



OP-0682

Characterization of circulating Hepatitis B virus RNAs *in vitro* and chronic hepatitis B patients

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Lyon 1

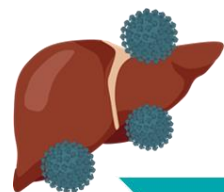
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New biomarkers to HBV cure



Therapy & Cure definition

Partial cure → Functional cure → Complete cure → Sterilizing cure



cccDNA

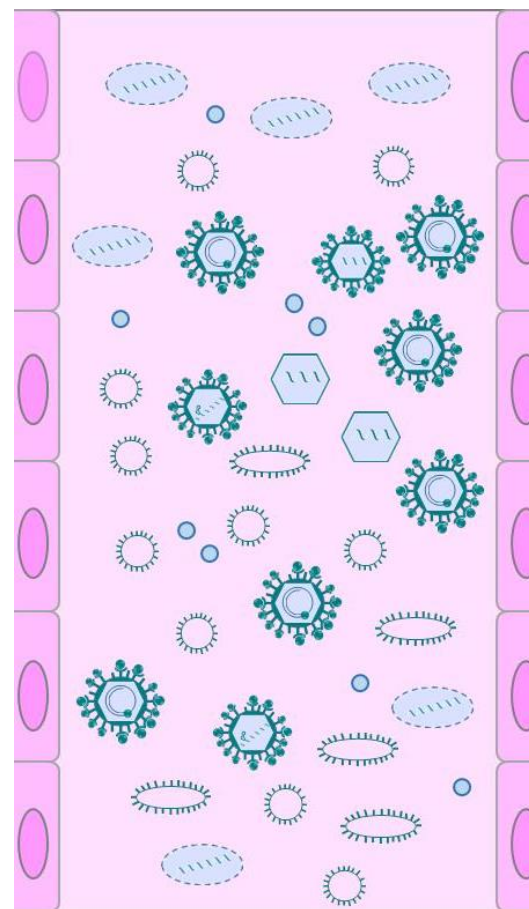


HBV DNA integration



Ideal HBV biomarker?

- Non-invasive
- Reflects intrahepatic cccDNA pool and activity
- Predicts “HBV cure”



CHB patient Bloodstream

Vehicle components in the bloodstream

Classical biomarker



HBV DNA



HBeAg



HBsAg

Circulating HBV RNA (CirB-RNAs)



Virion Like-Particles

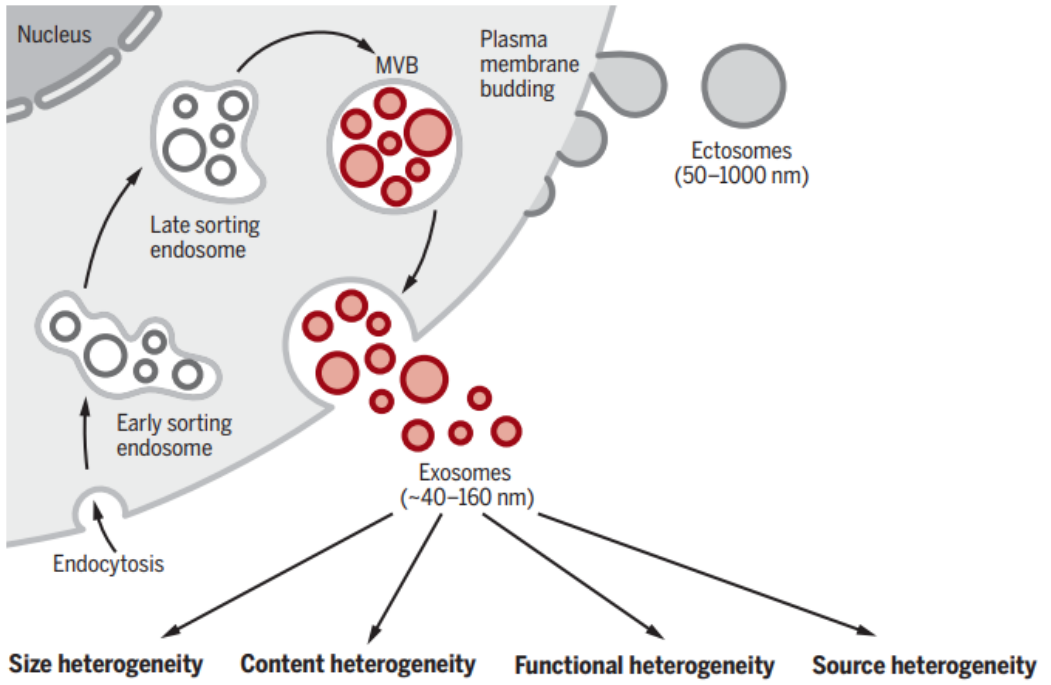
(Jie W, J Hepatol, 2016)



Extracellular Vesicle



Extracellular Vesicles (EVs): Exosome



EVs

- Role: Cell to Cell communicator
- Cargo: DNA/RNA, protein, and miRNA
- Diameter: 40 - 160 nm
- Marker: CD9 and CD81

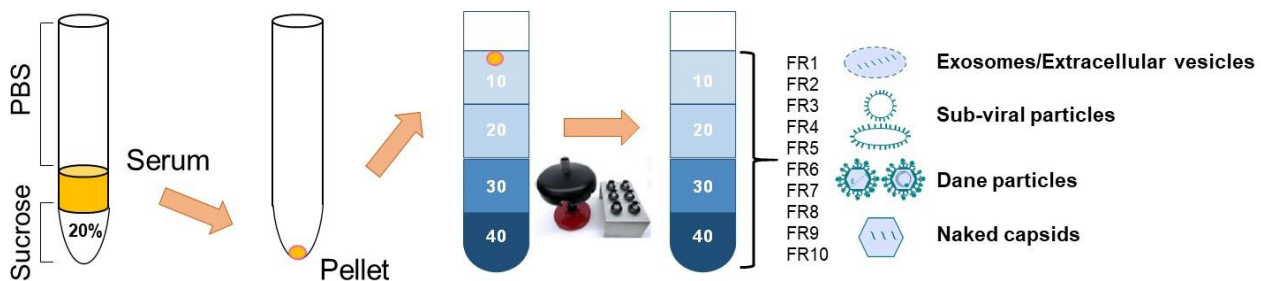
Kalluri et al, Science, 2020

Trojan EVs hypothesis

The Trojan EVs hypothesis is illustrated with three colored circles and a central image of a Trojan horse. The red circle represents **HBV DNA/RNA spread and pathogenesis**, supported by references: *Yang et al, Cell & Mol Immunol 2016*, *Sanada et al, CMGH 2017*, *Kapoor et al, Virus Res 2017*, and *Shuang et al, Biomed Res Int. 2019*. The blue circle represents **IMMUNE system regulation**, supported by *Masatoshi et al, Plos One 2018*. The green circle represents **Chemoresistance**, supported by *Liu et al, Oncol Lett 2019*. The central image shows a Trojan horse, symbolizing the delivery of viral components into cells via EVs.



Method: Density gradient ultracentrifugation and Patients' Information



Experimental Method

1. Concentration: Ultracentrifugation for 5 hours at 35000rpm
2. Density gradient ultracentrifugation for 16h at 35000rpm
-10-40% Iodixanol/Sucrose
3. Total of 10 fractions, 500ul each

Patients' Information

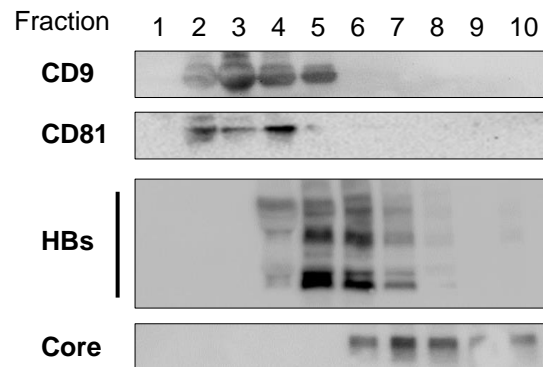
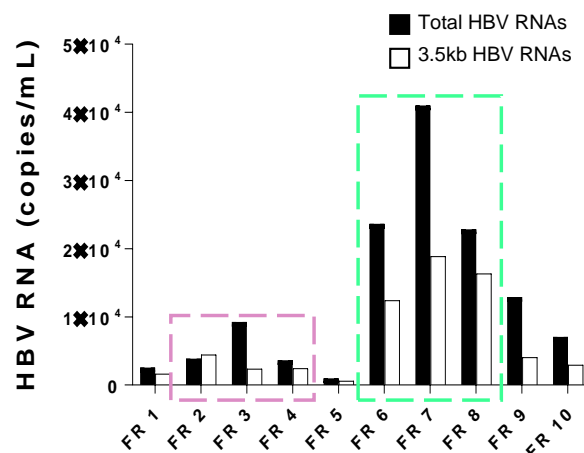
| Sample Name | HBV load (log IU/ml) | HBeAg | HBsAg (log IU/ml) | ALT | Anti-HBV Treatment | CHB phase |
|--------------------|----------------------|-------|-------------------|-----|---------------------|-----------|
| Patient 1 | 8.5 | (+) | 81000 | 36 | NO | HBeAg+ CI |
| Patient 2 | 5.6 | (-) | 570 | 158 | NO | HBeAg- CH |
| Patient 3 sample 1 | 8.3 | (+) | 37000 | 201 | NO | HBeAg+ CH |
| Patient 3 sample 2 | 3.1 | (+) | 270 | 27 | Tenofovir 6 months | |
| Patient 3 sample 3 | <LLoQ | (+) | 240 | 24 | Tenofovir 10 months | |

According to EASL CPG 2017

Where are CirB-RNAs distributed in CHB patients?

A. Patient 1

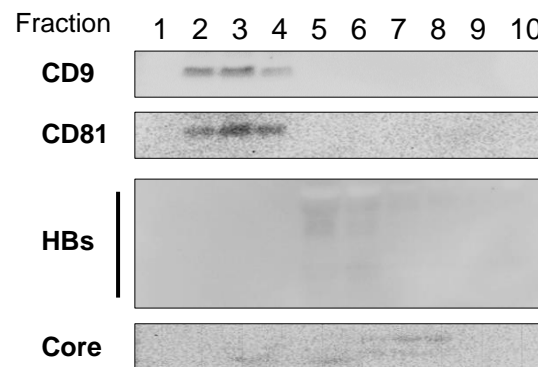
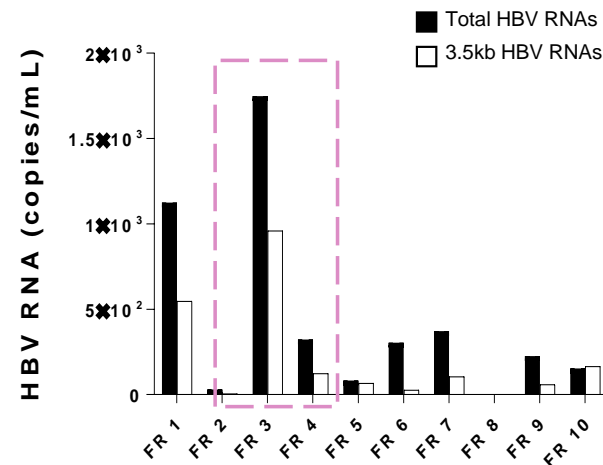
HBeAg(+), high viral load



CirB-RNAs are detected in EVs (FR2-4) and VLPs (FR6-8) fractions.

B. Patient 2

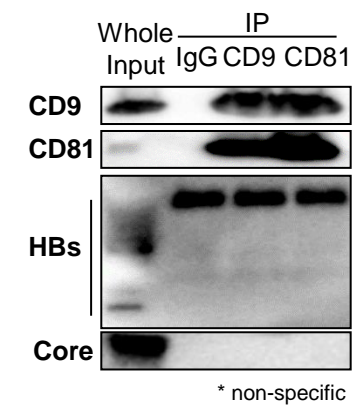
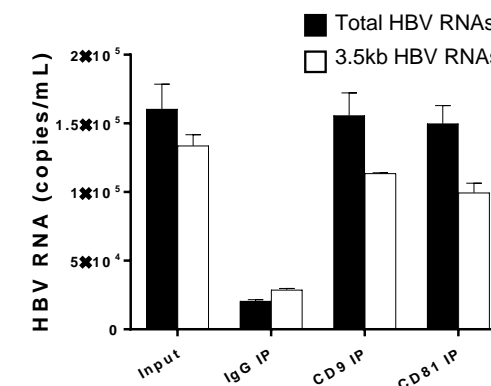
HBeAg(-), low viral load and HBsAg



CirB-RNAs are mostly detected in EVs (FR2-4) fractions.

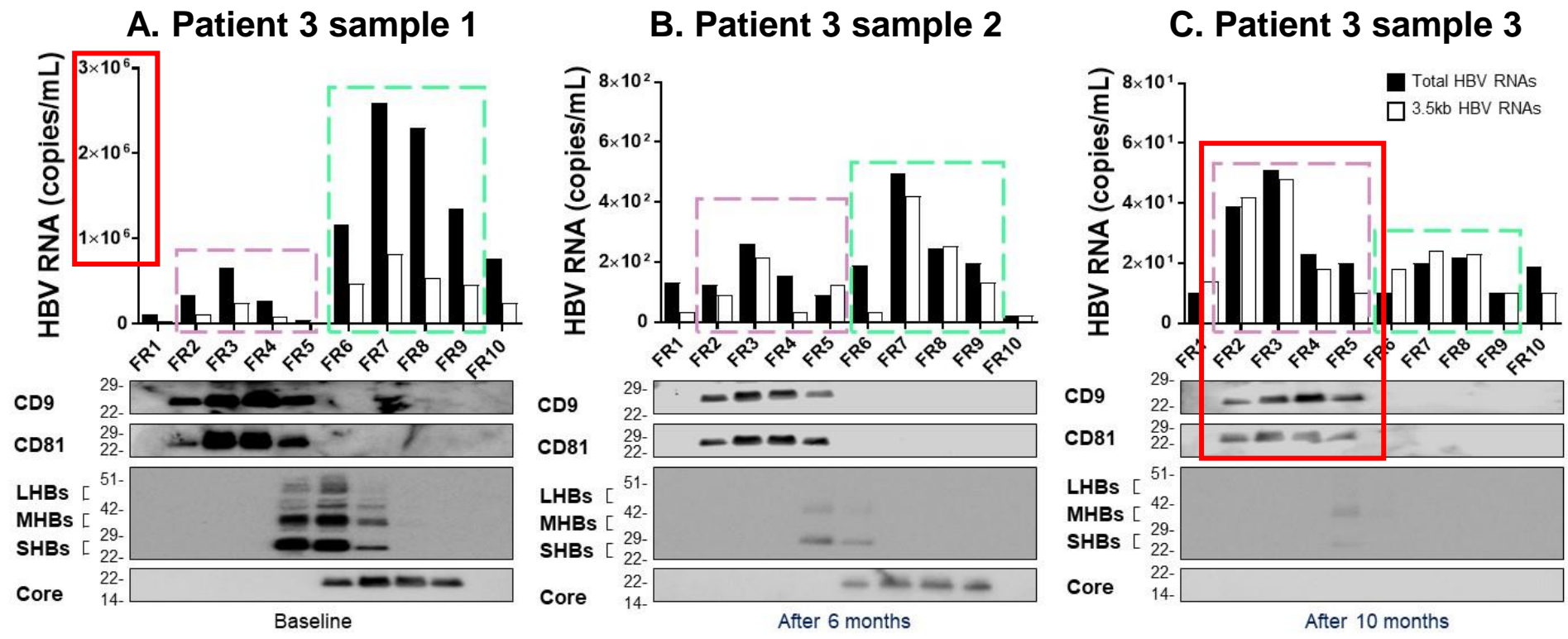
C. Patient 1

IP assay



CirB-RNAs are detected in EVs.

Longitudinal study of CHB patient's treatment with TDF

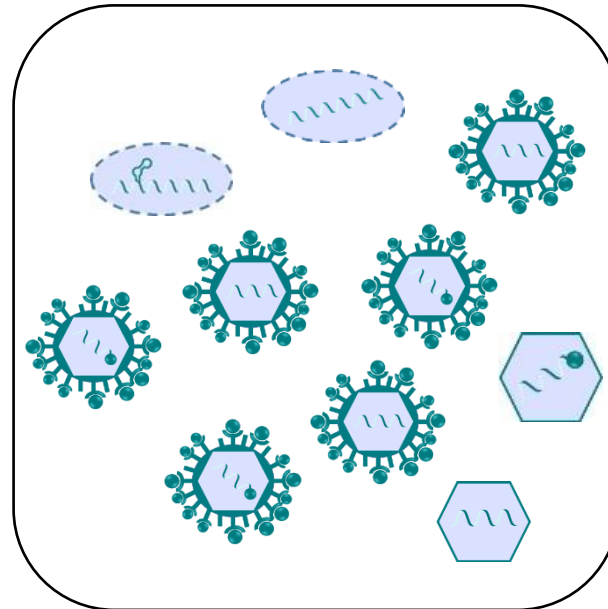


the distribution of cirB-RNAs changed from VLPs to EVs during TDF treatment

Conclusions

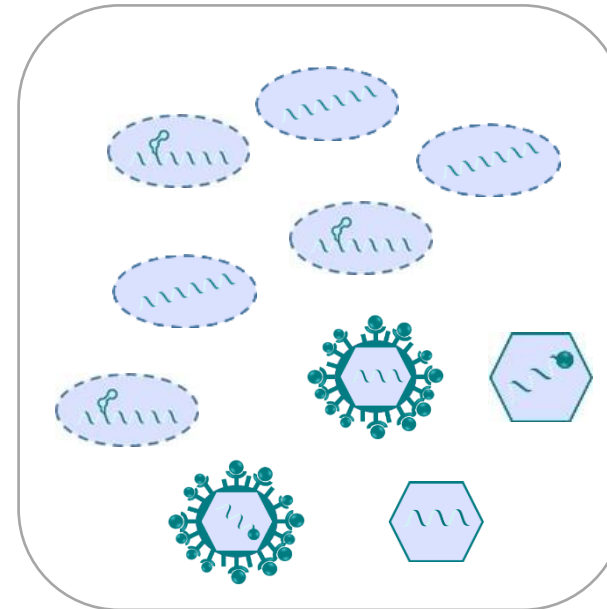
High viremia

HBeAg (+), High viral load,
High HBsAg



Low viremia

HBeAg (-), Low viral load,
Low HBsAg



Altogether, our results significantly contribute to the characterization of **cirB-RNAs** as **new viral biomarker**

ACKNOWLEDGEMENTS



INSERM U1052- RHU's Team

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