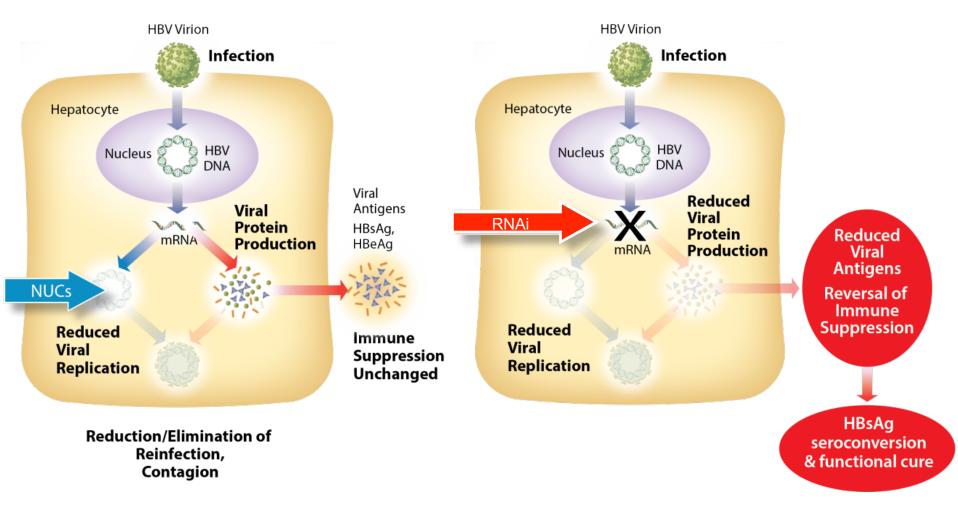
Integrated HBV DNA implicated in maintaining circulating HBsAg levels in chronically infected chimpanzees

November 15, 2015 The Liver Meeting

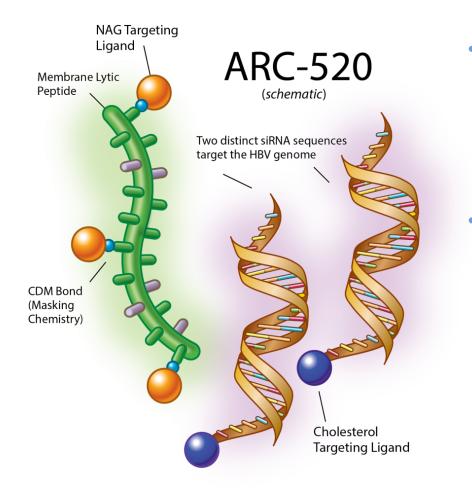
Christine I. Wooddell, Ph.D.

RNAi therapeutics vs. reverse transcriptase inhibitors (NUCs) for treatment of chronic HBV



RNA interference therapeutic ARC-520 for chronic HBV infection

Designed to reduce all transcripts from HBV cccDNA



ARC-520 Excipient

Hepatocyte-targeted
DynamicPolyConjugate[™] peptide
(NAG-MLP) to enhance siRNA
delivery

ARC-520 API

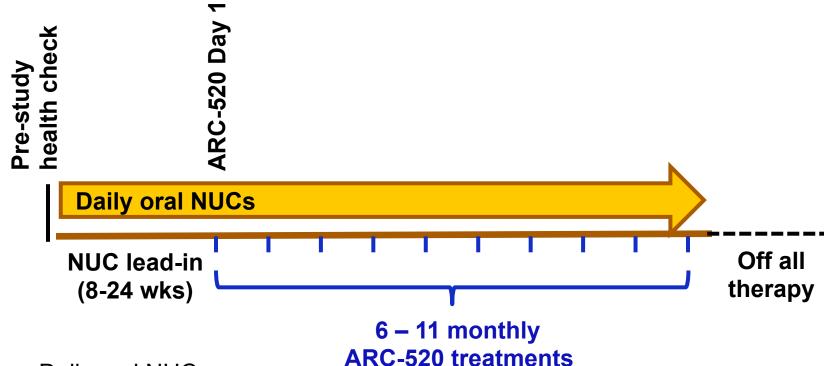
- Mixture of 2 cholesterolconjugated siRNAs
- Inclusion of two siRNAs gives broader genotype coverage (>99%)

Treatment of chimpanzees with RNAi therapeutic ARC-520

9 chimpanzees used in the study:

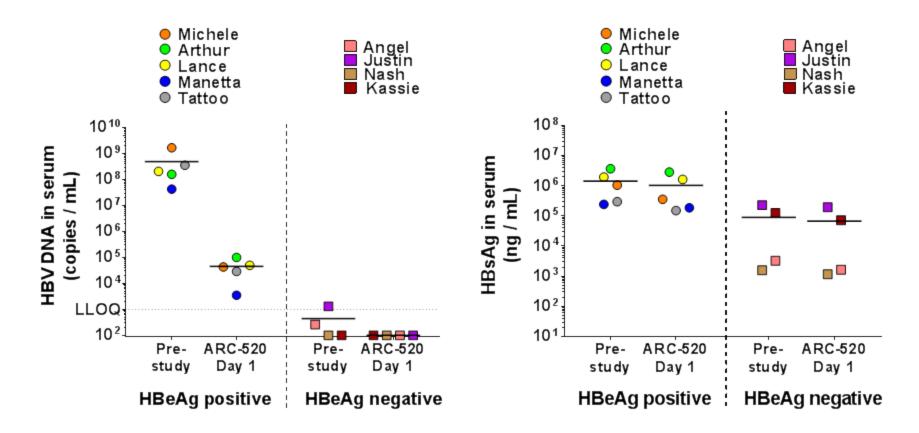
- 5 HBeAg positive and 4 HBeAg negative
- 5 males, 4 females; 9-37 years old at start of study
- Had been chronically infected many years, most from birth
- Deep sequencing and phylogenetic analysis indicated sequence was chimp variant of human HBV

Chimp dosing and sampling timeline



- Daily oral NUCs:
 - 0.5 or 1.0 mg entecavir (ETV)
 - 300 mg tenofovir added at week 15 for chimp Michele
- 2, 3 or 4 mg/kg ARC-520 intravenous injections
- Monitor safety and efficacy
 - Blood collection performed regularly throughout study
 - Periodic liver needle biopsies

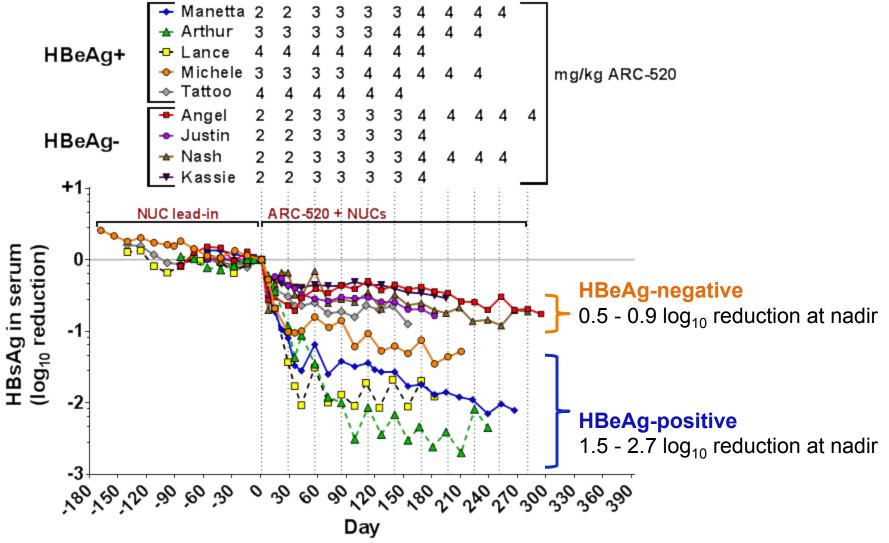
NUC lead-in: HBV serum DNA but not HBsAg decreased



Chimp response to NUC therapy similar to humans:

- Deep decrease in serum HBV DNA in HBeAg+ chimps
- Serum HBV DNA in HBeAg- chimps dropped below the LLOQ
- NUC treatment had minimal effect on HBsAg levels

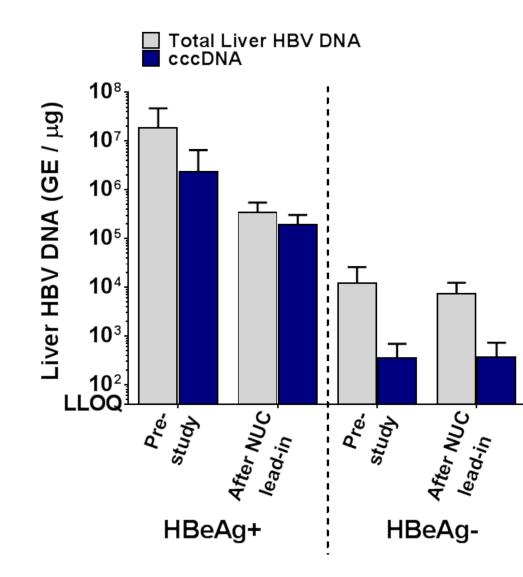
Differences in degree of HBsAg reduction correlated with HBeAg status



Analysis of HBV DNA from chimp liver biopsies

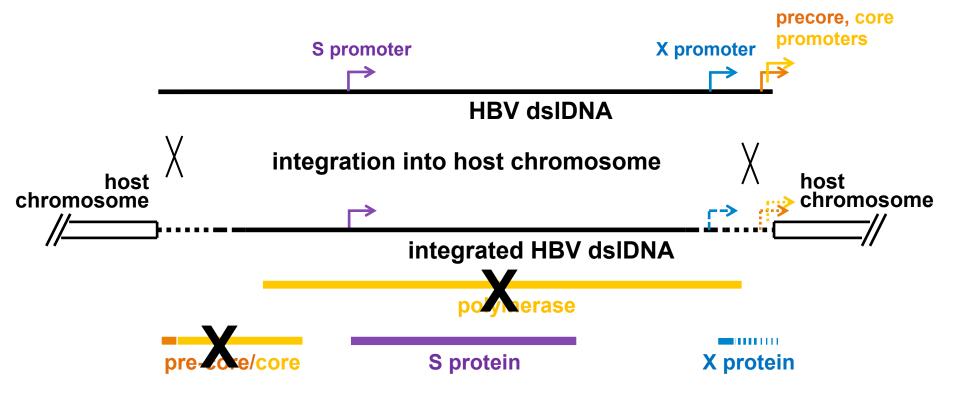
- <u>Total HBV DNA</u> (cccDNA, rcDNA and integrated HBV DNA) measured by qPCR using primers in the core gene.
- <u>cccDNA</u> measured by qPCR using same primers after digestion of rcDNA and linear (including chromosomal) DNA with Plasmid-Safe DNase.

Predominant form of liver HBV DNA differs in HBeAg- vs. HBeAg+

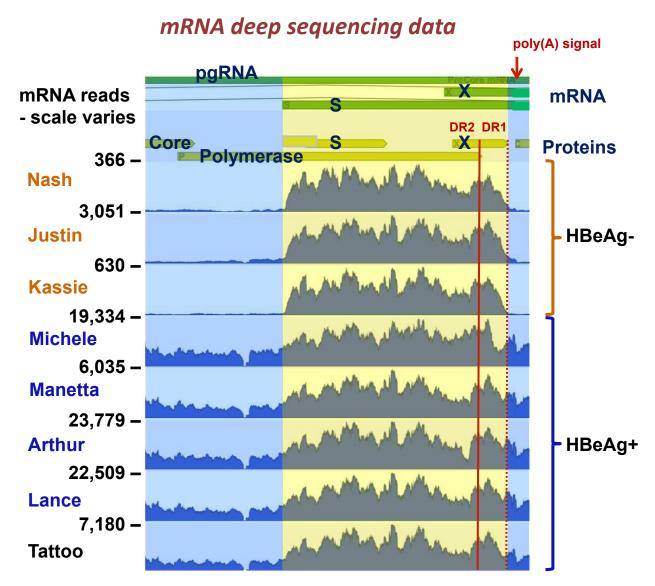


Conclusion: DNA profile in HBeAgchimps suggests most of the HBV DNA is integrated.

Process of HBV dsIDNA integration and theoretical production of HBsAg

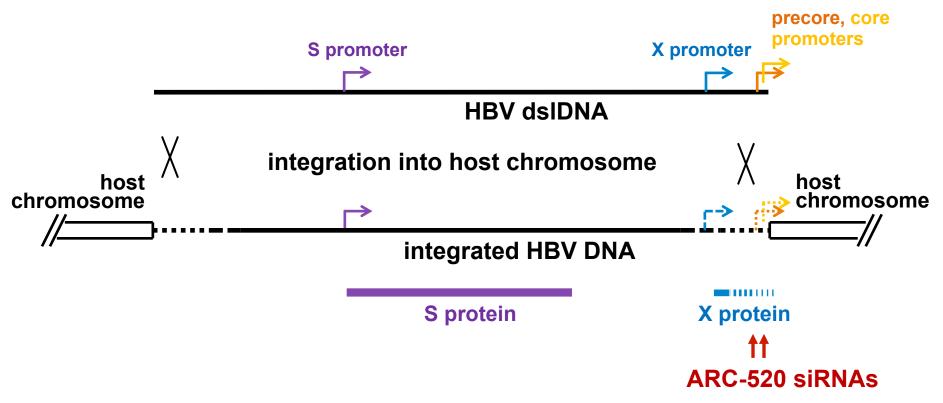


HBV transcript profiles differ between HBeAgand HBeAg+ chimps



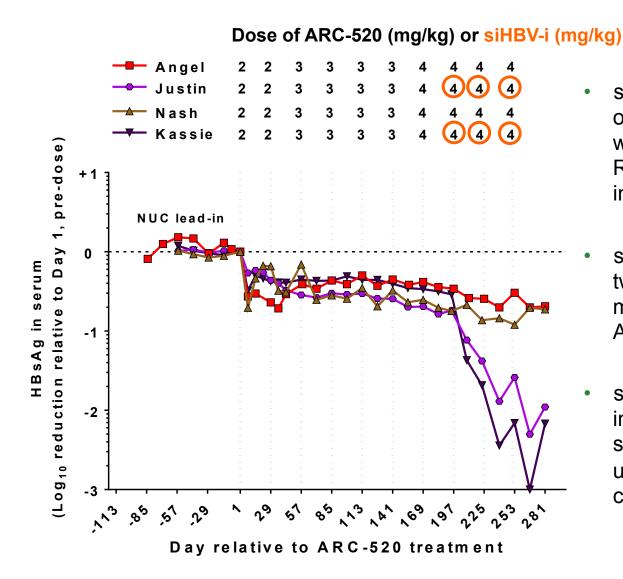
In HBeAg- chimps, HBsAg gene transcript reads are strongly reduced in a region surrounding the DR1 site.

Process of HBV dsIDNA integration and theoretical production of HBsAg



- dsIDNA integration explains persistent expression of HBsAg despite very low cccDNA in HBeAg- chimps.
- Loss of ARC-520 target sites explains lower KD of HBsAg in HBeAgchimps.

siRNA designed to target RNA derived from HBV integration products in HBeAg- chimps



- siRNA with target sequence outside of DR1-DR2 (siHBV-i) was designed to target HBV RNA expressed from integrated HBV DNA.
- siHBV-i was administered to two HBeAg- chimps once a month for 3 months following ARC-520 therapy.
- siHBV-i gave deep reductions in HBsAg in HBeAg- chimps, similar to those observed using ARC-520 in HBeAg+ chimps.

Conclusions

- ARC-520 well tolerated after multiple doses up to 4 mg/ kg (highest dose tested)
- Treatment with ARC-520 reduced HBsAg in all chimps
 - Greatest response in HBeAg+ chimps: **up to 2.7 log reduction**
 - Lower response in HBeAg- chimps: up to 0.9 log reduction
- Integrated HBV DNA is likely a significant source of HBsAg, especially in HBeAg negative chimps
 - Liver HBV DNA profiles differ between HBeAg+ vs. HBeAg- chimps
 - HBV RNA profiles in HBeAg- chimps consistent with transcripts arising from dsIDNA
- siRNA targeting integrant-derived transcripts results in deep HBsAg reduction in HBeAg- chimps

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See poster #2022 on Tuesday