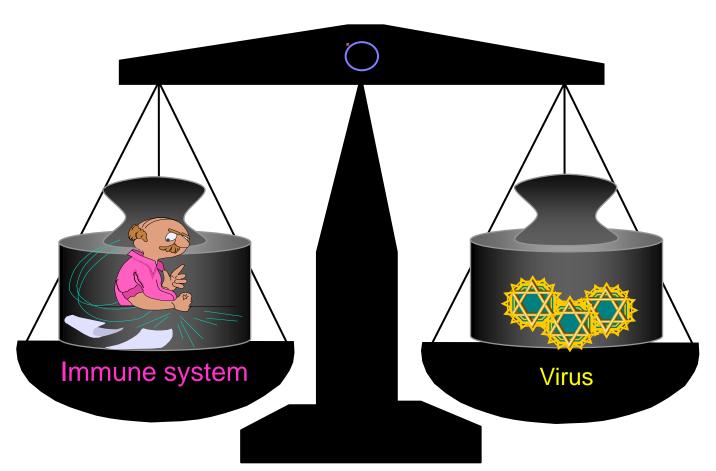
# Epigenetics of liver diseases: potential clinical application of the study of serum miRNA profiling in chronic carriers of Hepatitis B Virus

Maurizia Rossana Brunetto

UO Epatologia - Centro di Riferimento Regionale per "la Diagnosi e il Trattamento delle Epatopatie Croniche e del Tumore di Fegato" Azienda Ospedaliero Universitaria Pisana - Pisa-Italy

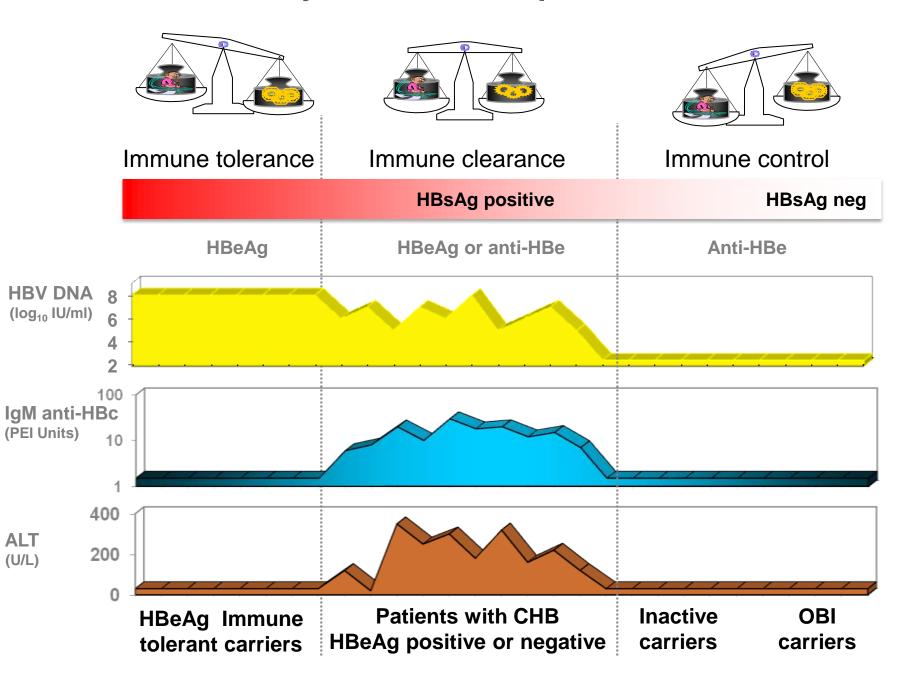
# Outcome of HBV infection and liver disease



A coordinated humoral and cellular immune response may control HBV infection, allowing its persistence without liver damage

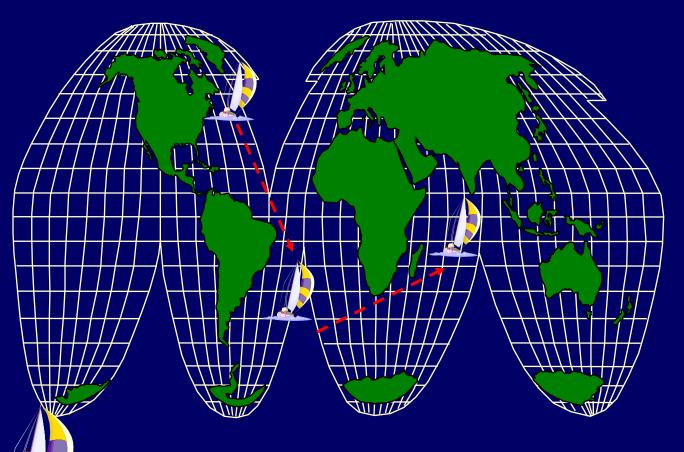
The unique HBV biology in the contest of liver physiology, allows the virus to replicate without liver damage

### **Natural history of Chronic Hepatitis B Virus Infection**



### **Management of HBV carriers**

Phases of infection and response to therapy are defined by **quantitative measurement of viral constituents** (nucleic acid, *HBV-DNA* and proteins, *HBeAg* and *HBsAg*) and **antiviral immune response** (antiviral antibody levels, IgM and Total anti-HBc)



Latitude Longitude Time Date

ALT
HBV-DNA
IgM anti-HBc
HBsAg

32°09'18"N, O 40°07'13"; 8.15 a.m., 19.01.01 29°00'S, E 24°00'; 8.15 a.m., 29.01.01 3°15'N, E 73°00' 8.15 a.m., 3.2.02

However, serum biomarkers of the virus/host interplay are influenced by many interfering factors, such as host and virus heterogeneity and have to be repeatedly measured overtime to accurately define the extent of immune control or of the antiviral treatment effectiveness



Unmet needs are biomarkers of the achievement of a sustained control of HBV infection by the host's immune system





Virions + defective particles (exceeding virions by a factor of 10<sup>2</sup>-10<sup>5</sup>)

replication cccDNA transcription/ mRNAs translation

#### **HBsAg serum levels reflect:**

- in HBeAg positive patients, the overall amount of cccDNA
- in HBeAg negative carriers, the transcriptionally active cccDNA

Brunetto et al, J Hepatol 2010

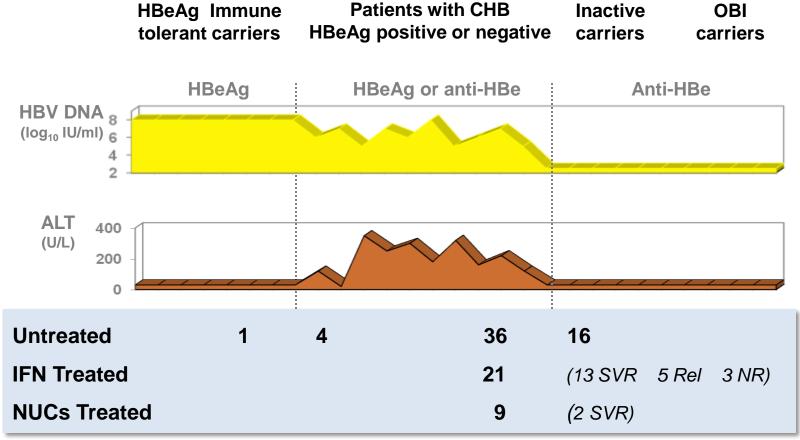
#### Circulating HBsAg particles carry human miRNAs

Liver Specific: miR-27a, miR-30b, miR-122, miR-126, miR-145

Immune regulatory: miR-106b, miR-223

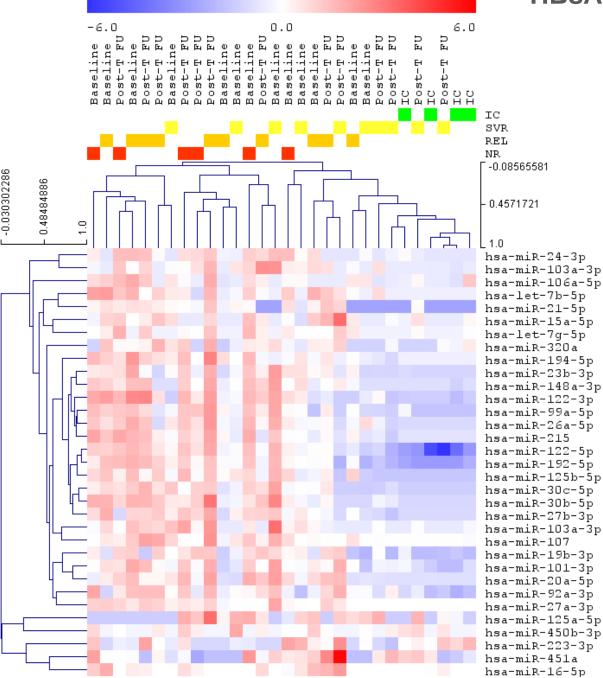
# A serum microRNA signature is associated with the immune control of chronic hepatitis B virus infection

Dynamic variation of miRNA profiling was characterized in **141** sera and immunoprecipitated HBsAg-particles of **61** HBsAg-carriers during different HBV infection phases and according to treatment response in **32** patients



(response to therapy; SVR= sustained virological responders)

#### **HBsAg particles miRNA profiling**

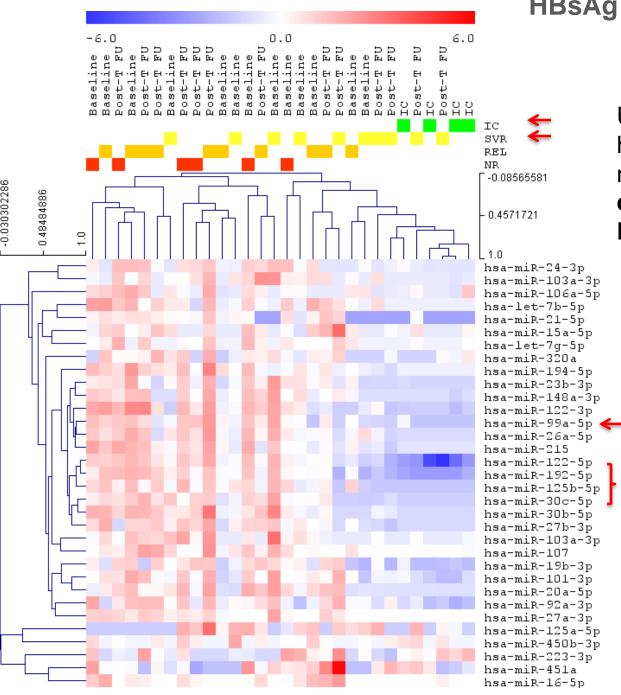


Each miRNA was assayed once by qPCR on the Ready-to-use microRNA-PCR panels I/II containing 739-miRNA-assays

Differences were observed in the mean variation of the average signal (ΔCq) between Inactive Carriers (IC) and BL- Chronic Hepatitis B and post-treatment follow-up of NR and REL

Brunetto MR, AASLD 2013, submitted

#### **HBsAg particles miRNA profiling**



Unsupervised two-way
hierarchical analysis of
miRNA showed the
clustering of post-Treatment
Follow-up of SVR with IC

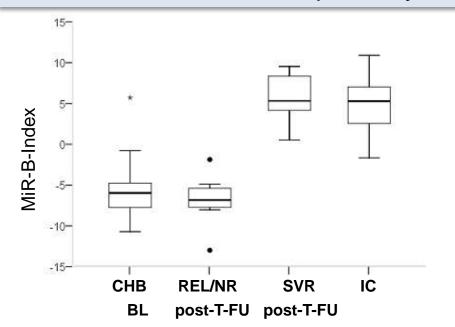
miRNA differentially expressed comparing BL-SVR-CHB to both post-Treatment-Follow-Up of SVR and IC

Brunetto MR, AASLD 2013, submitted

### Serum miRNA profiling

- ✓ 21 miRNAs were differentially expressed in Inactive Carriers (IC) and Chronic Hepatitis B (CHB) patients with the largest difference for miR-122-5p, miR-99a-5p and miR-192-5p.
- ✓ The 3 liver-miRNAs were significantly down-regulated during and after end of treatment in SVR.
- MiRNA-profiles of IC and SVR at post-TFU clustered in the heatmap.

Liver-miRNAs were combined with 3 additional miRNA as internal controls to build a MiR-B-Index which showed 100%-sensitivity and 84.4%-specificity in identifying IC.



Mir-B-Index improved during therapy and post-treatment-follow-up reaching IC-like values in IFN/NUCs-SVR (5.28,-1.65/10.91 vs 5.33, 0.54/9.53, P=.553)

- Dynamic change of a miRNA signature may identify the natural occurring and therapy induced immune control of HBV infection.
- The same signature qualifies as new diagnostic biomarker to satisfy the unmet need of the early identification of the sustained switch from chronic active hepatitis to the inactive HBV infection in patients treated with antivirals.
- The in vivo study of circulating HBsAg-associated hepatocellular miRNAs provide a unique model to study the epigenetics of liver- physiopathology.





The implementation of new diagnostic tools warrant a more accurate management of chronic HBV carriers.

However, because of the highly dynamic course of both infection and disease, the appropriate therapeutic intervention requires an integration of all biological and medical data to accurately simulate the interplay between HBV and host's immune response during antiviral

therapy, as we already done for chronic hepatitis C.

(Colombatto et al. Clin Phar & Ther 2008)

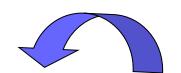




## Hepatology Unit & Laboratory of Molecular Genetics and Pathology of Hepatitis Viruses

University Hospital of Pisa, Italy

Antonella Cristofani Simonetta Ferretti (part-time) Simona Giannetti Teresa Crisponi Isabella Rossi (Chief)



Pietro Ciccorossi
Barbara Coco
Piero Colombatto
Filippo Oliveri
Veronica Romagnoli
Carlotta Rastelli
Beatrice Cherubini

#### **Nurses**





#### **Medical Doctors**

#### **Administrative Personel**

at the moment missing...

#### **Biologists**





Ranieri Bizzarri



Ferruccio Bonino
Gatroenterology Chair
Pisa University