



Medical Faculty Heidelberg

An update on some virological aspects of HBV/HDV Biology

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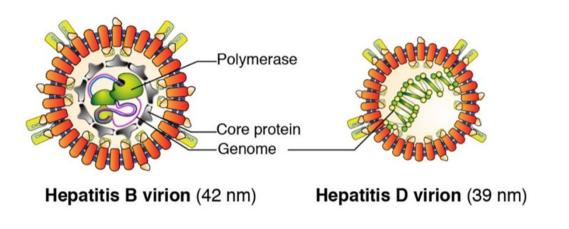


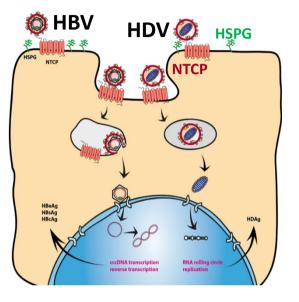
Disclosures of Stephan Urban

- OPINIONS EXPRESSED DO NOT REFLECT ANYONE'S POSITION BUT MY OWN
- ASSEMBLY BIOSCIENCES, GILEAD SCIENCES, HEPATERA-LTD, MYR-GMBH, VIRBIO
- I AM PATENT HOLDER AND INVENTOR ON PATENTS PROTECTING HEPCLUDEX/BULEVIRTIDE



HBV, HDV and their modes to establish persistent infections





- HBV and HDV share the same envelope proteins and use the same receptors
- Both viruses establish circular episomes (cccDNA, circRNA) in the nucleus of infected hepatocytes
- Maintenance of cccDNA and circRNA is crucial for persistence
- HBV cccDNA gets lost during mitosis (must be replenished by de novo entry or "amplification")
- HDV can assemble and spread using HBV envelopes encoded in integrates (no cccDNA dependence)
- RNPs of HDV- and HDV-like agents efficiently spread via cell division

What are the consequences of HBV integration on HBV/HDV infection ?



Immediate integration of HBV ds-linear HBV DNA upon infection



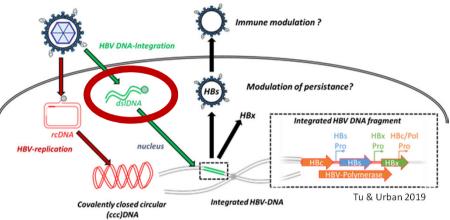
VIRUS-CELL INTERACTIONS



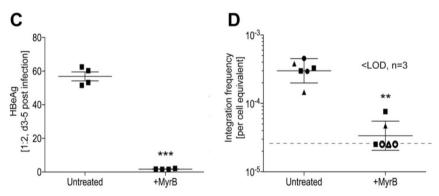
Hepatitis B Virus DNA Integration Occurs Early in the Viral Life Cycle in an *In Vitro* Infection Model via Sodium Taurocholate Cotransporting Polypeptide-Dependent Uptake of Enveloped Virus Particles

[©]Thomas Tu,^a Magdalena A. Budzinska,^b Florian W. R. Vondran,^{c,d} Nicholas A. Shackel,^{b,e,f} Stephan Urban^{a,g}





Inhibition of infection and integration by BLV/MyrB



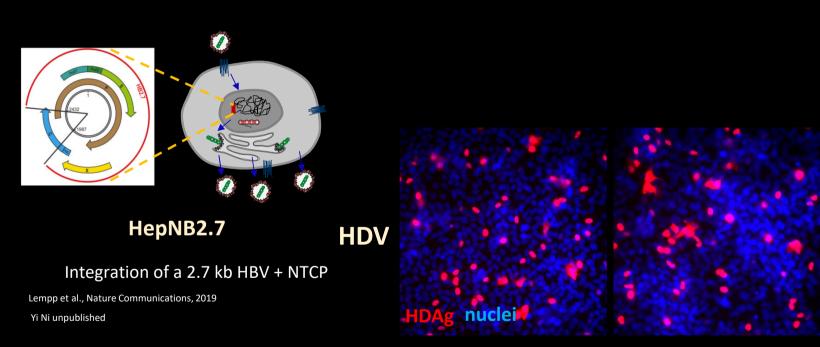
- 5-10 % of enveloped "virions" contain double stranded linear (dsl) HBV DNA.
- dsIDNA containing particles enter hepatocytes via NTCP (blocked by BLV/MyrB).
- dsIDNA instantly integrates into chromosomes.
- Integrated HBV DNA can propagate via cell division and clonal expansion.

\Rightarrow How does HBV integration affect HBV/HDV replication ?



Abrogation of HBV infection by integrated HBV DNA (encoding envelope proteins)

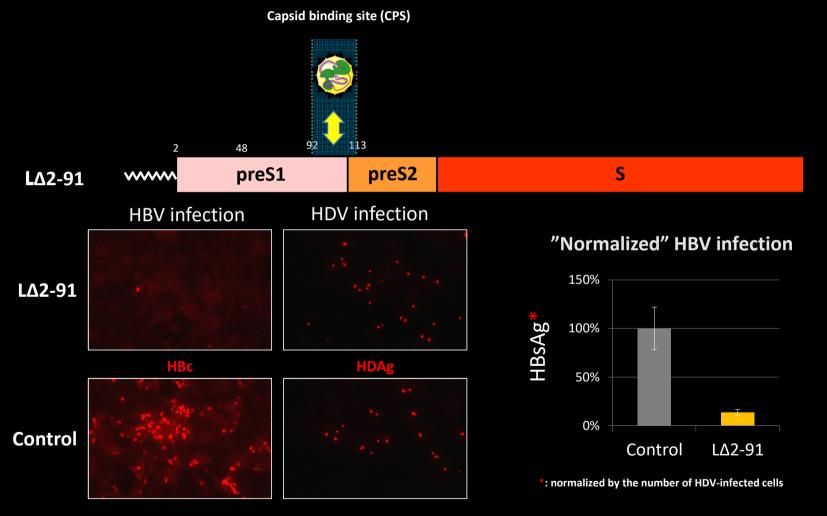
HepG2-NTCP no integrate HepNB2.7 artificial integrate



Cells expressing HBV proteins from integrated HBV-DNA support HDV infection but are resistant to HBV infection

⇒ Clonal expansion of integrants reduces HBV replication space but provides additional space for HDV replication !

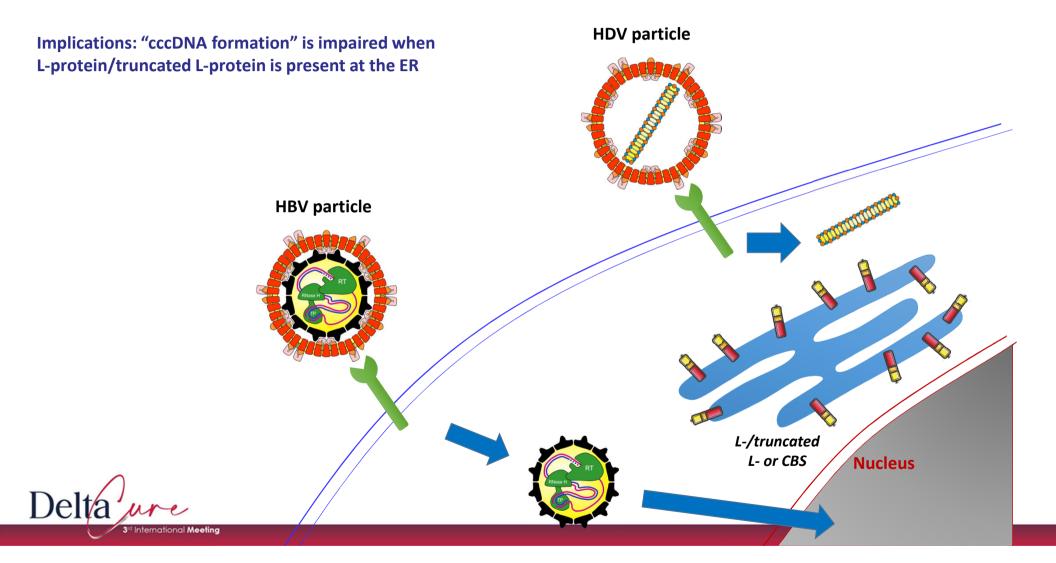
The capsid binding domain of L-protein is responsible for abrogation of HBV infection



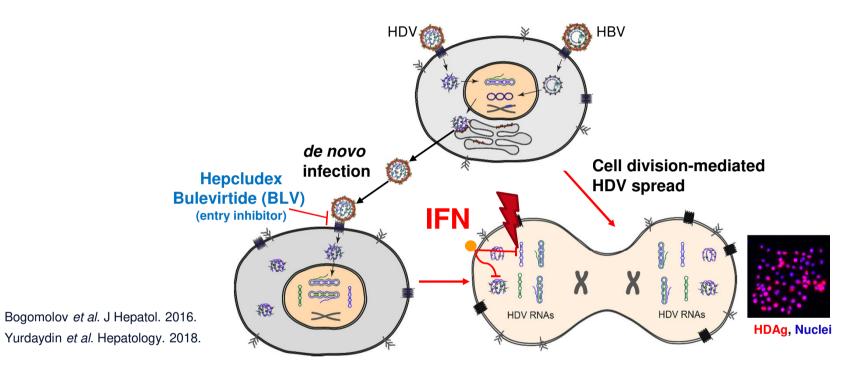
Yi Ni, unpublished

⇒ Intracellular/cytoplasmic expression of the HBV Capsid binding site blocks establishment of HBV infection

<u>The model</u>: Presence of the nucleocapsid binding site within preS prevents nuclear import of capsids and redirects both, incoming and newly synthesized NCs into the secretion pathway



HDV spreading pathways and persistence



Giersch *et al.* Gut. 2019. Zhang *et al.* J Hepatol. 2022. Zhang, *et al.* Viruses 2020.

- HBV-env-dependent *de novo* infection.
- Cell division-mediated HDV spread.

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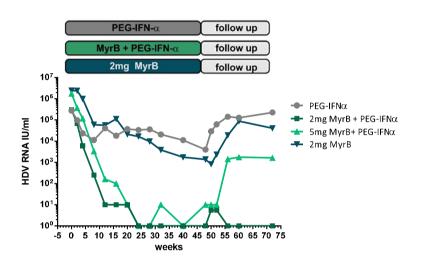
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Cell division mediated spread is inhibited by IFNs Extracellular spread is inhibited by BLV or mAbs

Therapeutic interference with both pathways result in synergistic effects (Myr-203)

Clinical findings: Myr-203 and 204 study (HDV serum RNA)

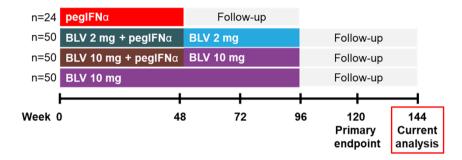
Myr-203: Median HDV RNA levels from 15 patients



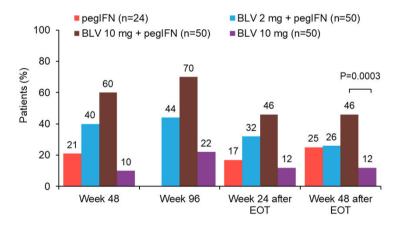
Strong synergism of IFN α and BLV on HDV serum RNA

Wedemeyer, et al. EASL ILC 2020.

Myr-204: study design



Undetectable HDV RNA through 48 weeks after EOT

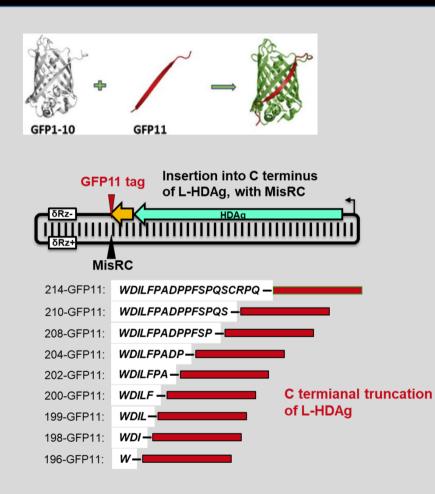


46% of patients remain HDV RNA negative 48w after EOT (10 mg BLV + pegIFN α)

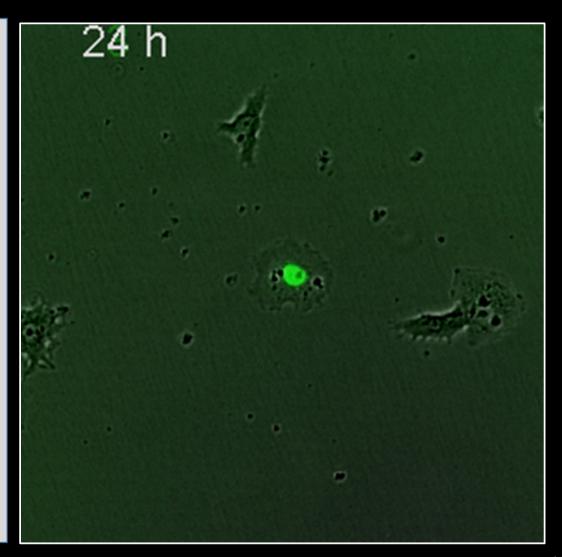
Delta une 3ª International Meeting

Asselah T, et al. EASL ILC 2024. GS-002.

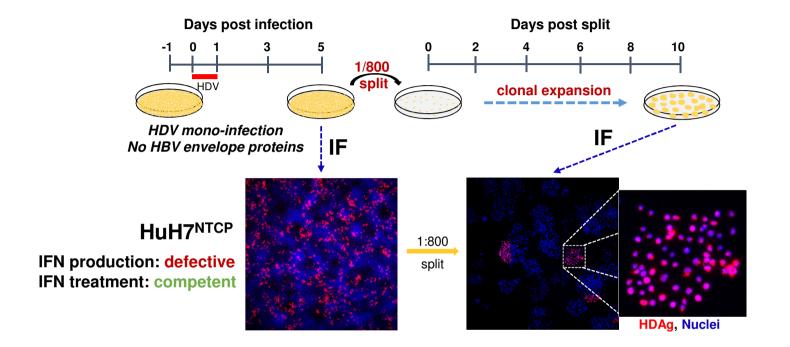
Live cell imaging of HDV-spread by cell division



Zhenfeng Zhang, unpublished



Efficacy of cell-division-mediated HDV spread depends on the innate immune competence of cells

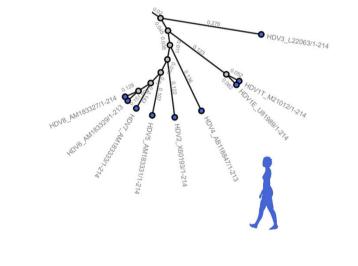


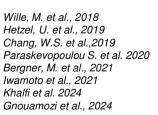


HDV evolution and discovery of HDV-like agents

HDV and HDV-like agents (DLA)

- Encode homologous small delta antigens (SDAg)
- No (farnesylated) large delta antigen expression
- No evidence for hepadnaviral helper function / no liver tropism





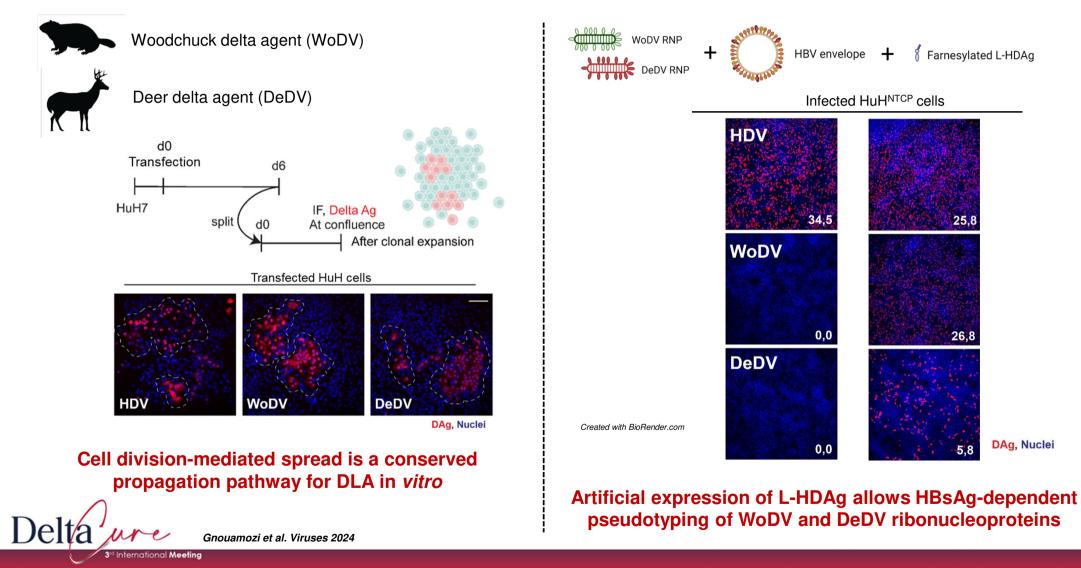
Replication, spreading pathways and host counteractions are still unknown

Primate

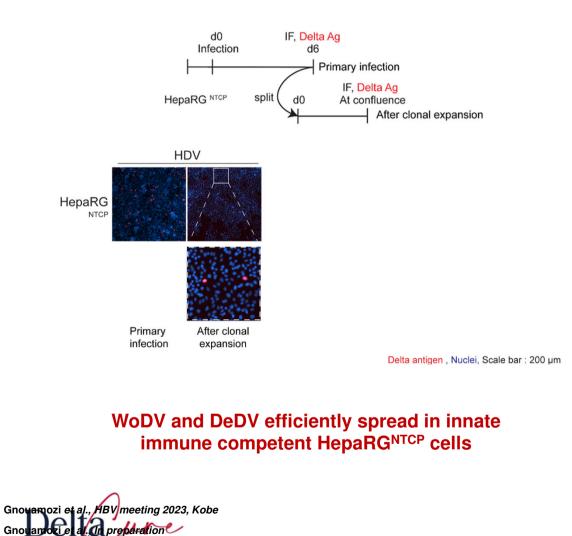


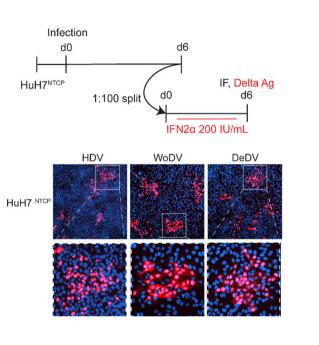
Gnimah Eva Gnouamozi

Woodchuck and Deer delta-like agents: persistence and pseudotyping



Effect of intrinsic and exogenous interferon stimulation on spread

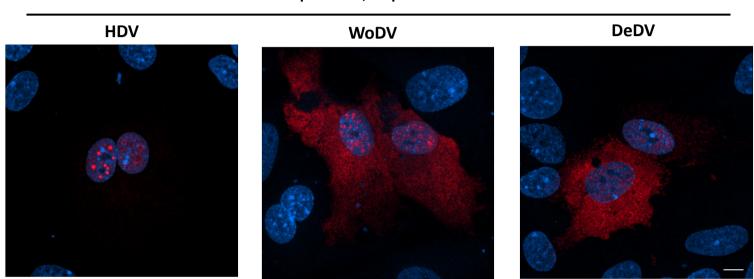




Delta antigen , Nuclei, Scale bar : 200 µm

IFN does not inhibit spread in HuH7^{NTCP} cells

Viral antigen subcellular localization in infected HepaRG^{NTCP} cells



HepaRG^{NTCP}, d6 post-infection

Delta antigen Nuclei Scale bar: 10 µm

- The HDV HDAg exhibits predominantly nuclear staining in infected cells
- WoDV and DeDV DAg are also widely distributed throughout the cytoplasm

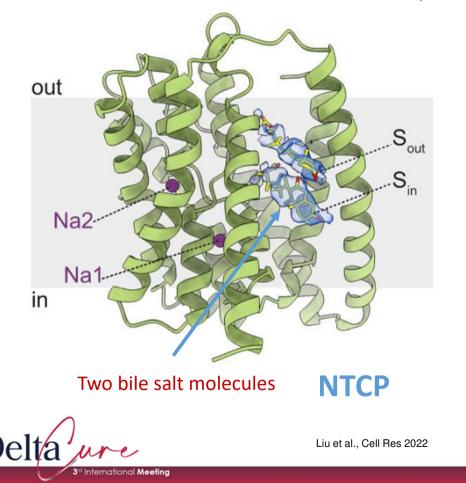
A possible role for DAg in counteracting the innate immune system.



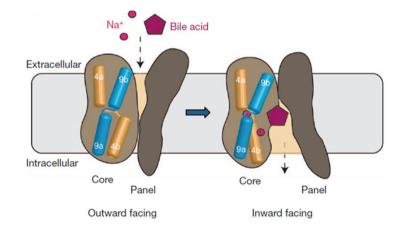
Structure model and CryoEM-structure of the HBV/HDV receptor NTCP (SLC10A1)

Model of the structure of **sodium Taurocholate Co-transporting Polypeptide NTCP**:

with the location of a double bile salt pocket

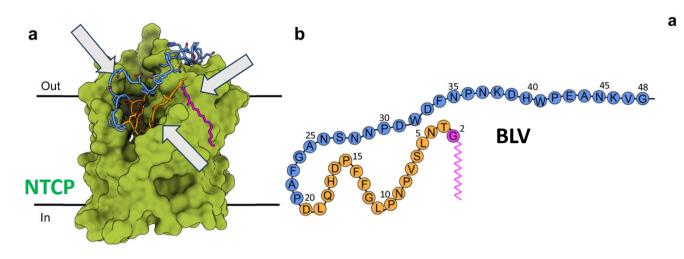


- Integral transmembrane protein (9 TM helices)
- Transports bile salts through PM into hepatocytes
- Conformational change driven by sodium Ion gradient



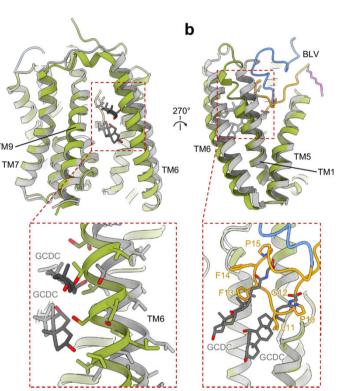
Goutam et al., Nature 2022; Park et al., Nature 2022; Asami et al., Nature 2022; Liu et al., Cell Res 2022

Cryo-EM structure of bulevirtide/Hepcludex bound to NTCP



- Bulevirtide: myristoyl anchor, plug domain and string domain
- The myristoyl anchor locates sidewise intruding the lipid bilayer and positioning Glycin-2 at the external entrance of the tunnel.
- The **plug domain (Gly2-Asp20)** wedges inside the translocation tunnel, deeply intruding the protein by making several specific contacts.

• The string domain (Pro21-Gly48) covers the surface of the plug and bridges the extracellular surface of NTCP ("clasping" the molecule)



Superposed BLV-bound NTCP (green) and substratebound NTCP (gray).

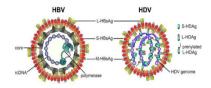
Binding of BLV to NTCP displaces the core alpha helices

- \rightarrow Resulting in a shift of the panel domain.
- \rightarrow blocking bile acid and sodium ion binding.



Liu et al., Nat Communication 2024

Summary, conclusions, take home messages



- HBV and HDV share the same envelope proteins which are provided by HBV integrates or cccDNA.
- HBV DNA readily integrates into host chromosomes and mediates superinfection exclusion of HBV but not HDV
- Expression of envelope proteins restricts the replenishment of cccDNA by preventing NC-(re)import
- HDV genomes disseminate by an extracellular route (intrahepatic and between hosts) and by cell-division mediated spread (intrahepatic).
- Cell division mediated spread of HDV but not WoDV and DeDV are sensitive to IFNs.

Clinical implications:

In addition to entry inhibition with BLV (or upcoming neutralizing antibodies), acceleration of HDV clearance by inhibitors that prevent celldivision mediated spread may allow elimination of HDV even in the presence of HBsAg.



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3rd International Meeting

MOLECULAR VIROLOGY H E I D E L B E R G

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