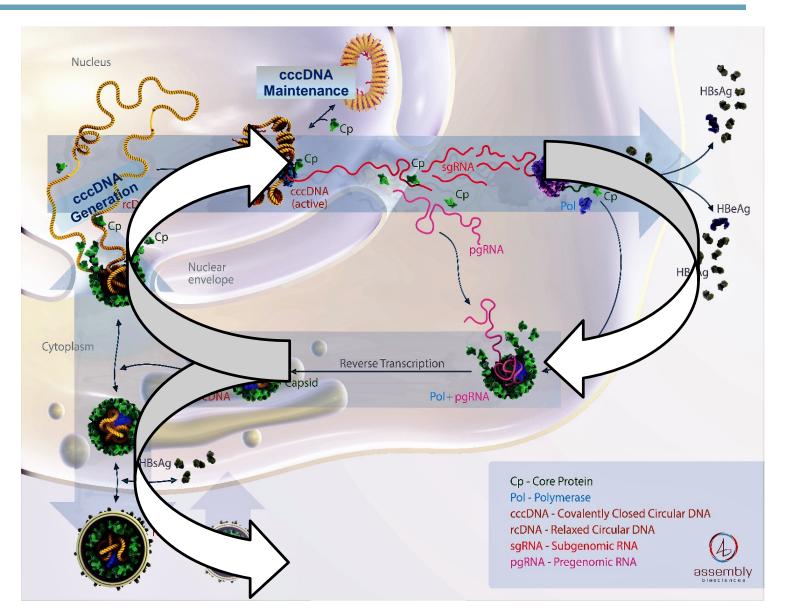
* Combined p = 0.0002

Unfortunately, this is not what happens in HBV patients with prolonged therapy

[‡]Historical control. Tennant, et al. Viral Hepatitis and Liver Disease 1988: 462-464 Colonno, et al. JID 2001;184:1236-45

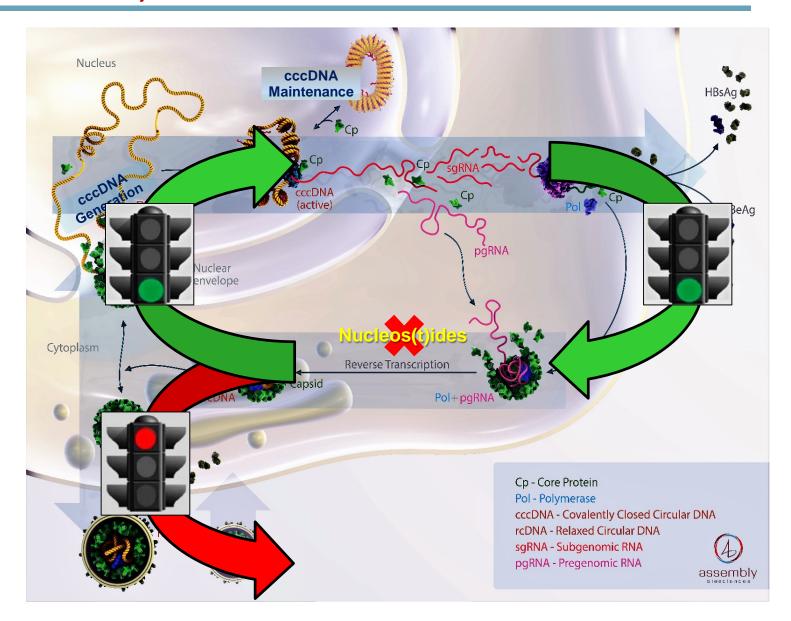
HBV Life Cycle: Complex and Poorly Understood





HBV Life Cycle: Failure of Nucs to Inhibit cccDNA





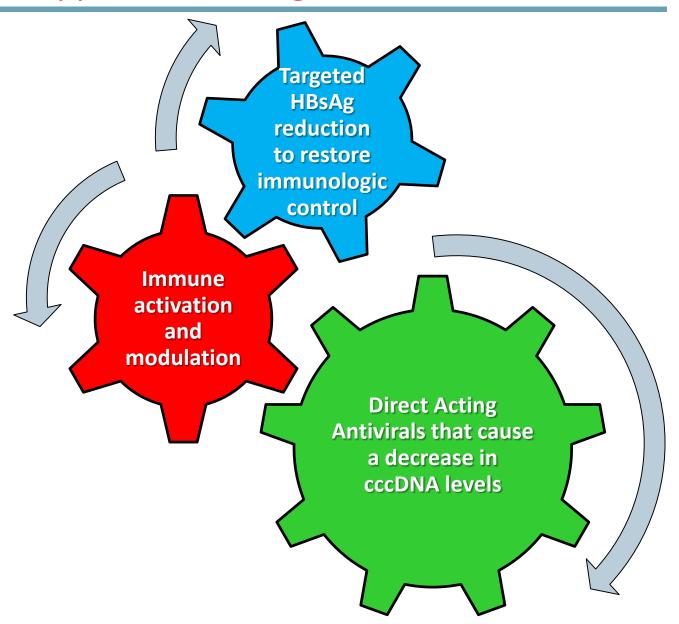
Aspirational Objectives for Clinical Cure in Humans



- We want what we achieved in woodchucks!
- Must cause depletion of cccDNA pools
 - Inhibit generation of new cccDNA
 - Direct silencing or elimination of existing cccDNA (more challenging)
- Decrease HBsAg levels to restore/enhance immune response
- Treatment of less than 2 years
 - Convenient dosing (QD?) and low pill burden
 - Excellent safety profile, with minimal side effects
- Sustained remission off therapy
 - Viral DNA replication remains undetectable
 - Elimination of cccDNA reservoirs
- Clinical efficacy
 - HBsAg loss and ideally, seroconversion
 - Reversal of liver damage, lack of hepatic inflammation
 - Significant reduction in the risk of future HCC development

Strategic Approaches Being Pursued for Cure





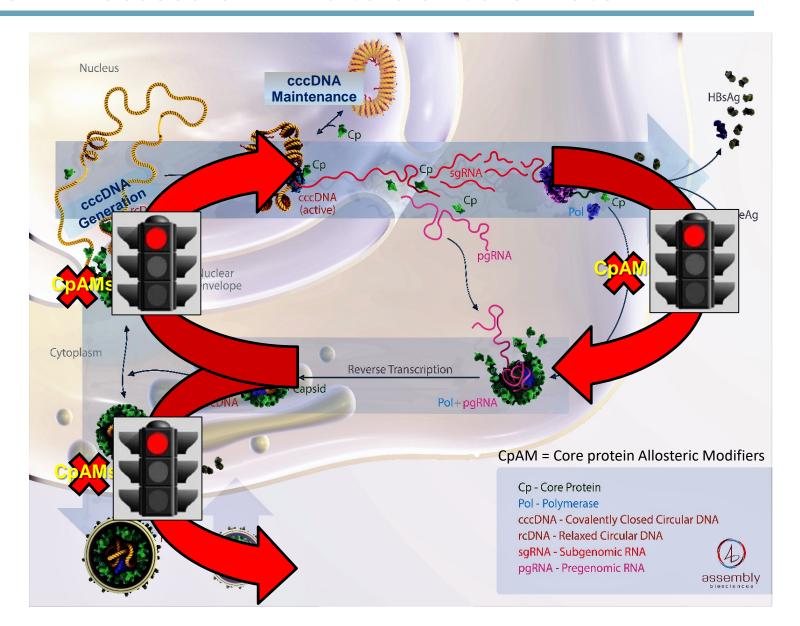


Assembly Biosciences Focused on Targeting Core Protein to Decrease cccDNA Levels



ASMB Focused on Inhibitors of Core Protein





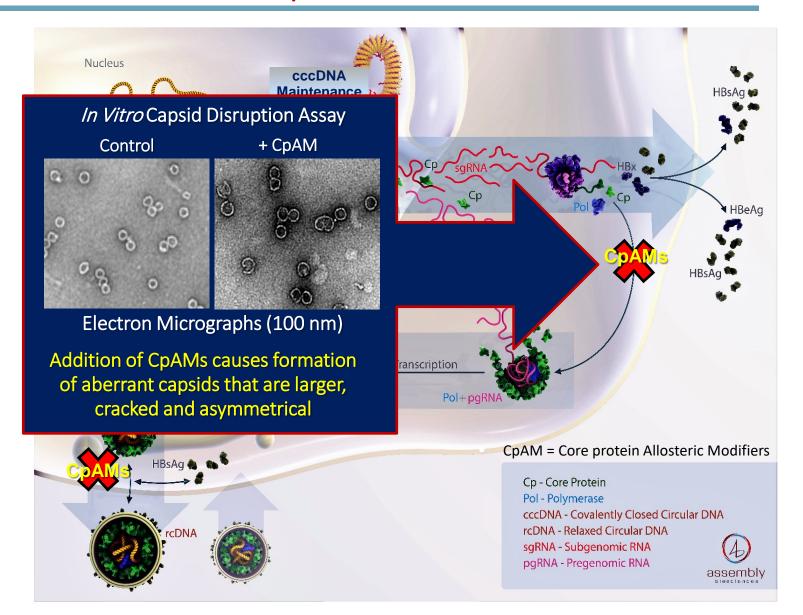
CpAMs Inhibit HBV Life Cycle at Several Steps



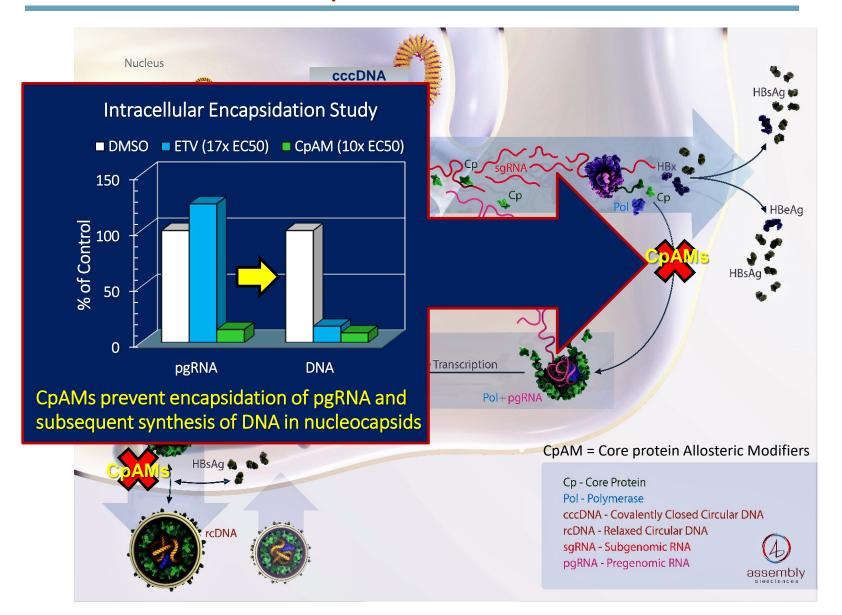
- CpAMs (Core protein Allosteric Modifiers) target Core protein and inhibit key functional steps required for to cccDNA generation
- ❖ Bind at dimer-dimer interface and disrupt the orderly folding of Core protein dimers into functional nucleocapsids, generating empty aberrant capsids
- Prevent encapsidation of Pol and pgRNA, a pre-requisite for RT activity and generation of rcDNA
- Prevent maturation of infectious viral particles
- Prevent trafficking of encapsidated rcDNA to nucleus for conversion to cccDNA
- Alter phosphorylation levels of Core protein, and may also lead to Core protein elimination in infected cells
- Ability to disrupt existing nucleocapsids

Because their distinct inhibitory mechanism(s), CpAMs and Nucs together should exhibit enhanced antiviral potency and have the potential to reduce levels of both cccDNA and HBsAg

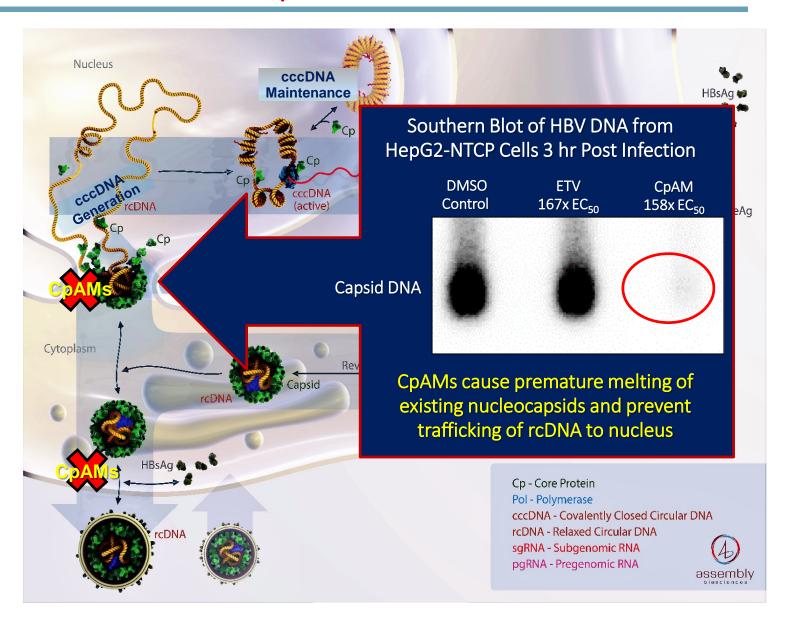




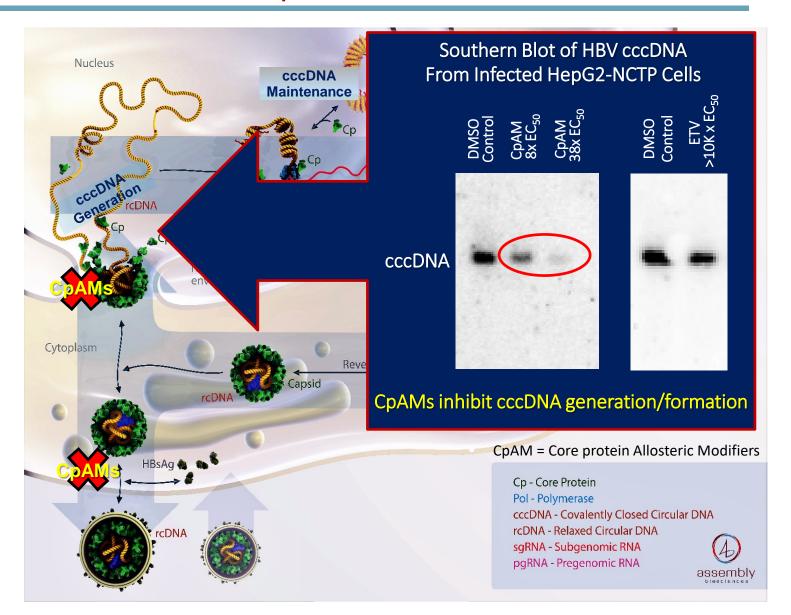








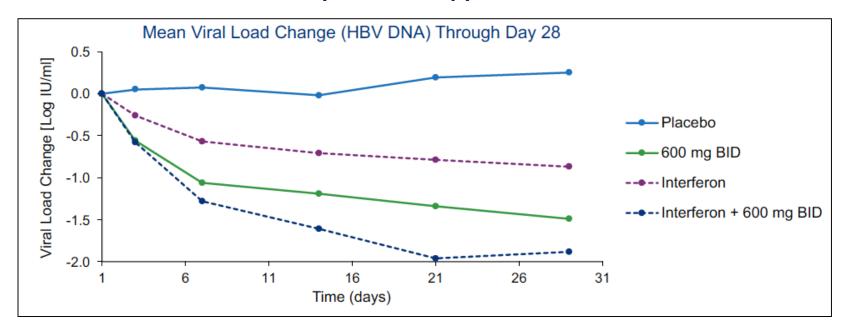




Clinical POC of CpAMs as Effective HBV Antivirals



NVR 3-778 Phase 1b 28-Day Monotherapy Results in HBV-Infected Patients



- Satisfactory safety at all dose levels, no pattern of treatment-related clinical AEs or lab abnormalities
- Oral doses of 600 mg BD in chronic HBV patients resulted in a potent antiviral response in 28 days
 - Mean 1.72 log₁₀ IU/mL HBV DNA reduction
 - Mean 0.86 log₁₀ copies/ml reduction in serum HBV RNA levels





Assembly Biosciences Establishing A Pipeline of CpAMs

Drugable Properties Will Play an Important Role



- Higher concentrations of CpAMs will be required to decrease cccDNA levels relative to inhibiting viral DNA levels
- Premium will be placed on CpAMs with favorable DMPK properties that are able to achieve and maintain higher effective concentrations in infected hepatocytes
- Convenient oral dosing (frequency and pill burden)
- Metabolic stability in hepatocytes to enable maximal sustained inhibition
- Good PK profile
 - Oral bioavailability, half-life, C_{max} and C_{min}
 - Rapid achievement of steady state
- Controlled and predictable liver:plasma concentrations, minimal accumulation in liver
- Limited drug-drug interactions
- ❖ Well tolerated good safety profile with prolonged dosing

Assembly Pipeline of CpAMs



HBV CpAM	Discovery	Lead Optimization	IND Enabling	Phase 1a	Phase 1b	Phase 2
ABI-H0731						
ABI-H2158						
Next Generation						

- ABI-H0731 has completed Phase 1a evaluation, and is currently being studied in HBV patients in a 28-day, double-blind, placebo-controlled Phase 1b study
- ABI-H2158 selected as our next generation CpAM clinical candidate and is currently undergoing IND-enabling studies for initiation of Phase 1a studies
- Plan to identify and select a third CpAM clinical candidate by year end
- All ASMB CpAMs derived from distinct proprietary chemical scaffolds unrelated to previous HAP and Novira-like molecules

ABI-H0731 Preclinical Overview



- Selective targeting of dimer-dimer interface of HBV Core protein, leading to inhibition of cccDNA generation in infected cell assays
- ❖ Pangenotypic coverage of HBV genotypes (A − D)
- No significant activity noted against a panel of other viruses or in cytotoxicity assays utilizing a panel of cell types
- ❖ Additive to synergistic in combination with Nuc
- Highly favorable DMPK properties
 - High (62-95%) oral bioavailability observed in all animal species tested
 - T½ predictive of human QD dosing
 - Limited accumulation with repeat dosing
 - Enhanced liver concentrations relative to plasma levels, parallel half-lives
 - Highly stable, excretion predominantly as intact compound
- Clean safety profile in a panel of GLP toxicology studies, no organ toxicities identified

ABI-H0731 Clinical Overview



Phase 1a Completed

- Single oral doses from 100 to 1,000 mg, multiple doses of 800 mg QD and 800 mg BID x 7 days evaluated
- ❖ Favorable PK profile with a half-life consistent with QD dosing
- Well absorbed, with achievement of concentrations believed to be sufficient to suppress viral replication and cccDNA generation
- No SAEs, no clinically significant AEs and no withdrawals due to AEs
- Treatment emergent AEs deemed "possibly related," such as headache and rash, were mild and transient, and only observed at the highest doses
- No clinically significant treatment emergent laboratory abnormalities, vital sign changes or ECG findings

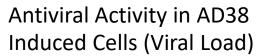
Data to be presented at AASLD, October 2017 in Washington, DC

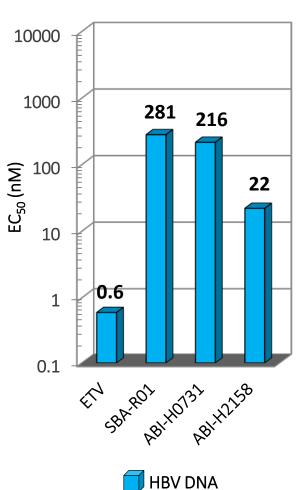
Phase 1b In Progress

28-Day monotherapy dosing in HBV patients

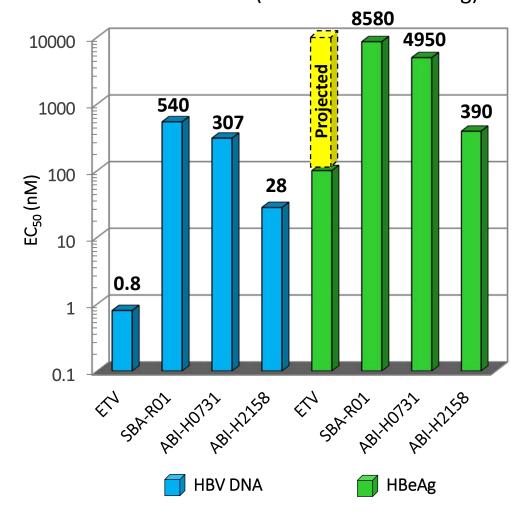
ABI-H2158 - Next Generation CpAM





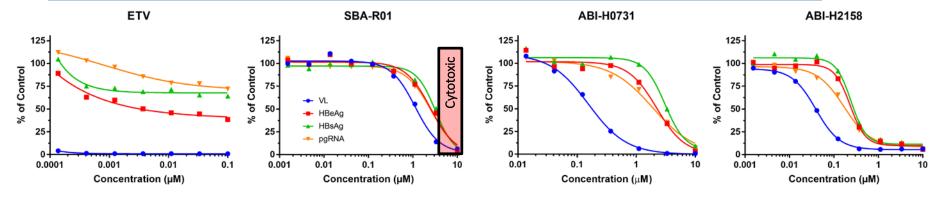


Antiviral Activity in HepG2-NTCP Infected Cells (Viral Load and HBeAg)



CpAMs Profiling in Primary Human Hepatocytes





СрАМ	HBV DNA EC ₅₀ (nM)	HBeAg EC ₅₀ (nM)	HBsAg EC ₅₀ (nM)	pgRNA EC ₅₀ (nM)
ETV	<0.1	Est >100	Est >100	Est >100
SBA-R01	1,130	2,502	3,192	2,554
ABI-H0731	154	2,210	3,000	1,840
ABI-H2158	39	230	242	169

- CpAMs reduced viral HBV DNA levels and known surrogate markers for cccDNA (HBeAg, HBsAg and pgRNA)
- ETV was highly effective at inhibiting HBV DNA levels, but exhibited limited effect on cccDNA surrogate markers

Data to be presented at AASLD, October 2017 in Washington, DC

Profile of ASMB Clinical Candidates



Virology Parameters	SBA-R01	ABI-H0731	ABI-H2158
AD38 VL EC ₅₀ (nM)	281	170	14
HC9AT HBeAg EC ₅₀ (nM)	8,580	4,950	390
PHH VL EC ₅₀ (nM)	1,130	150	23
PHH HBeAg EC ₅₀ (nM)	2,502	2,210	230
DMPK Parameters			
Human Liver Microsomes (% remaining at 45 min)	100	87	91
CYP Profile (IC ₅₀)	All > 10 μM	All >10 μM	All ≥10 μM
Protein Binding (%)	98	97	97
Rat PK %F		95	50
(1 mg/kg) T _{1/2} (hr)		6.1	2.9
C _{max} (ng/mL)		162	536
Oral AUC _{last} (hr*ng/mL)		1,470	3,671

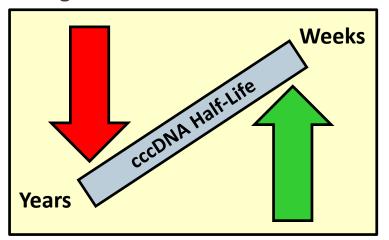
- Unclear what potency/exposure levels are required to suppress cccDNA generation
- ❖ Liver concentrations of ABI-H0731 projected to be ~25x plasma concentrations
- If needed, ABI-H2158 has superior antiviral potency to ABI-H0731, while maintaining favorable DMPK properties

Will CpAM Treatment Result in Higher Cure Rates?

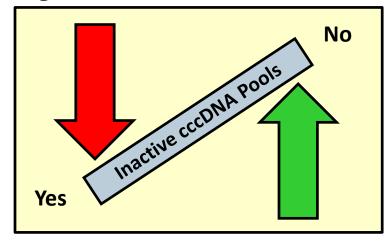


Potential Challenges

Long Intracellular cccDNA Half-life



Significant Pools of Inactive cccDNA



Studies underway using patient samples to gain insight into both topics

- Lamivudine (LVD) and Telbivudine (LdT) treatment results in rapid emergence of resistant variants due to their low barrier to resistance
- LVD/LdT resistance maps to L180M and M204I substitutions in the Pol gene
- Used these genetic markers to monitor the turnover of cccDNA, pgRNA and viral DNA in patient samples

cccDNA Studies – Initial Results



- All biological molecules have a half-life, including cccDNA
- Longitudinal study conducted on samples (serum and biopsies) from patients with emerging resistance to LVD and TBV
- Appearance and enrichment of resistant mutations used as a genetic marker in monitoring populations of viral DNA, pgRNA and cccDNA
- Results demonstrated rapid establishment of newly formed cccDNA pools harboring Nuc-resistant mutations
- Significant turnover of wt pgRNA molecules within months suggests that existing cccDNA may decay faster than previously predicted in the absence of any gross inflammation
- Little evidence for the existence and maintenance of substantial pools of inactive wt cccDNA in patient samples

Data to be presented at AASLD, October 2017 in Washington, DC

CpAM Summary



- Mechanism-based studies demonstrated that CpAMs bind to Core protein and disrupt viral replication at multiple steps
- Importantly, and distinct from Nucs, CpAMs appear to block the generation of new cccDNA molecules!
- ❖ ASMB's CpAM pipeline consists of candidate compounds selected and optimized from distinct and proprietary chemical series
- While the precise level of intrinsic potency needed for cccDNA inhibition in patients is yet to be established, emphasis has been placed on increasing potency while maintaining favorable DMPK properties
- Lead candidate ABI-H0731 has completed Phase 1a with favorable safety and PK properties predictive of QD dosing in patients, and is currently undergoing evaluation in chronically-infected patients (Phase 1b)
- Second generation candidate ABI-H2158 exhibits enhanced potency while retaining the favorable DMPK properties of ABI-H0731
- Future combinations of CpAMs and Nucs should result in enhanced antiviral activity, have a high resistance barrier and most importantly, decrease cccDNA levels

Acknowledgements



Assembly Biosciences HBV Team

Virology

Qi Huang
Dawei Cai
Ran Yan
Yi Zhou
Yuhua Zong
Alex Mercier
Pao-Chen Li
Emily Connelly
Lida Guo
Lichun Li
Esteban Carabajal
Xuman Tang

Chemistry/DMPK

Leping Li
Bill Turner
Simon Haydar
Lynn Bannen
Mark Bures
Roopa Rai
Kelvin Chan
Samson Francis

Ray Kauffman Lee Arnold

Clinical/Regulatory

Uri Lopatin Sandy Laiw Eric Ruby