



# New endpoints and biomarkers for treatment of CHB

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# Guidance for design and endpoints of clinical trials in chronic hepatitis B

## Report from the 2019 EASL-AASLD HBV Treatment Endpoints Conference

|                               | Sterilizing 'cure'         | Idealistic functional 'cure' | Realistic functional 'cure'          | Attainable Partial functional 'cure' |
|-------------------------------|----------------------------|------------------------------|--------------------------------------|--------------------------------------|
| Clinical scenario             | Never infected             | Recovery after acute HBV     | Chronic HBV with HBsAg loss          | Inactive carrier off treatment       |
| HBsAg                         | Negative                   | Negative                     | Negative                             | Positive                             |
| Anti-HBs                      | Negative/Positive          | Positive                     | Positive/negative                    | Negative                             |
| HBeAg                         | Negative                   | Negative                     | Negative                             | Negative                             |
| Serum HBV DNA                 | Not detected               | Not detected                 | Not detected                         | Low level or not detected            |
| Hepatic cccDNA, transcription | Not detected<br>Not active | Detected<br>Not active       | Detected<br>Not active               | Detected<br>Low level                |
| Integrated HBV DNA            | Not detected               | Detected?                    | Detected                             | Detected                             |
| Liver disease                 | None                       | None                         | Inactive, fibrosis regress over time | Inactive                             |
| Risk of HCC                   | Not increased              | Not increased                | Declines with time                   | Risk lower vs. active hepatitis      |

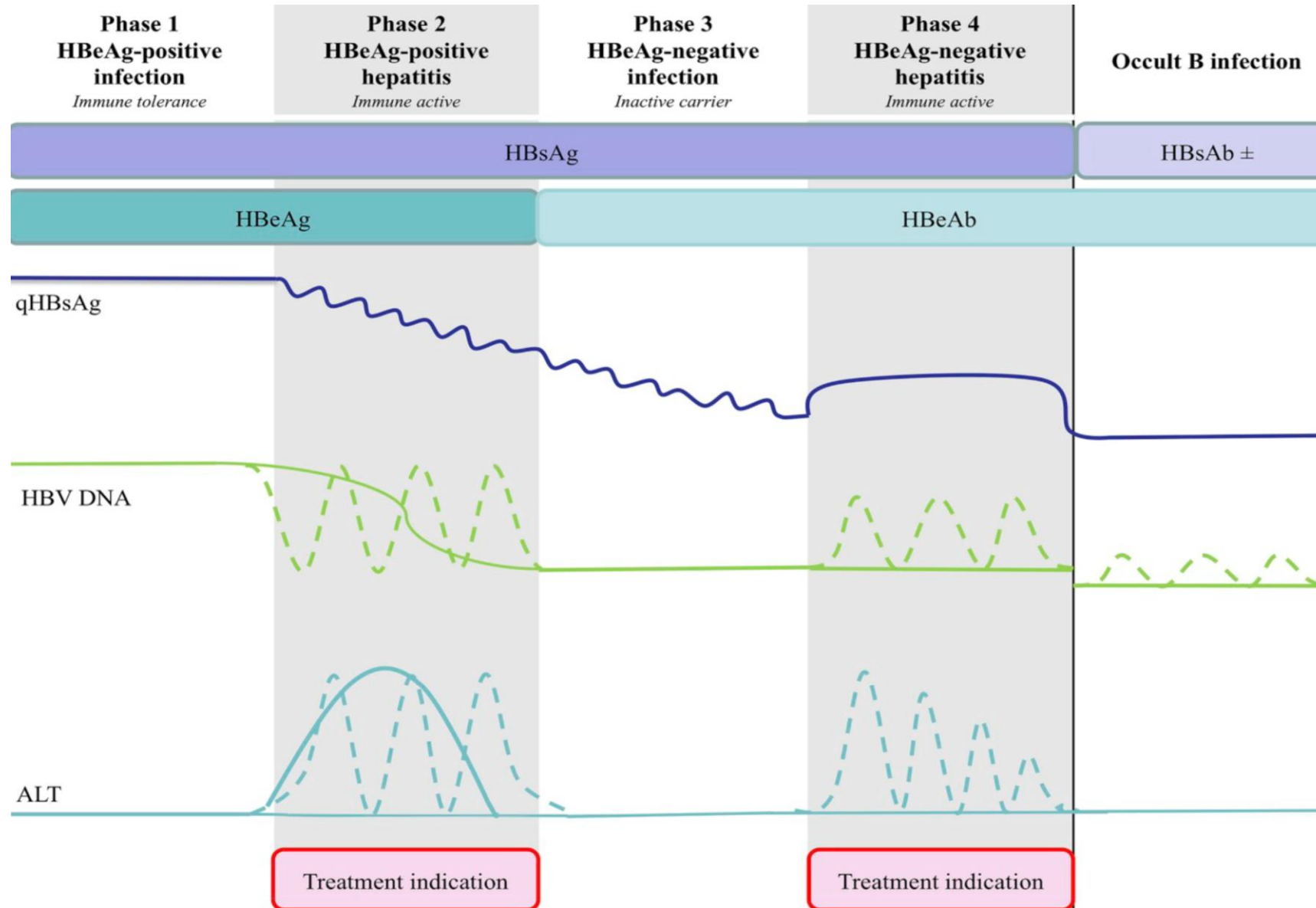
# EASL 2017 Clinical Practice Guidelines on the management of hepatitis B virus infection<sup>☆</sup>

European Association for the Study of the Liver<sup>\*</sup>

**HBsAg loss, with or without anti-HBs seroconversion, is an optimal endpoint, as it indicates profound suppression of HBV replication and viral protein expression**

**(Evidence level II-1, grade of recommendation 1)**

# Treatment indication in chronic HBV infection – The Phase matters





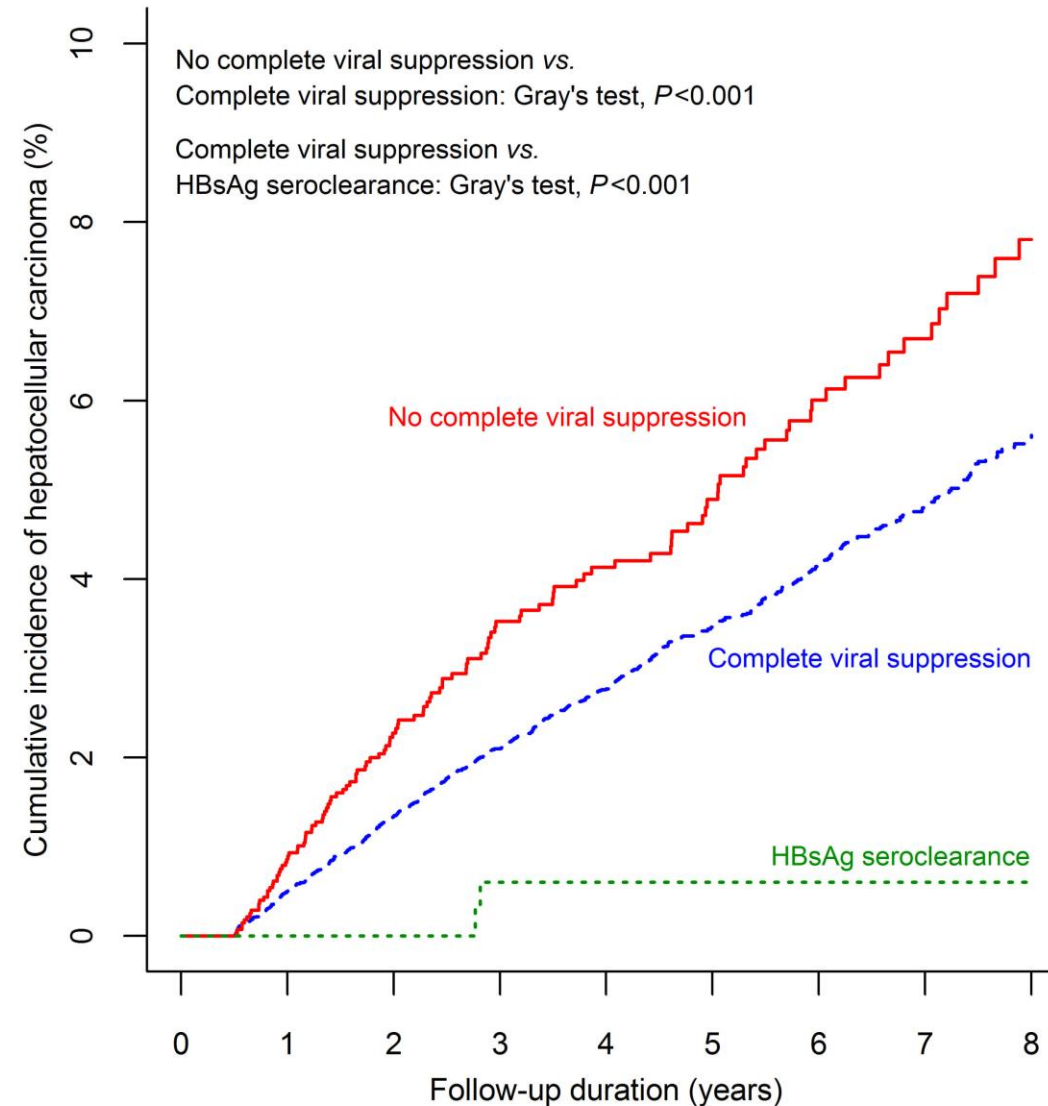
**Why we are aiming for HBsAg loss if HBsAg per se is not an indicator to start antiviral treatment?**

# **EASL CPG - Why aiming for HBsAg loss?**

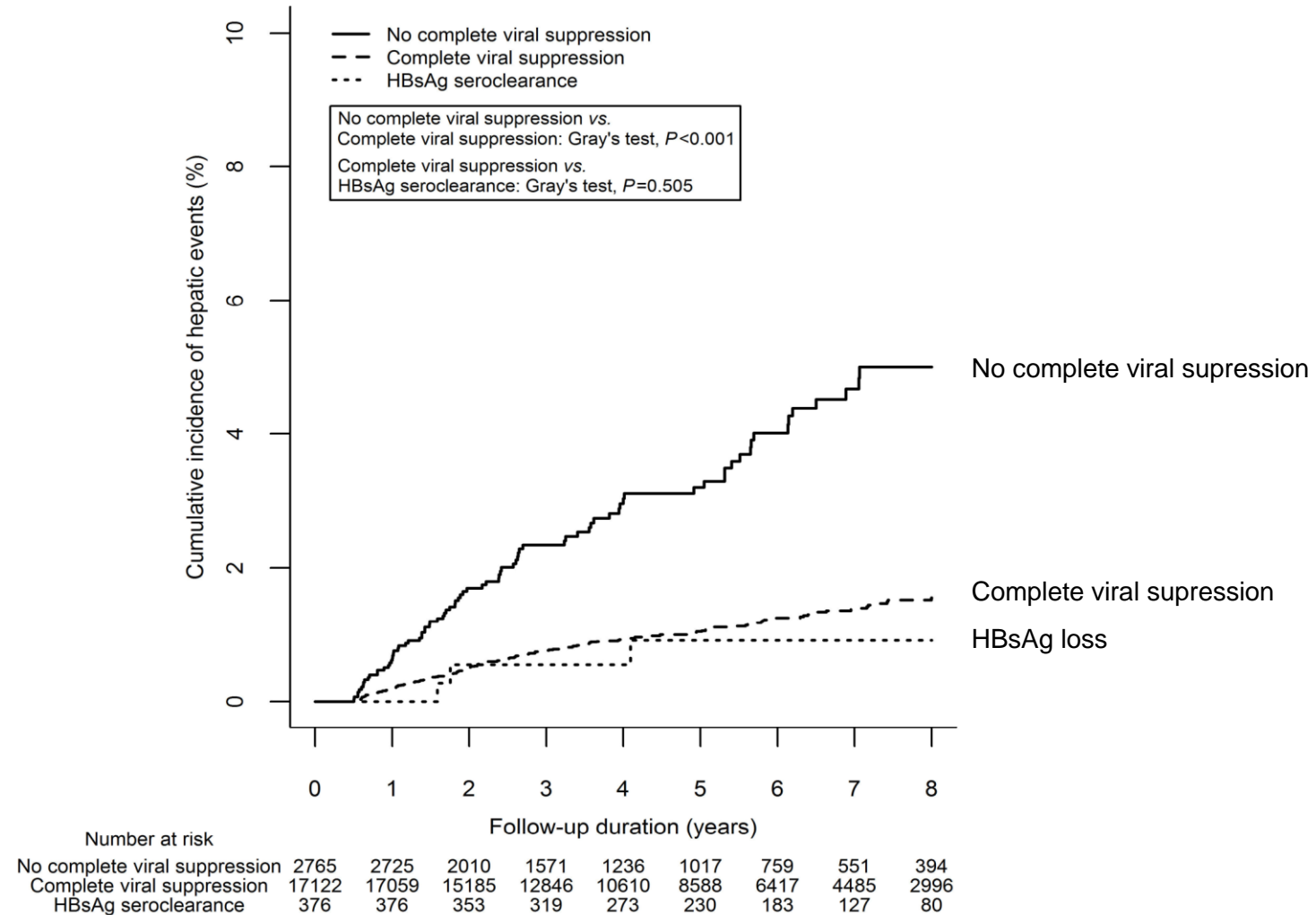
- As chronic HBV infection cannot be completely eradicated due to the persistence of cccDNA and integrated HBV DNA, it remains unclear whether HBsAg loss adds to the prevention of the long-term complications of chronic HBV infection beyond what can be achieved by the suppression of HBV DNA replication alone**
- HCC may still develop even after spontaneous HBsAg loss (annual rate approximately 0.55%)**
- The risk, however, is lower if HBsAg loss is achieved at a younger age and/or in the absence of significant fibrosis**

**Do we need to achieve HBsAg loss fully prevent  
any adverse outcome?**

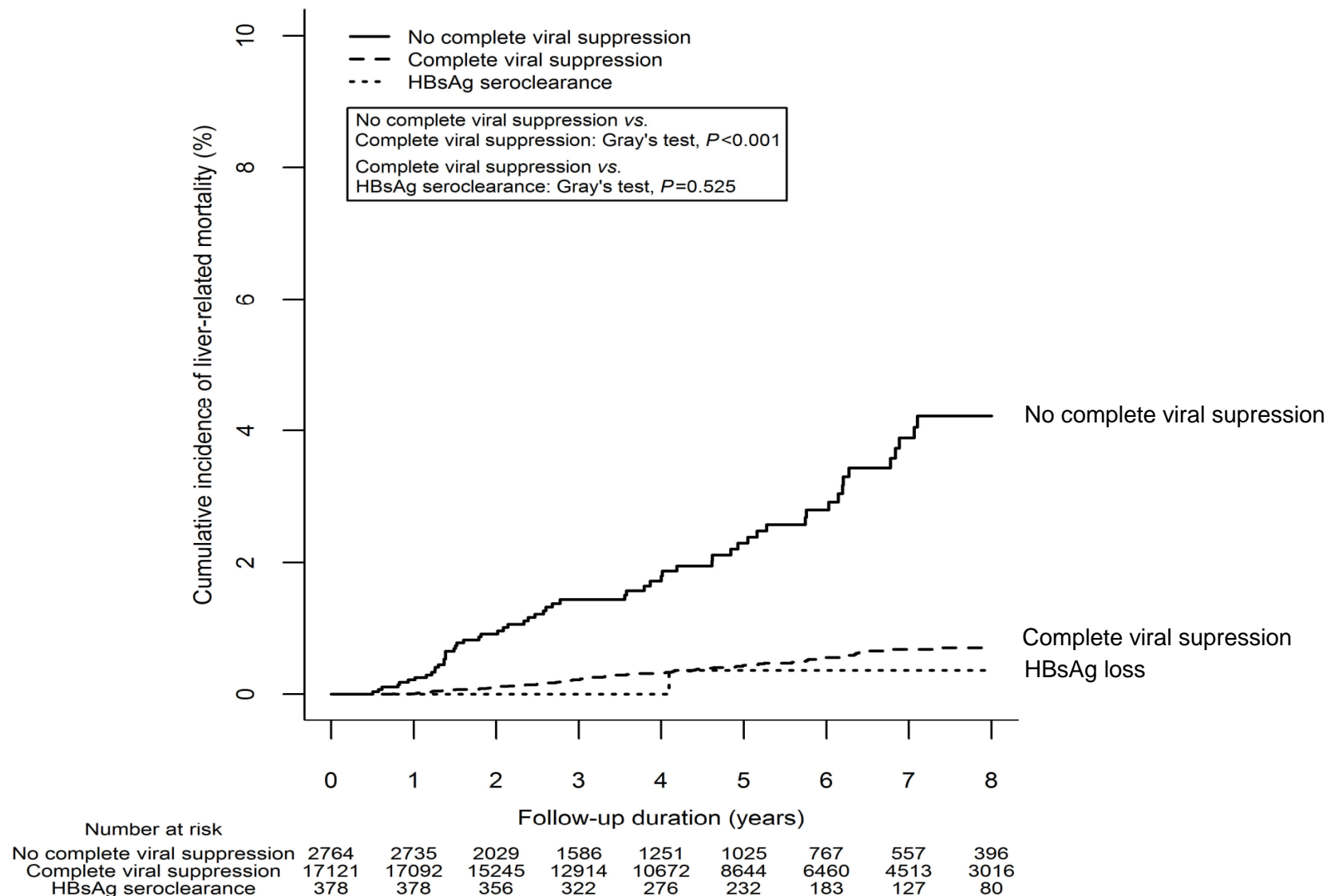
# HBsAg loss further reduces HCC risk after complete viral suppression



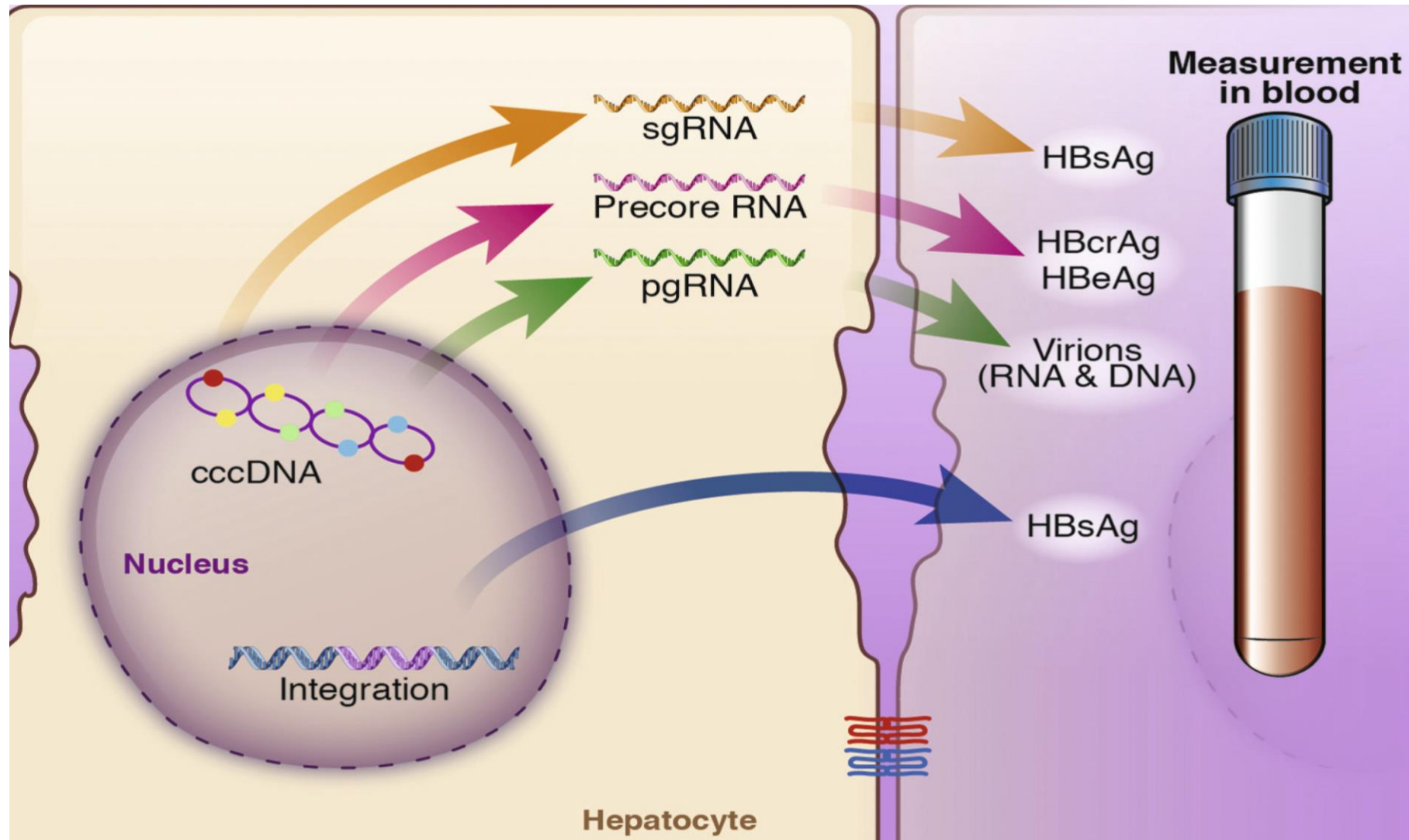
# Incidence of **hepatic events** in NA-treated patients according to level of viral suppression and HBsAg loss



# Incidence of **liver-related mortality** in NA-treated patients according to level of viral suppression and HBsAg loss



# HBsAg a valid endpoint for new antiviral compounds?





# Guidance for design and endpoints of clinical trials in chronic hepatitis B

## Aiming for attainable functional “cure”?

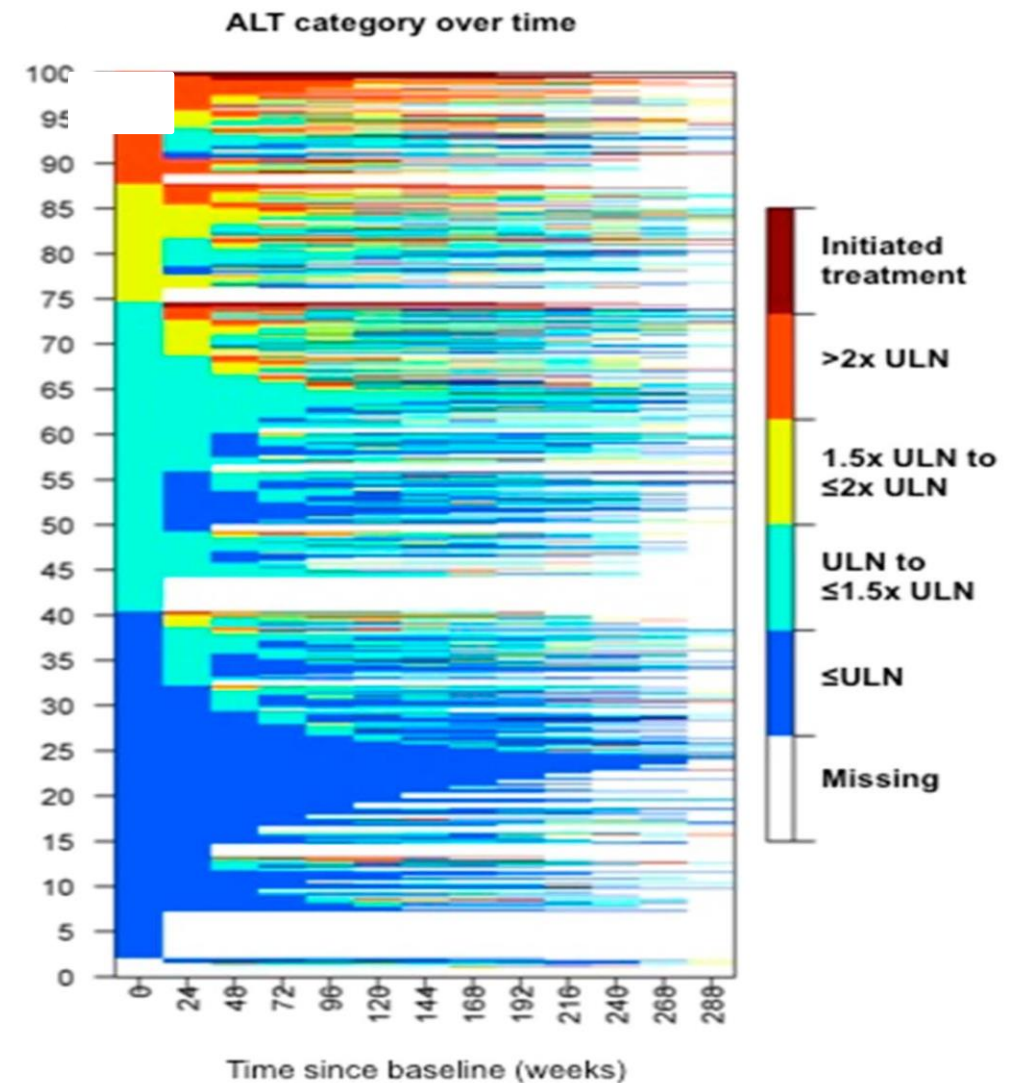
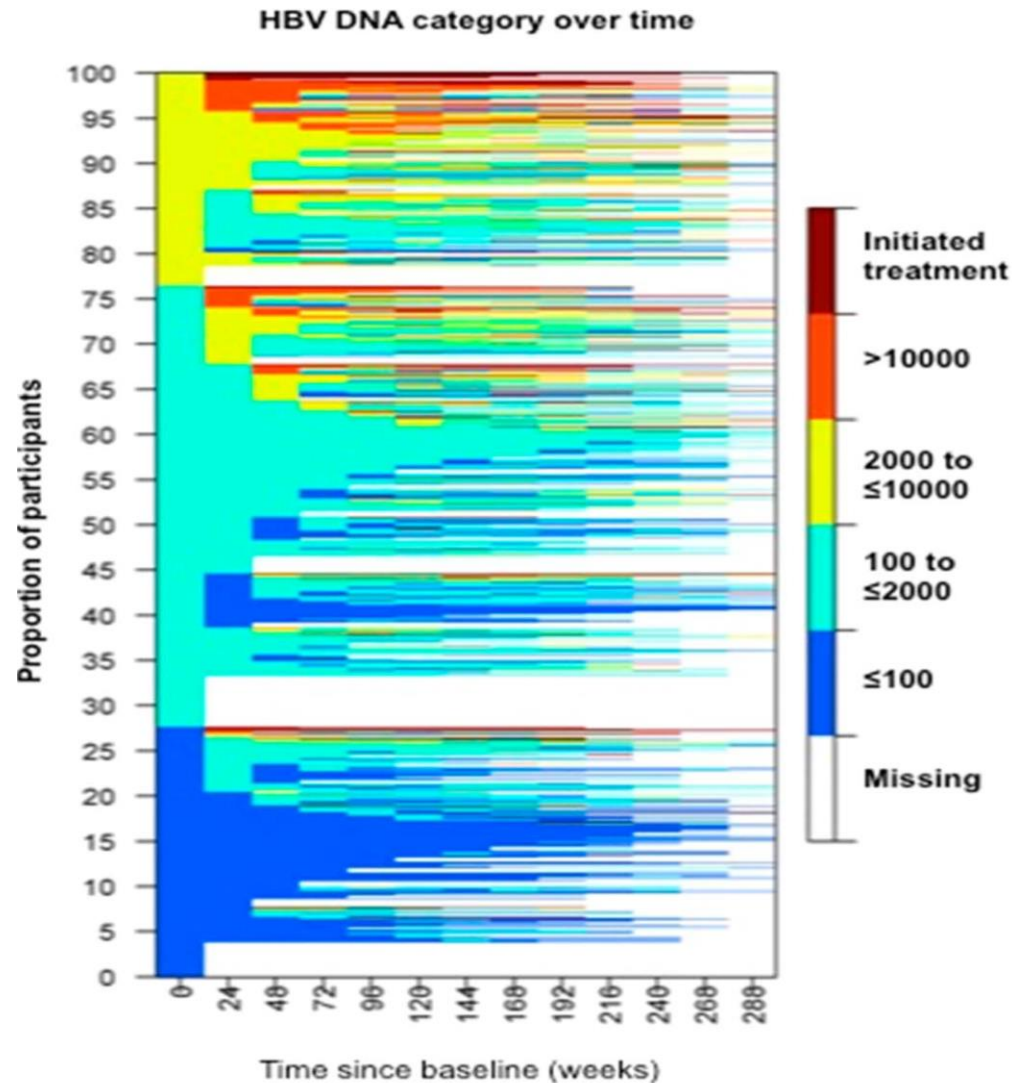
|                               | Sterilizing ‘cure’ | Idealistic functional ‘cure’ | Realistic functional ‘cure’ | Attainable Partial functional ‘cure’ |
|-------------------------------|--------------------|------------------------------|-----------------------------|--------------------------------------|
| Clinical scenario             |                    |                              |                             | Inactive carrier off treatment       |
| HBsAg                         |                    |                              |                             | Positive                             |
| Anti-HBs                      |                    |                              |                             | Negative                             |
| HBeAg                         |                    |                              |                             | Negative                             |
| Serum HBV DNA                 |                    |                              |                             | Low level or not detected            |
| Hepatic cccDNA, transcription |                    |                              |                             | Detected<br>Low level                |
| Integrated HBV DNA            |                    |                              |                             | Detected                             |
| Liver disease                 |                    |                              |                             | Inactive                             |
| Risk of HCC                   |                    |                              |                             | Risk lower vs. active hepatitis      |

# Guidance for design and endpoints of clinical trials in chronic hepatitis B

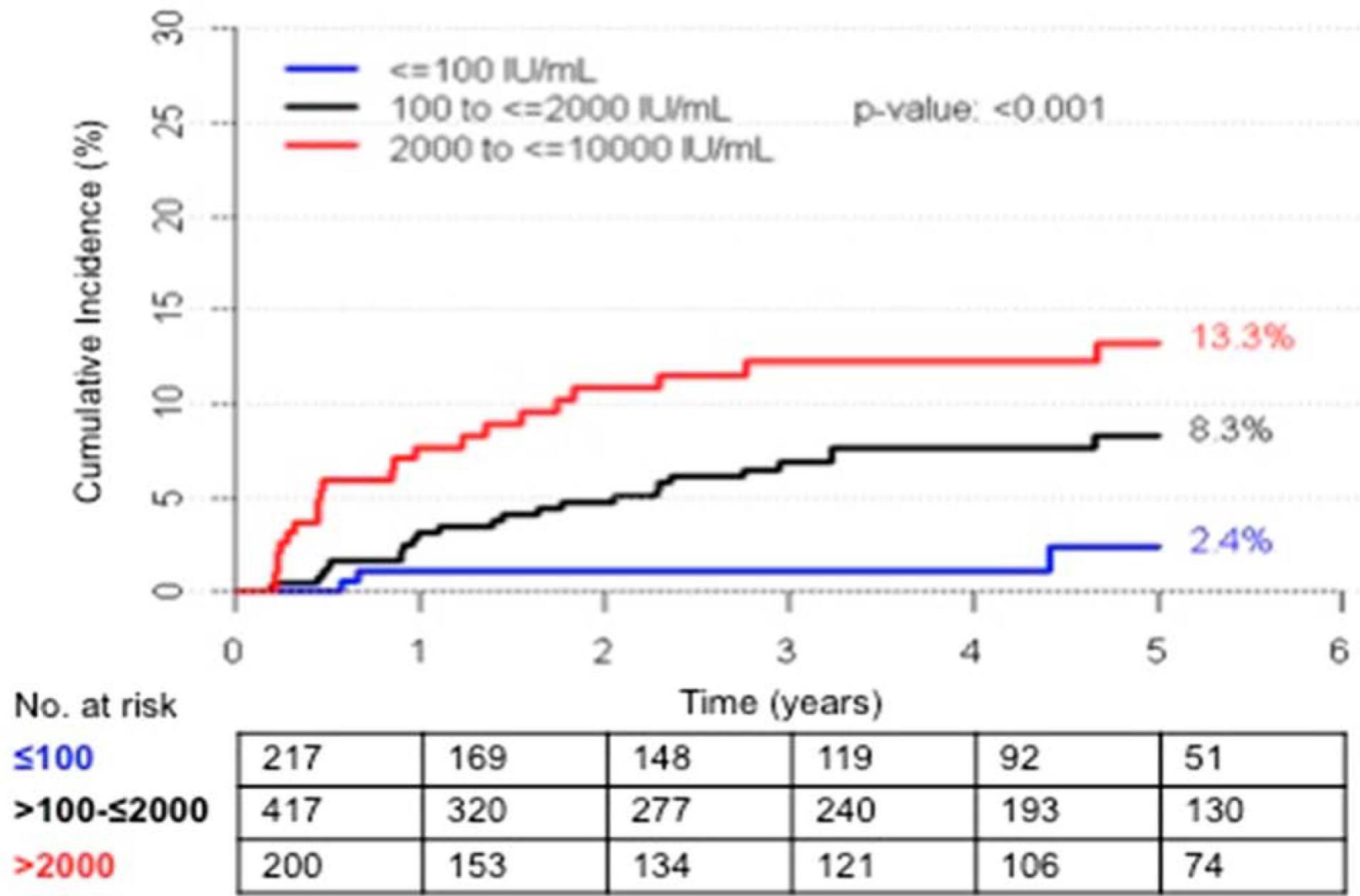
## Aiming for attainable functional “cure”?

|                               | Sterilizing ‘cure’  | Idealistic functional ‘cure’ | Realistic functional ‘cure’ | Attainable Partial functional ‘cure’ |
|-------------------------------|---|------------------------------|-----------------------------|--------------------------------------|
| Clinical scenario             | <div>Stability of the response?<br/>Disease reactivation risk?<br/>Long-term HCC risk?<br/>Extrahepatic manifestations?<br/>Long-term monitoring?</div> |                              |                             | Inactive carrier off treatment       |
| HBsAg                         |   |                              |                             | Positive                             |
| Anti-HBs                      |   |                              |                             | Negative                             |
| HBeAg                         |   |                              |                             | Negative                             |
| Serum HBV DNA                 |   |                              |                             | Low level or not detected            |
| Hepatic cccDNA, transcription |   |                              |                             | Detected<br>Low level                |
| Integrated HBV DNA            |   |                              |                             | Detected                             |
| Liver disease                 |   |                              |                             | Inactive                             |
| Risk of HCC                   |   |                              |                             | Risk lower vs. active hepatitis      |

# Phase transition in HBeAg negative infection (carriers) according to baseline HBV DNA and ALT levels



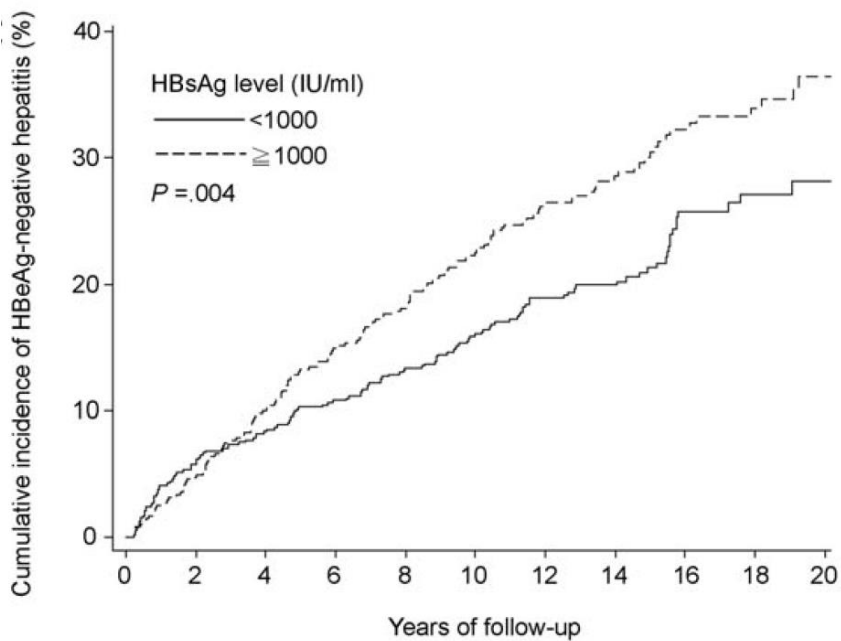
# Phase transition in HBeAg negative infection (carriers) according to baseline HBV DNA levels



**HBsAg – a disease progression marker?**  
**Does the level matters?**

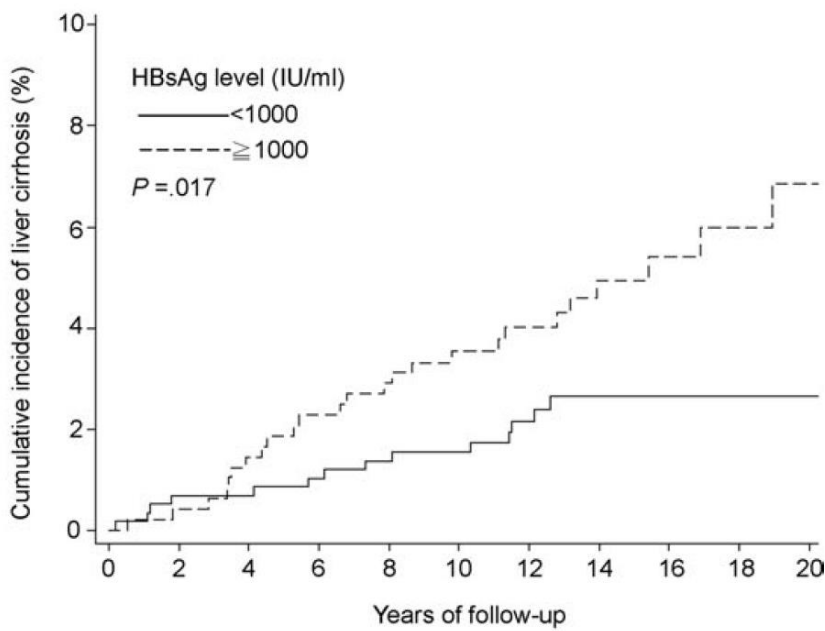
# HBsAg quantification – a biomarker to predict disease progression in patients with low baseline viral load (< 2000 IU/mL)

Reactivation of HBeAg negative hepatitis



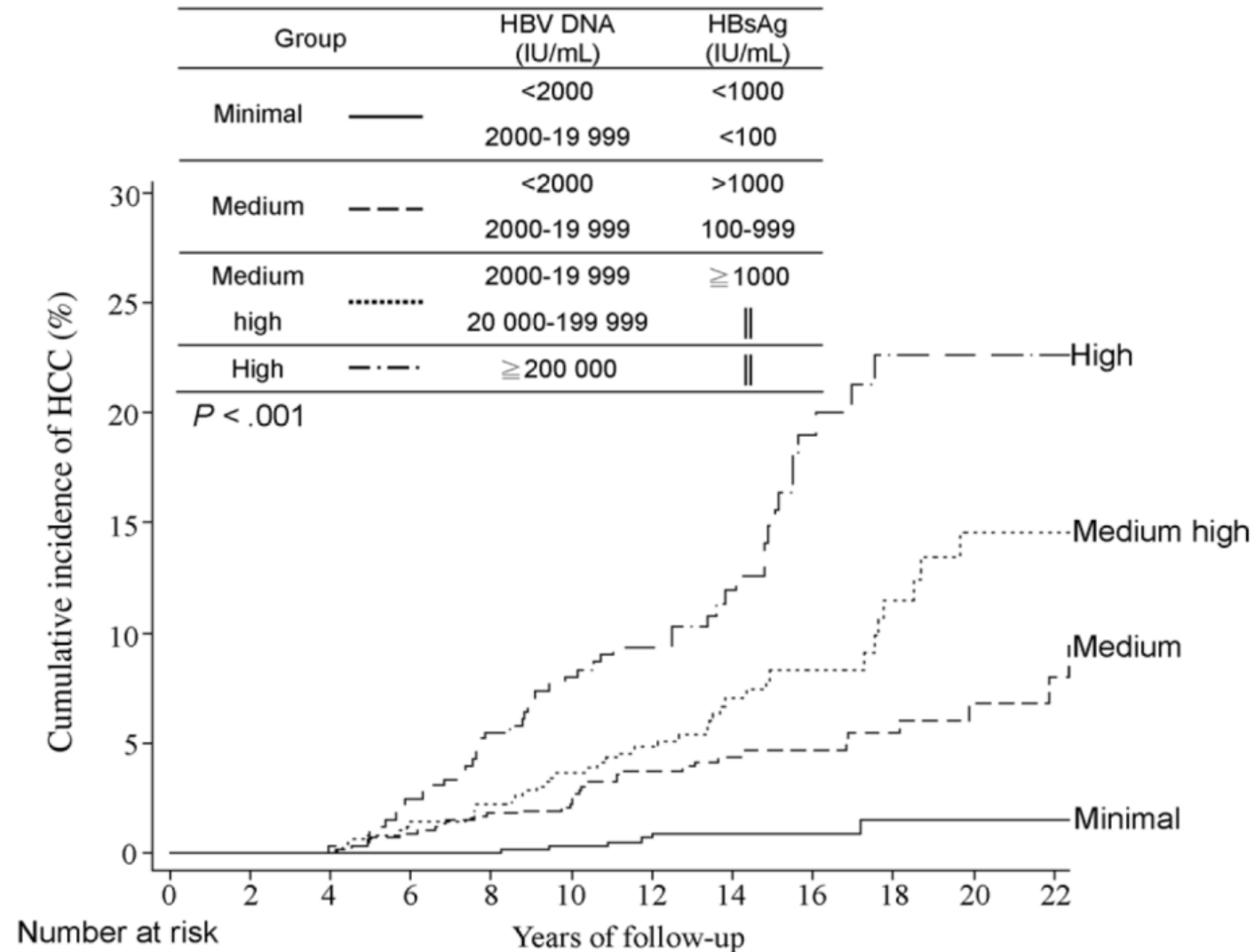
| Number at risk                         |     |     |     |     |     |     |     |     |     |     |    |
|--|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|
| Serum HBsAg levels at baseline (IU/ml) |     |     |     |     |     |     |     |     |     |     |    |
| <1000                                  | 585 | 551 | 536 | 521 | 503 | 454 | 357 | 265 | 146 | 90  | 58 |
| ≥ 1000                                 | 483 | 460 | 435 | 410 | 394 | 350 | 288 | 210 | 132 | 100 | 57 |

Cirrhosis development



| Number at risk                         |     |     |     |     |     |     |     |     |     |     |    |
|--|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|
| Serum HBsAg levels at baseline (IU/ml) |     |     |     |     |     |     |     |     |     |     |    |
| <1000                                  | 585 | 581 | 581 | 578 | 573 | 532 | 432 | 315 | 182 | 111 | 71 |
| ≥ 1000                                 | 483 | 481 | 476 | 471 | 466 | 436 | 373 | 276 | 182 | 131 | 81 |

# HCC risk stratification in HBeAg-negative HBV infection by HBV DNA and HBsAg levels



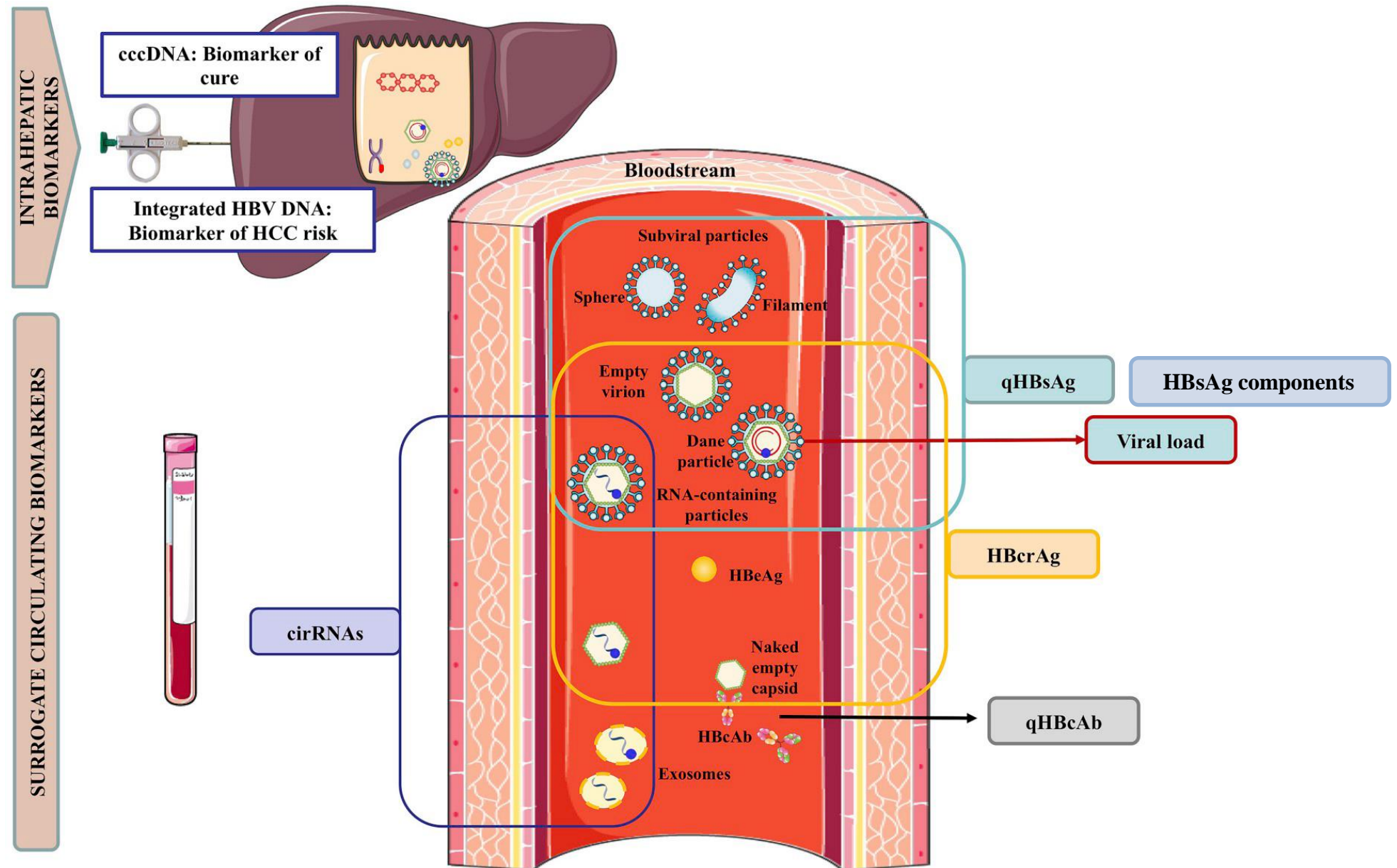


# Guidance for design and endpoints of clinical trials in chronic hepatitis B

## When aiming for attainable functional “cure” – do we need new biomarkers?

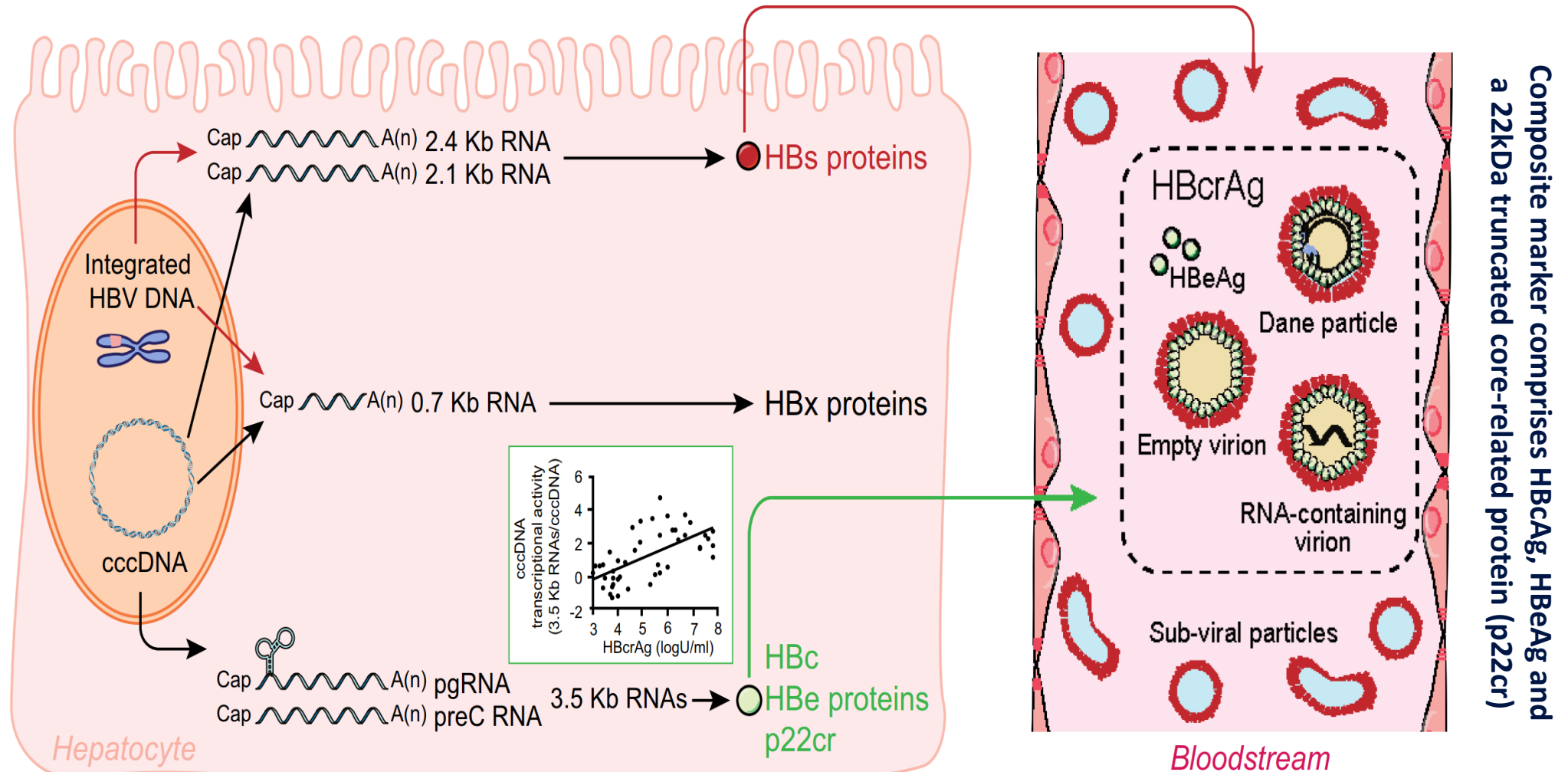
|                               | Sterilizing ‘cure’  | Idealistic functional ‘cure’ | Realistic functional ‘cure’ | Attainable Partial functional ‘cure’ |
|-------------------------------|---|------------------------------|-----------------------------|--------------------------------------|
| Clinical scenario             | <div>New HBV biomarkers</div> <div>Are they helpful for a more refined characterisation of the activity of the infection and predicting the risk of adverse outcomes?</div> |                              |                             | Inactive carrier off treatment       |
| HBsAg                         |   |                              |                             | Positive                             |
| Anti-HBs                      |   |                              |                             | Negative                             |
| HBeAg                         |   |                              |                             | Negative                             |
| Serum HBV DNA                 |   |                              |                             | Low level or not detected            |
| Hepatic cccDNA, transcription |   |                              |                             | Detected<br>Low level                |
| Integrated HBV DNA            |   |                              |                             | Detected                             |
| Liver disease                 |   |                              |                             | Inactive                             |
| Risk of HCC                   |   |                              |                             | Risk lower vs. active hepatitis      |

# Novel biomarkers for intrahepatic hepatitis B activity



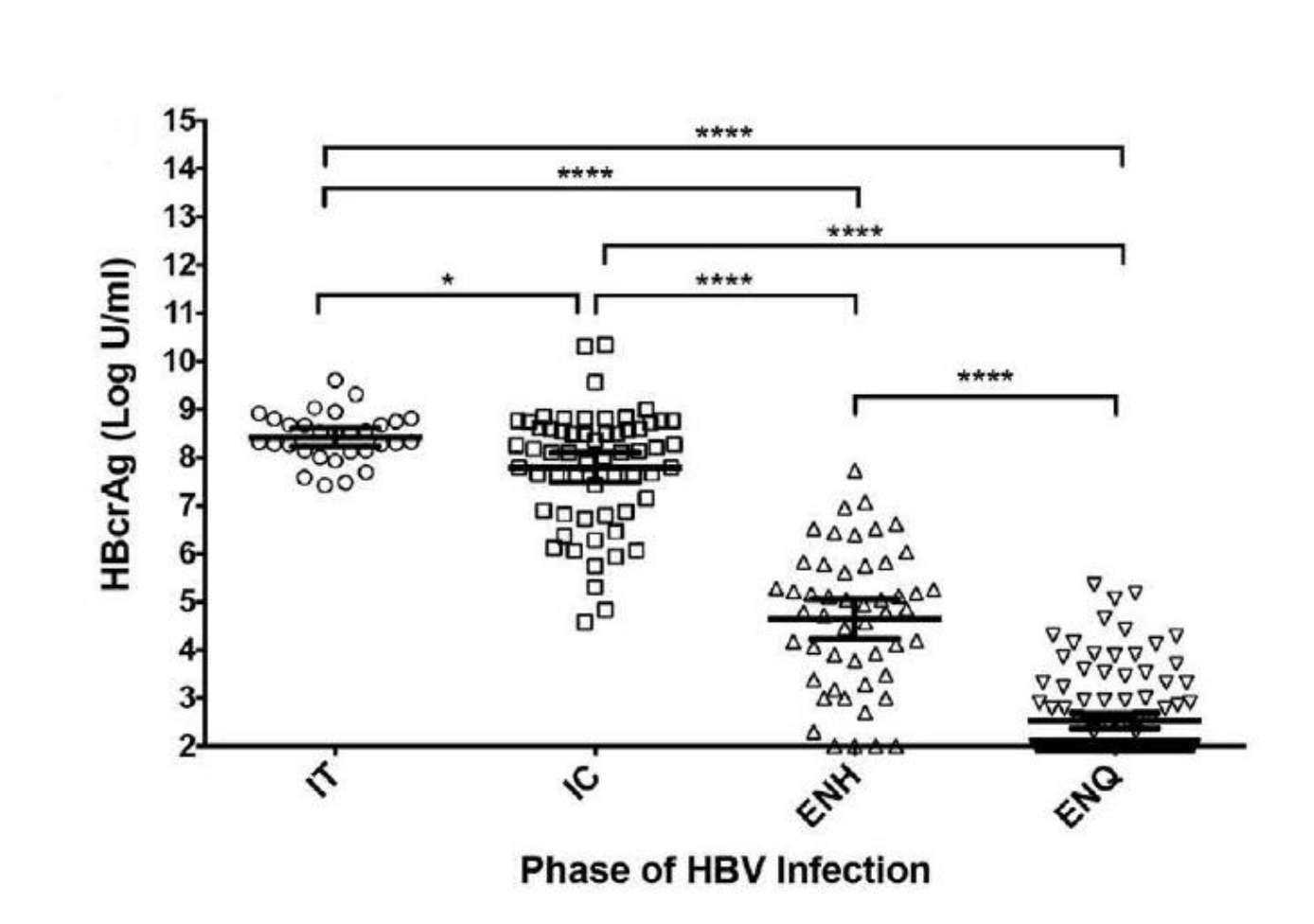
**HBcrAg**

# Hepatitis B core-related antigen (HBcrAg) a biomarker to predict transcriptional activity



**Not influenced by translation of integrated sequences**

# HBcrAg levels across phases of chronic hepatitis B



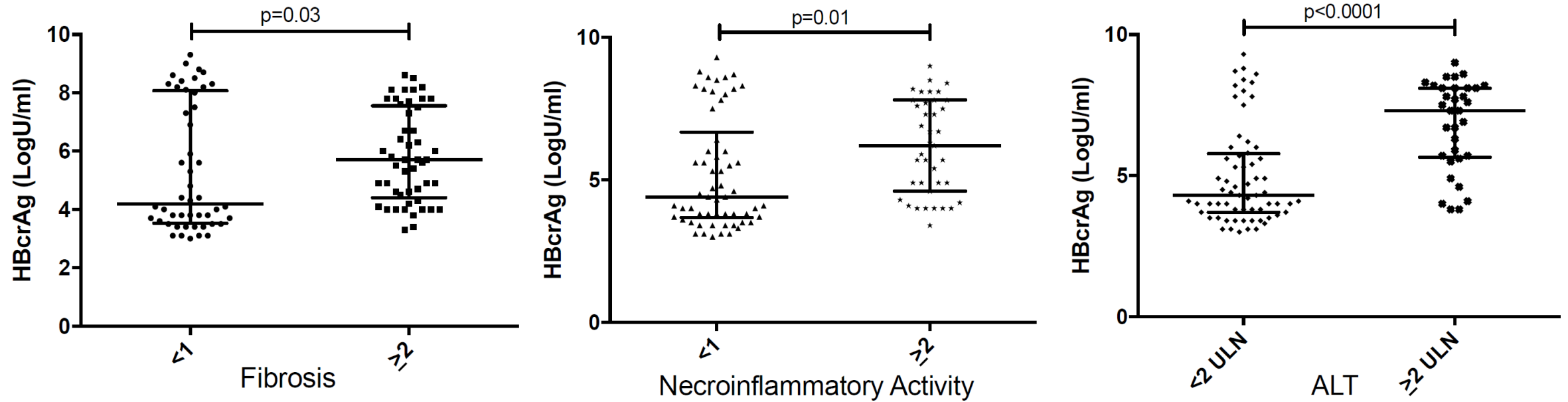
HBeAg-positive immune tolerance (IT)

HBeAg-positive immune clearance (IC)

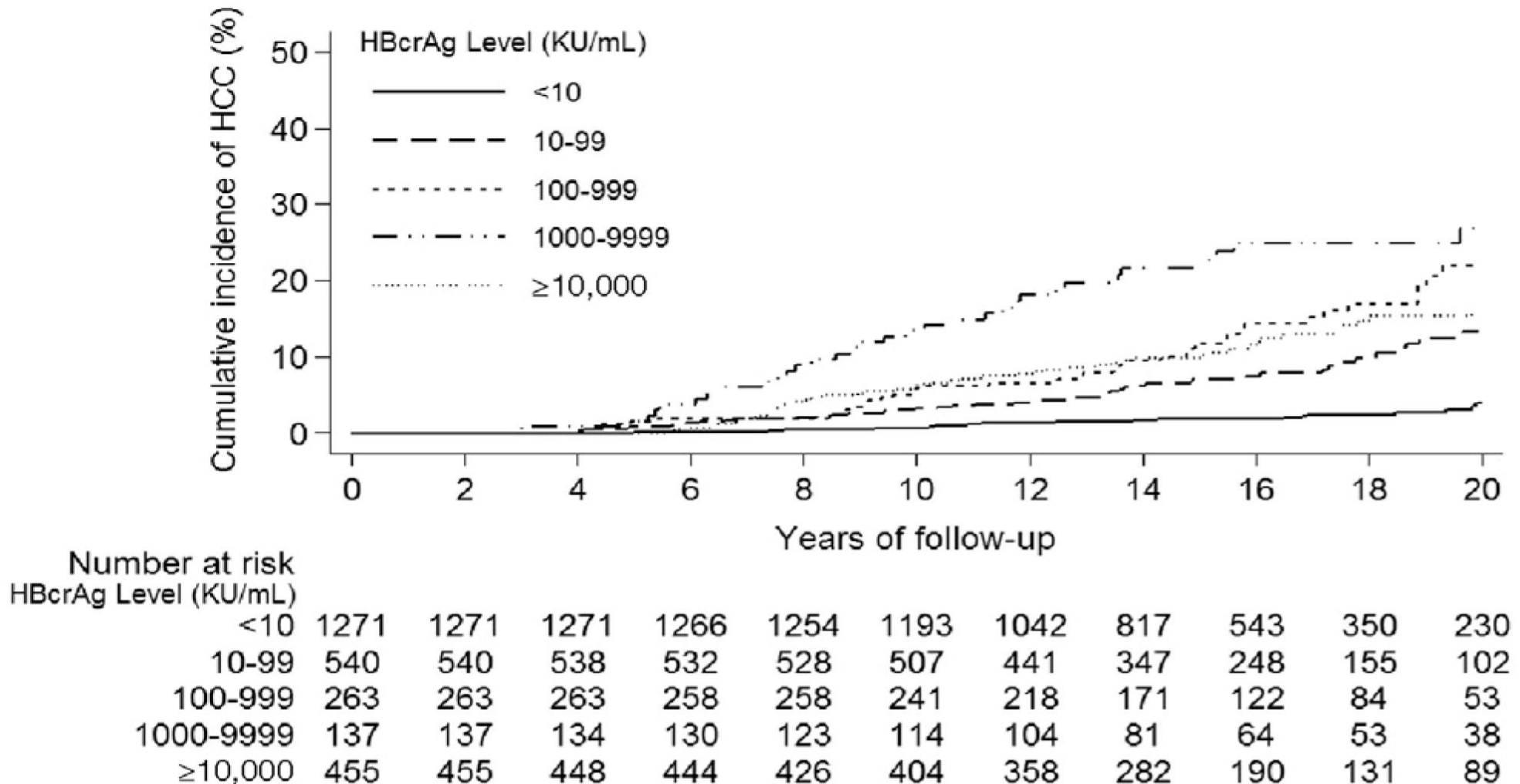
HBeAg-negative hepatitis (ENH)

inactive carriers (ENQ)

# HBcrAg levels according to liver injury markers

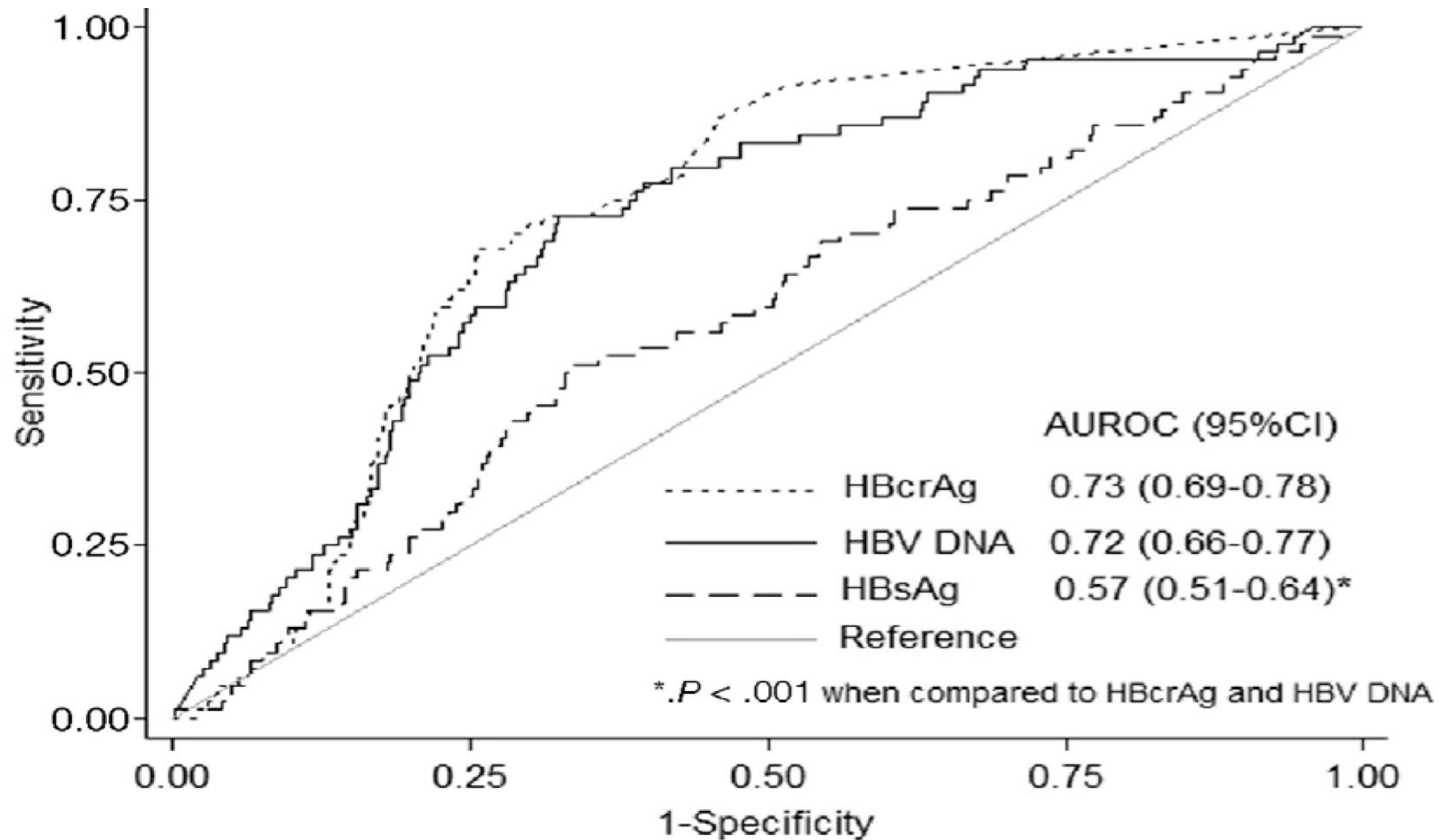


# Correlation between serum HBcrAg levels and HCC incidence

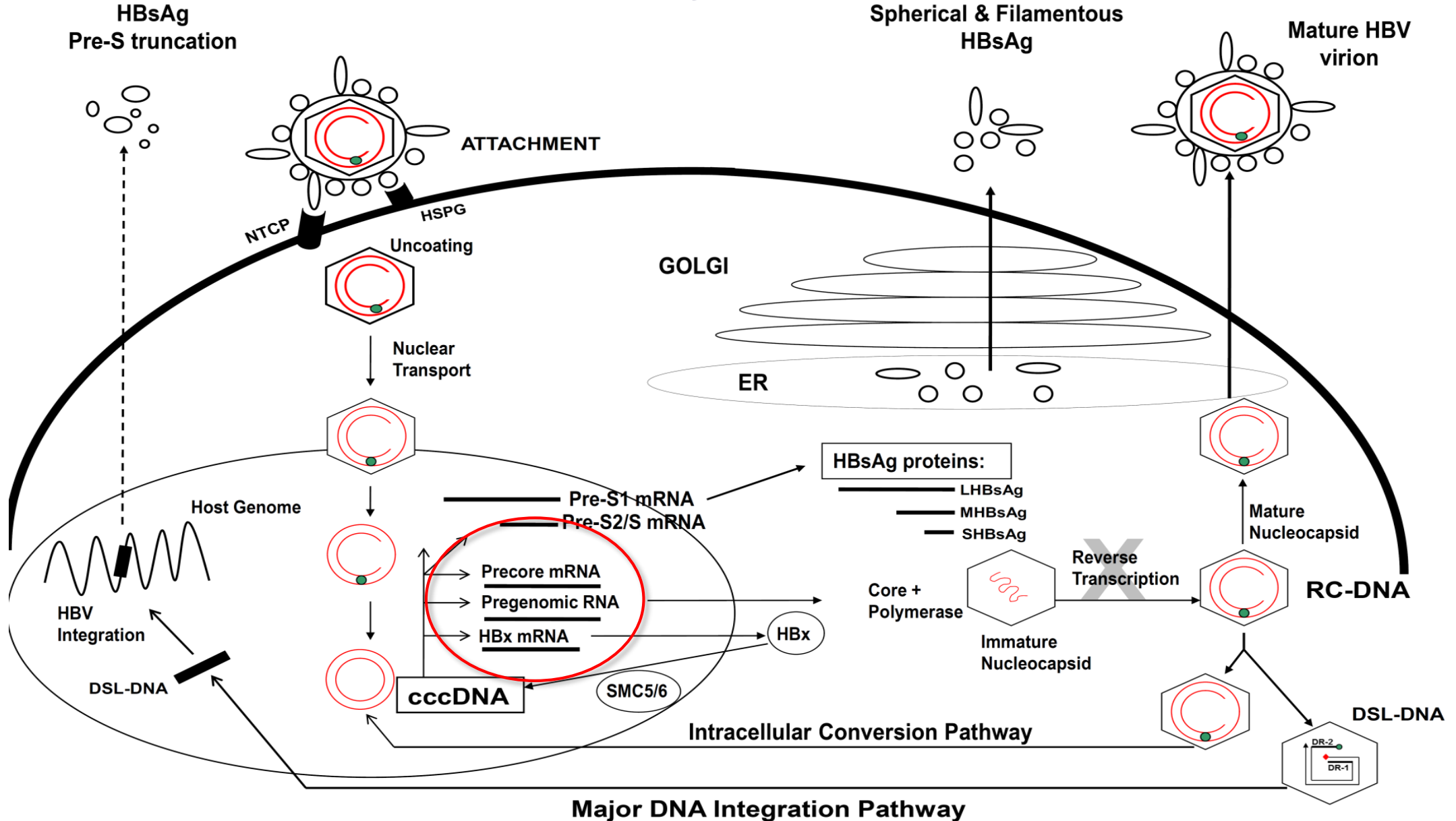




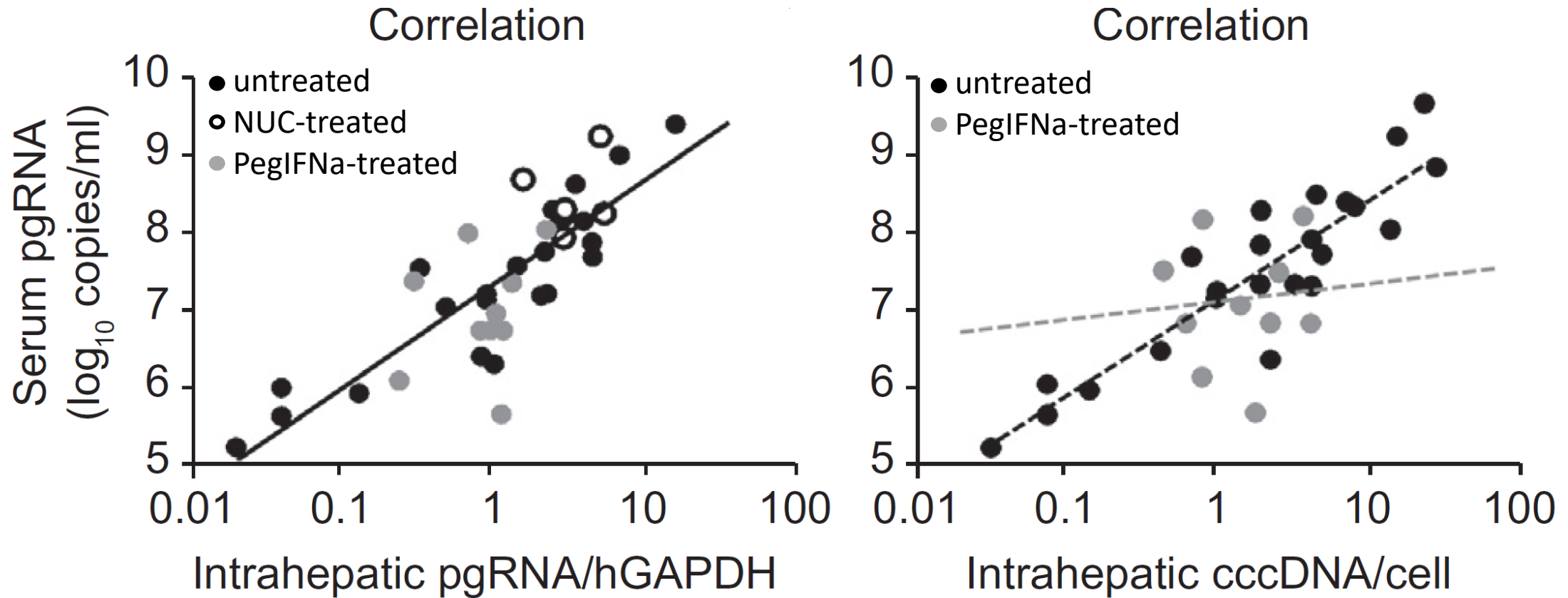
# Correlation between serum HBcrAg levels and HCC incidence



# HBV life cycle – HBV RNA



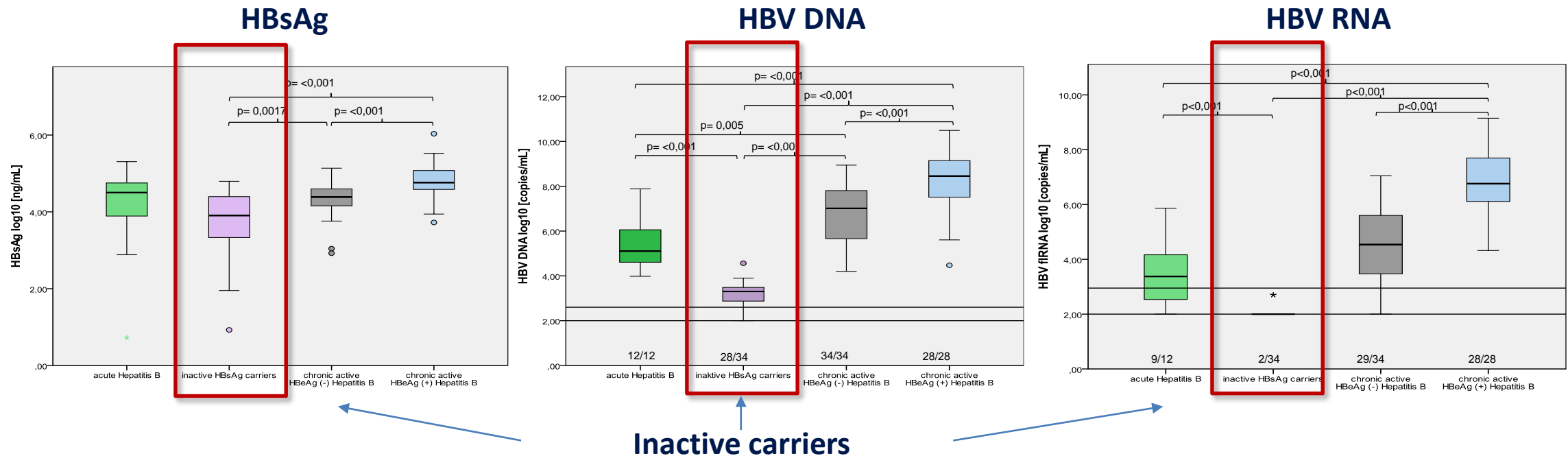
# Serum HBV pgRNA as a clinical marker for cccDNA activity



Humanized mouse model infected with HBeAg-positive wild-type HBV

**Serum HBV RNA - a disease activity marker?**

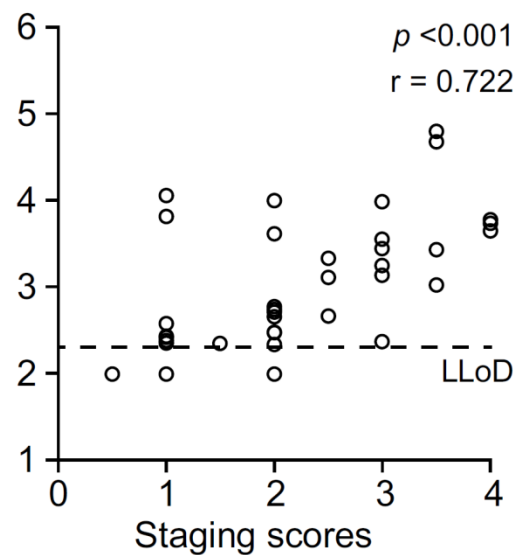
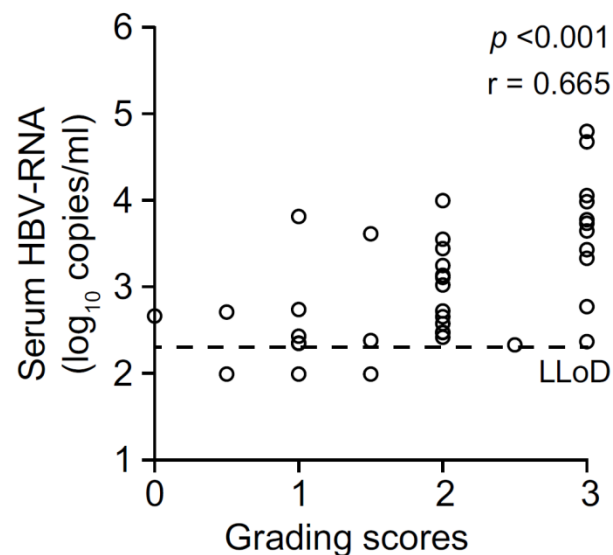
# Quantitative serum HBV RNA levels at different phases of the chronic HBV infection



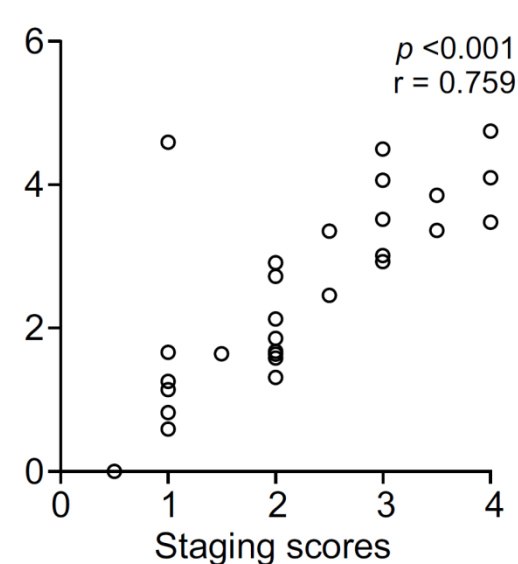
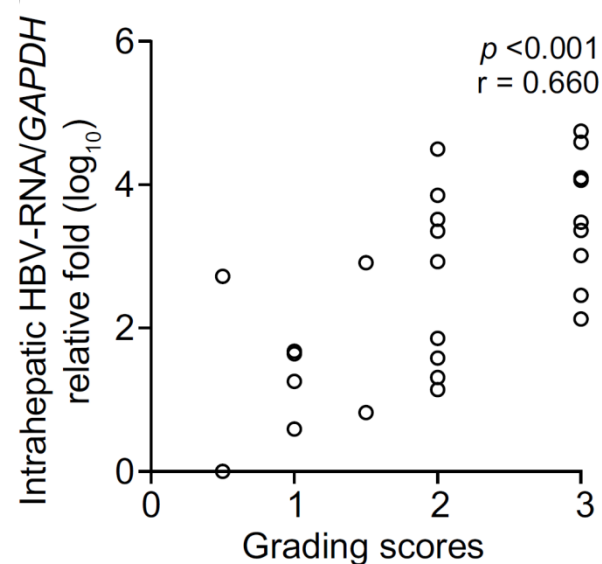
The upper and lower end of the bar features the 75- and 25-percentile. The mark inside the bar indicates the median. Significant results are given with significance level in the figure. The proportion of positive samples are indicated among the bars.

# Association between serum and intrahepatic HBV RNA levels and liver histopathology in patients under entecavir treatment

Serum HBV RNA



Intrahepatic HBV RNA

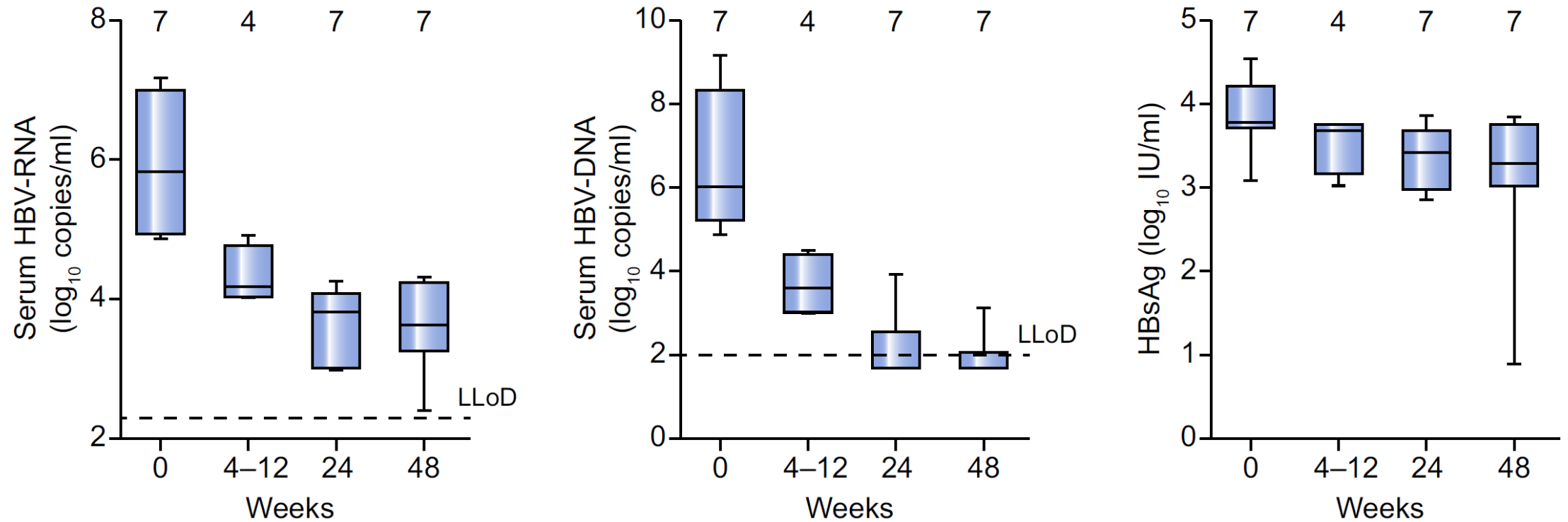


**A serum HBV-RNA level cut-off of 2.45 log<sub>10</sub> copies/mL had the highest accuracy for distinguishing mild (score <2) from severe liver histopathology**

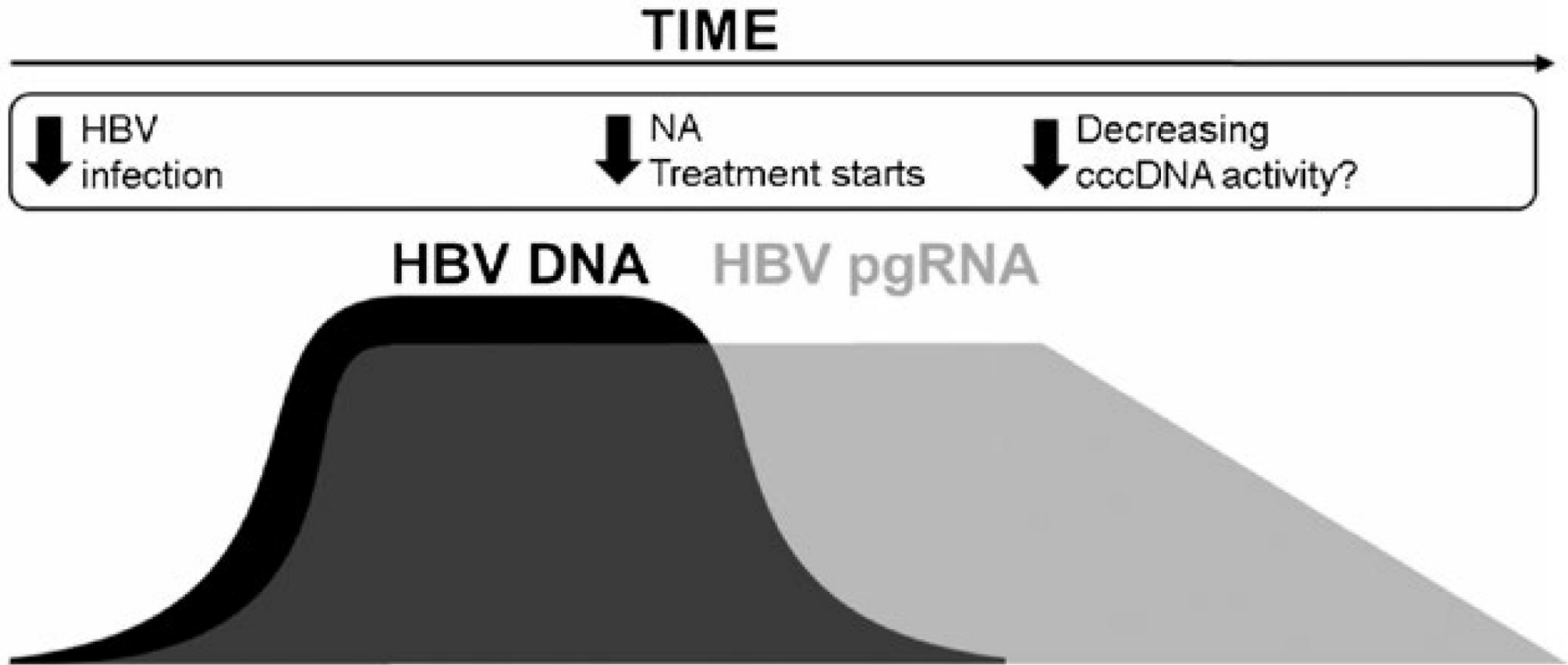
**Serum HBV RNA for predicting treatment response?**



# Kinetics of HBV serum markers during 48 weeks of entecavir therapy

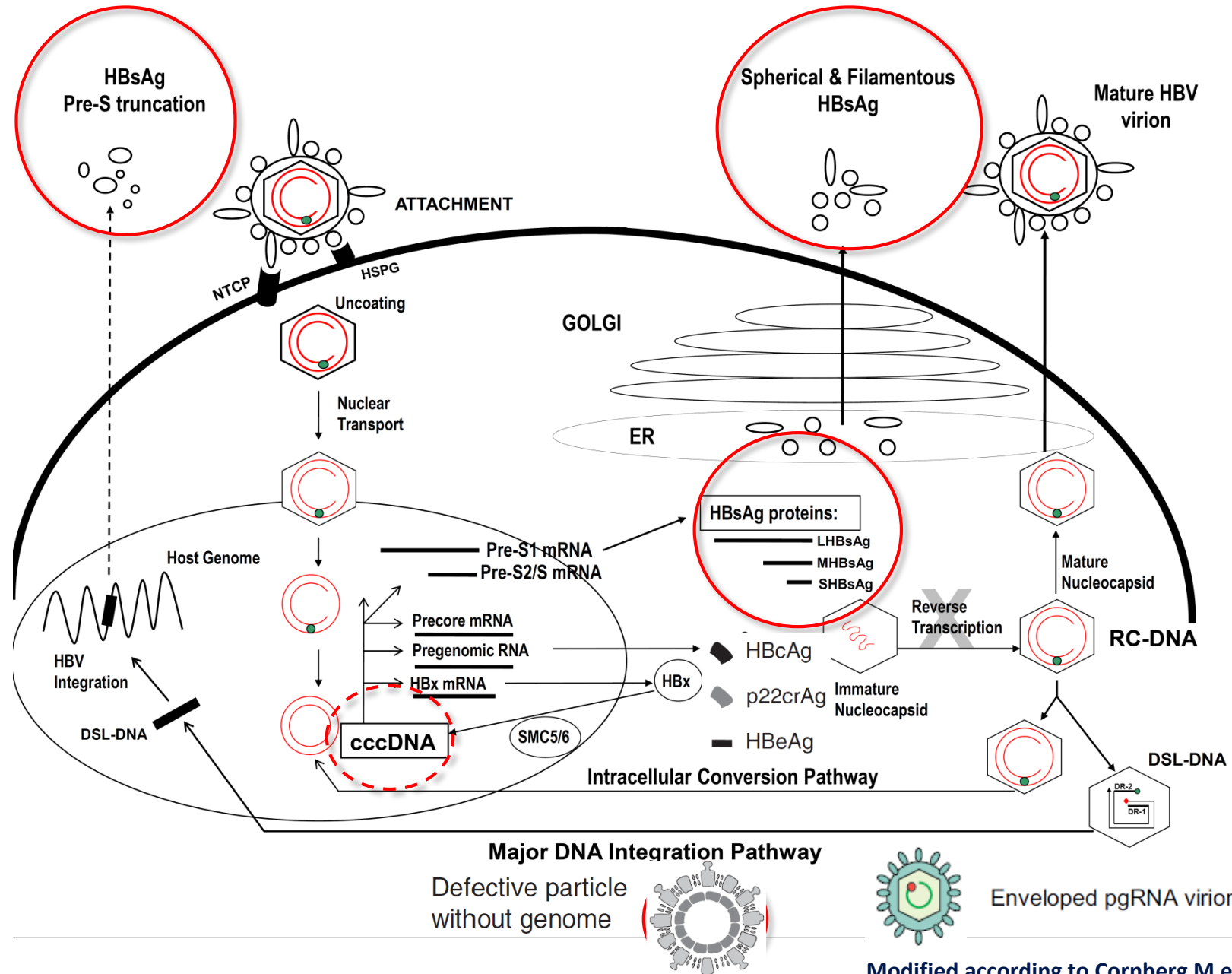


# Effect of NA treatment on serum HBV nucleic acids kinetics

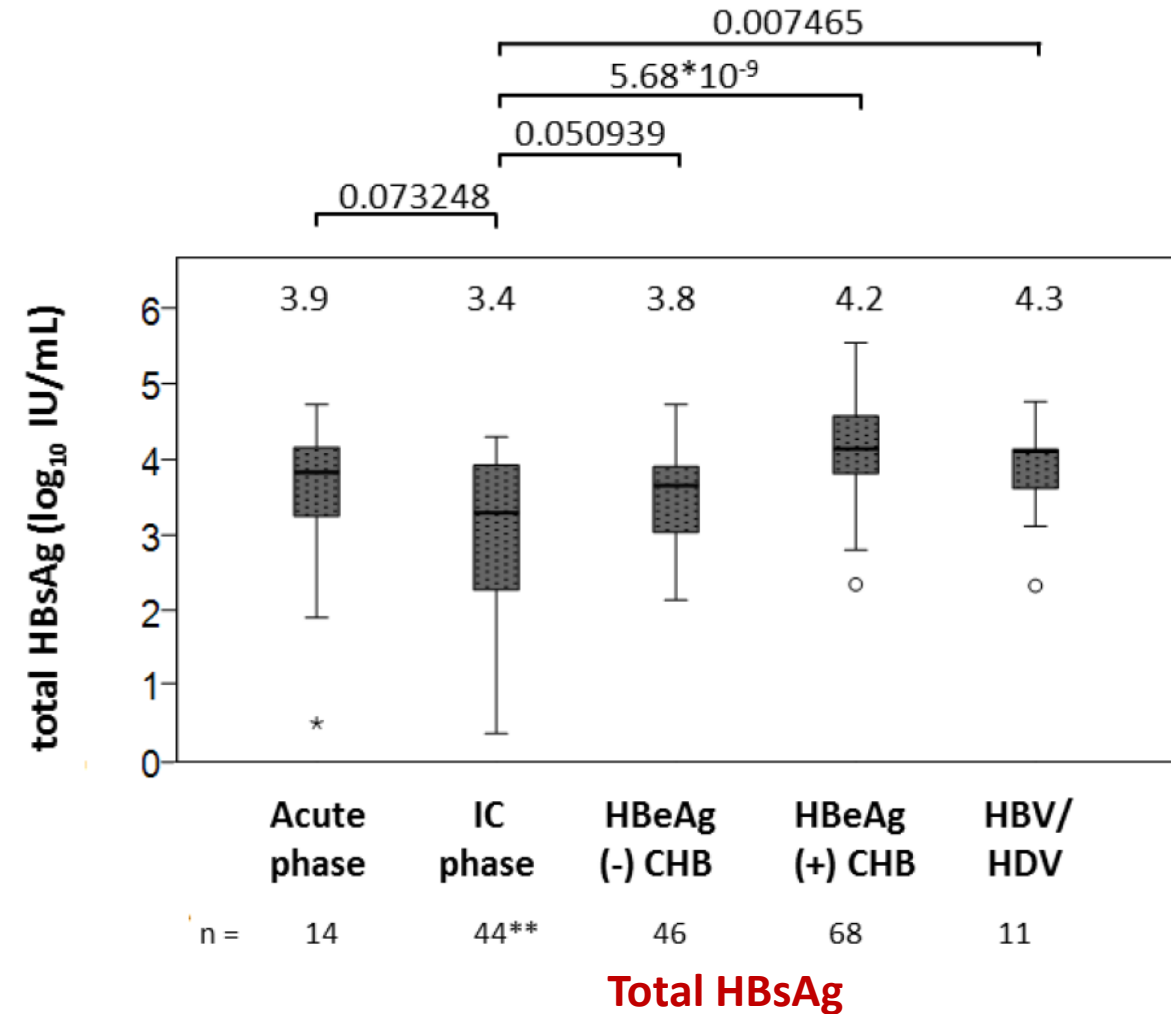
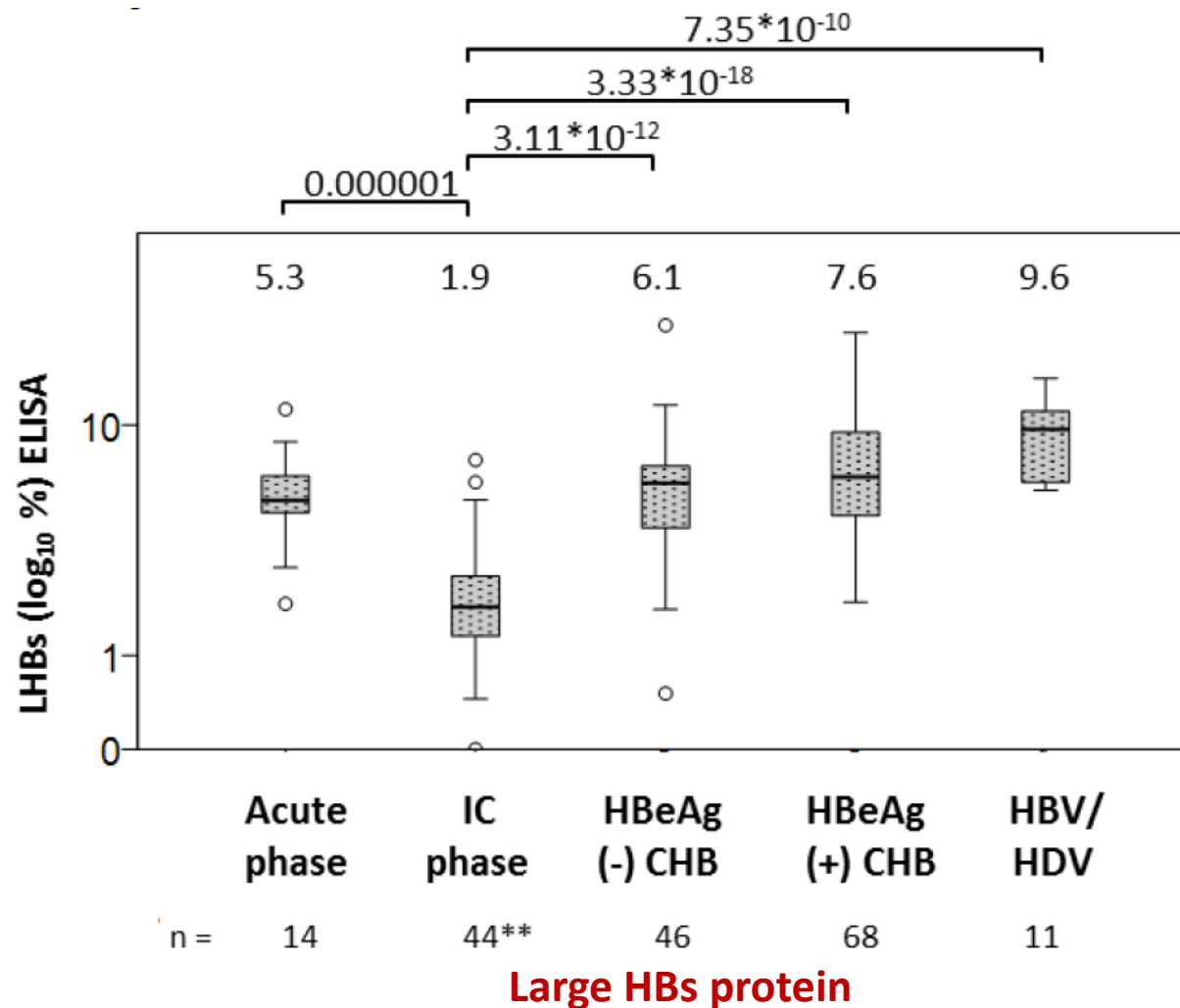


# **HBsAg components**

# Potential biomarkers for assessing endpoints – HBsAg components



# Low-level of large and middle HBsAg protein better describe inactive infection as compared to total HBsAg

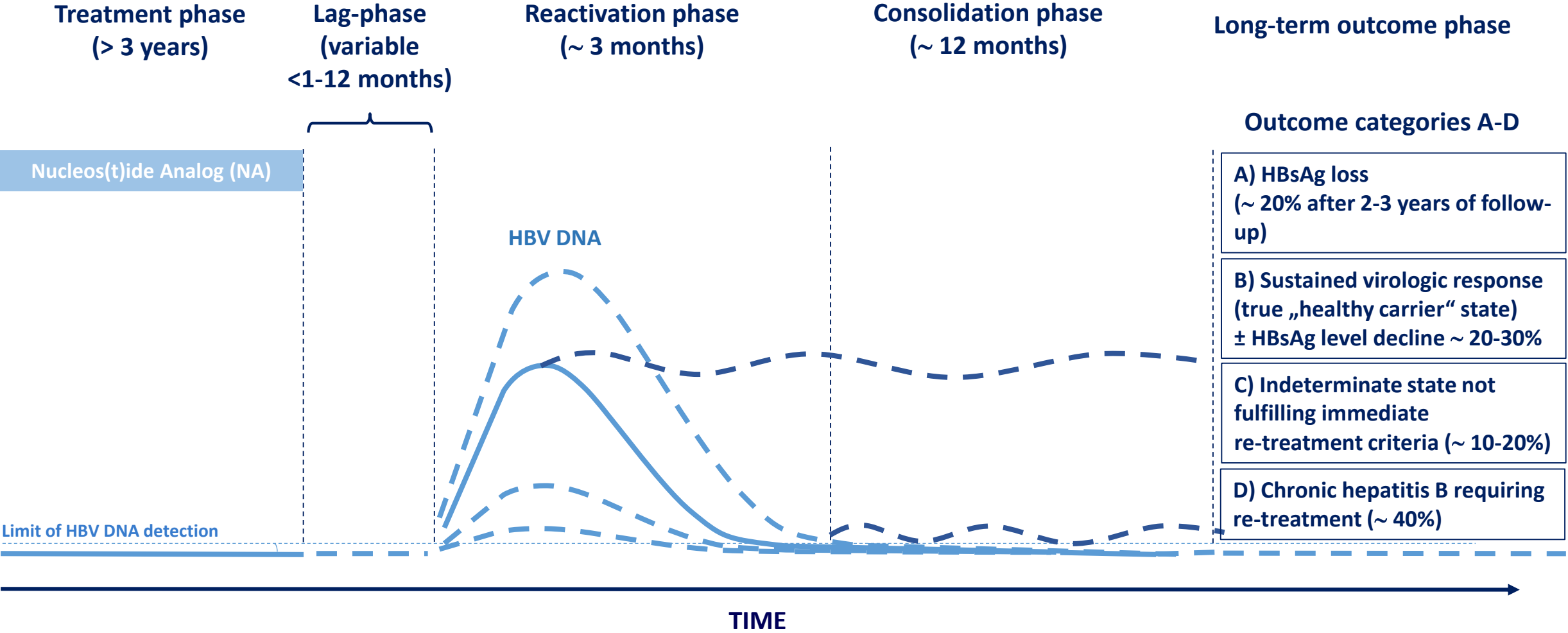


# **EASL Clinical Practice Guideline - Why aiming for HBsAg loss?**

- The main advantage of HBsAg loss is that it allows a safe discontinuation of antiviral therapy**

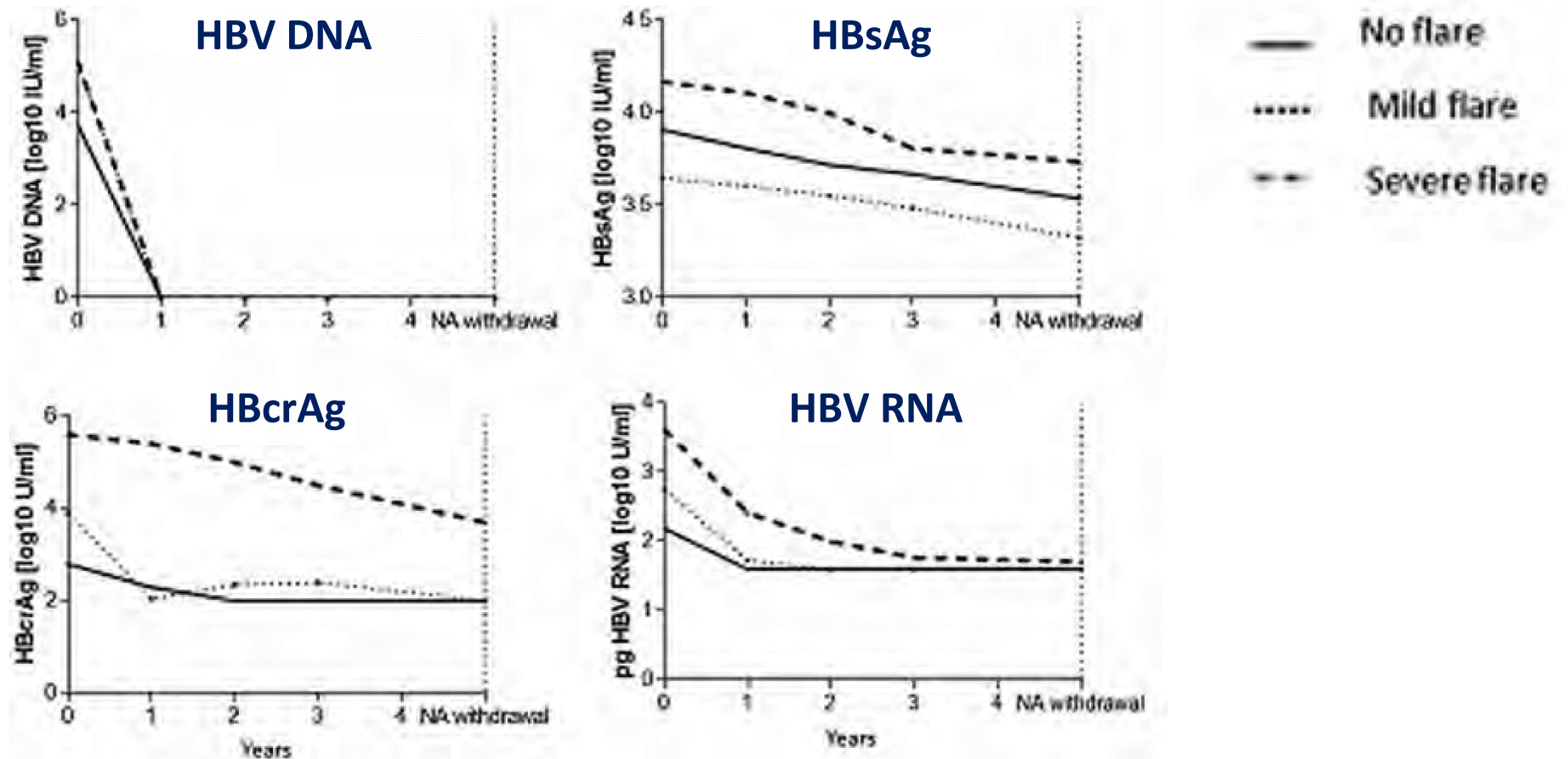
**New biomarkers helpful in predicting outcome after  
stopping a finite course of HBV treatment?**

# NA discontinuation frequently results in virologic and biochemical flares that runs through different phases





# The markers of HBV transcriptional activity-HBcrAg and pgHBV RNA during antiviral therapy with nucleos(t)ide analogue help to predict optimal timing of therapy withdrawal

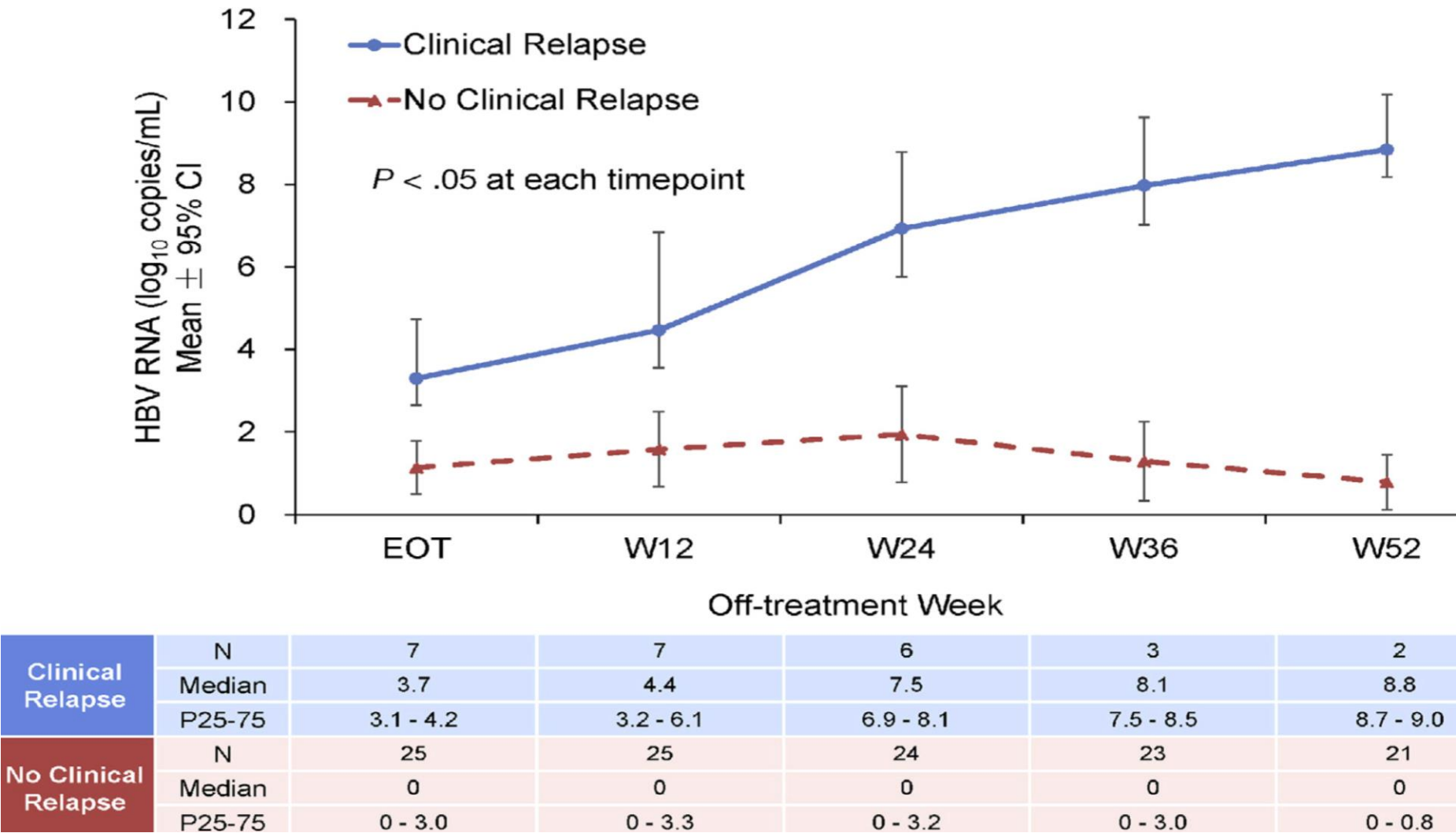


## Association of HBV RNA (pgRNA virion levels) and viral rebound after discontinuation of NUCs

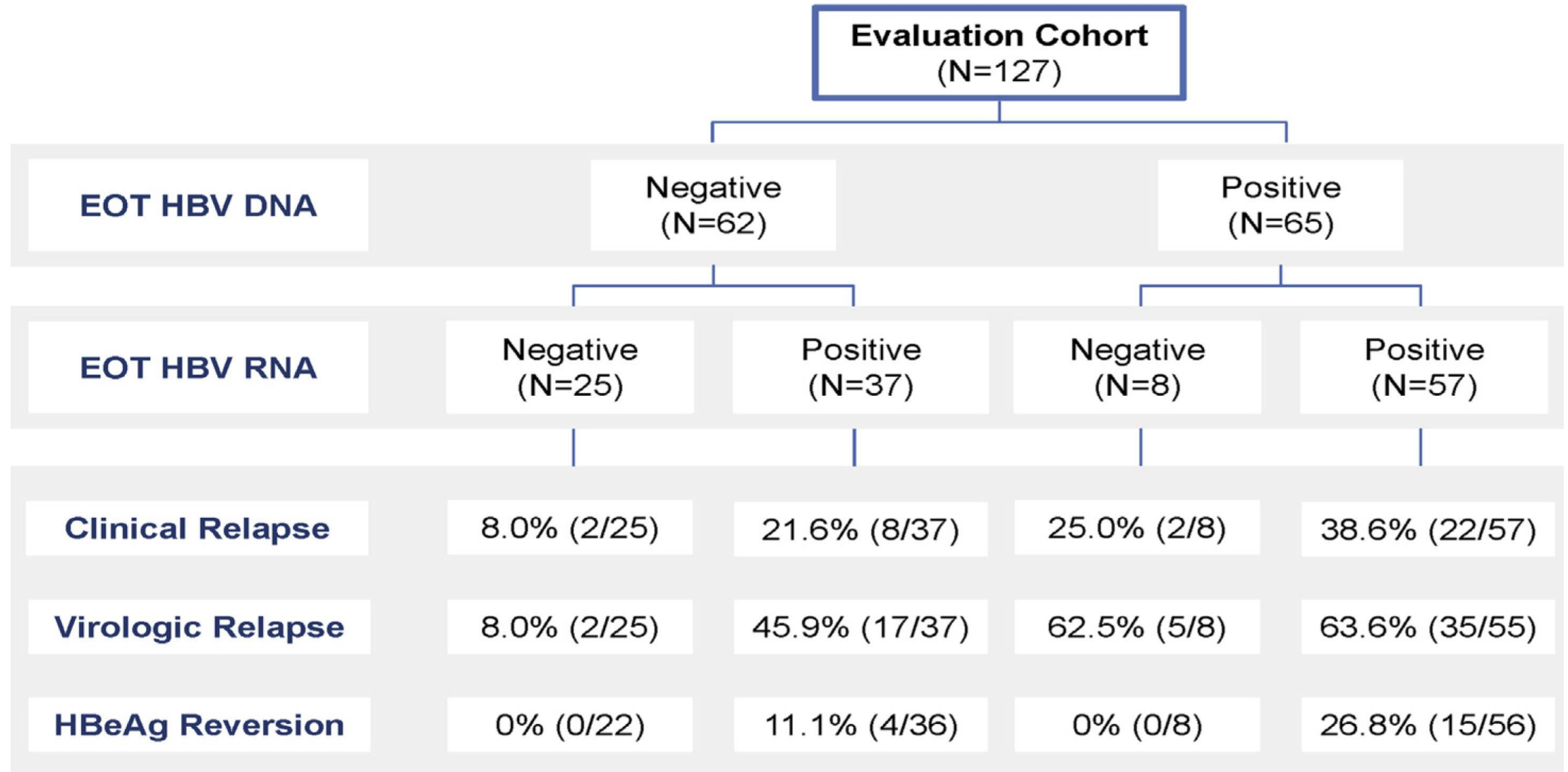
| HBV RNA       | Viral rebound (n) | No viral rebound (n) | Total (n) | * <i>p</i> value |
|---------------|-------------------|----------------------|-----------|------------------|
| Positive      | 21                | 0                    | 21        | 0.001            |
| Below the LoQ | 3                 | 9                    | 12        |                  |
| Total (n)     | 24                | 9                    | 33        |                  |

\*Chi-Square test; n, number of CHB patients.

# End-of-treatment HBV RNA levels predict the risk of relapse after stopping NA in HBeAg positive patients with seroconversion



# End-of-treatment HBV RNA levels predict the risk of relapse after stopping NA in HBeAg pos. patients with seroconversion



# Clinical utility of quantitative anti-HBc levels

- Anti-HBc levels are higher in HBsAg-positive patients as compared to those being anti-HBs-positive (but anti-HBc affinity was lower in chronic infection than during recovery)<sup>1</sup>
- Low anti-HBc levels are associated with HBsAg seroclearance (cut-off < 3log)<sup>2</sup>
- Serum levels of anti-HBc correlate with cccDNA positivity in OBI<sup>3</sup>
- Anti-HBc levels (cut-off  $\geq 6.41$  IU/ml) were significantly associated with high risk of HBV reactivation during immunosuppressive therapy<sup>4</sup>

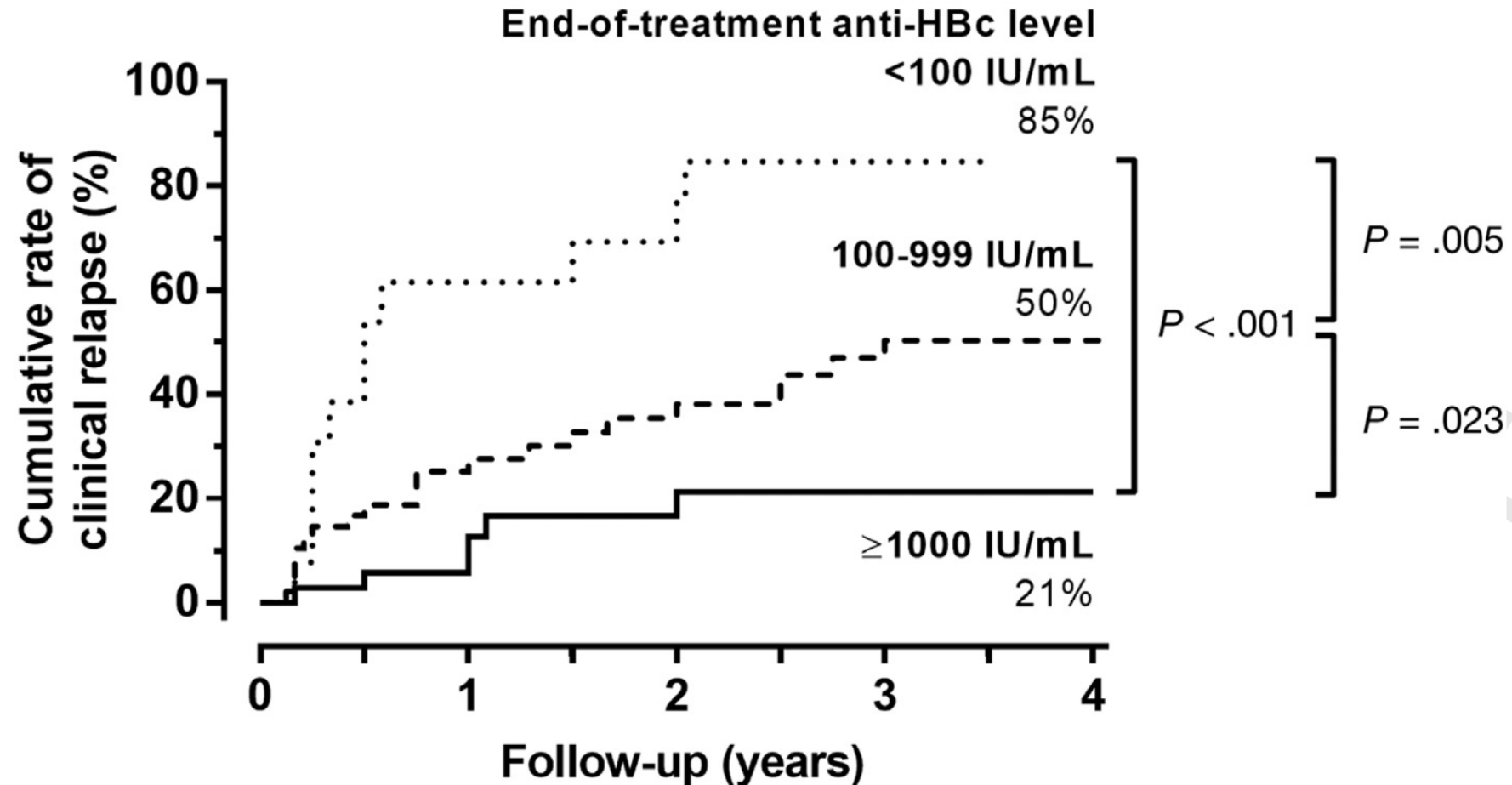
<sup>1</sup>Han et al. *J Clin Virol.* 2011;52(4):295-9

<sup>2</sup>Hu et al. *Clin Gastroenterol Hepatol.* 2018 [Epub ahead of print]

<sup>3</sup>Caviglia et al. *J Hepatol.* 2018;69(2):301-307

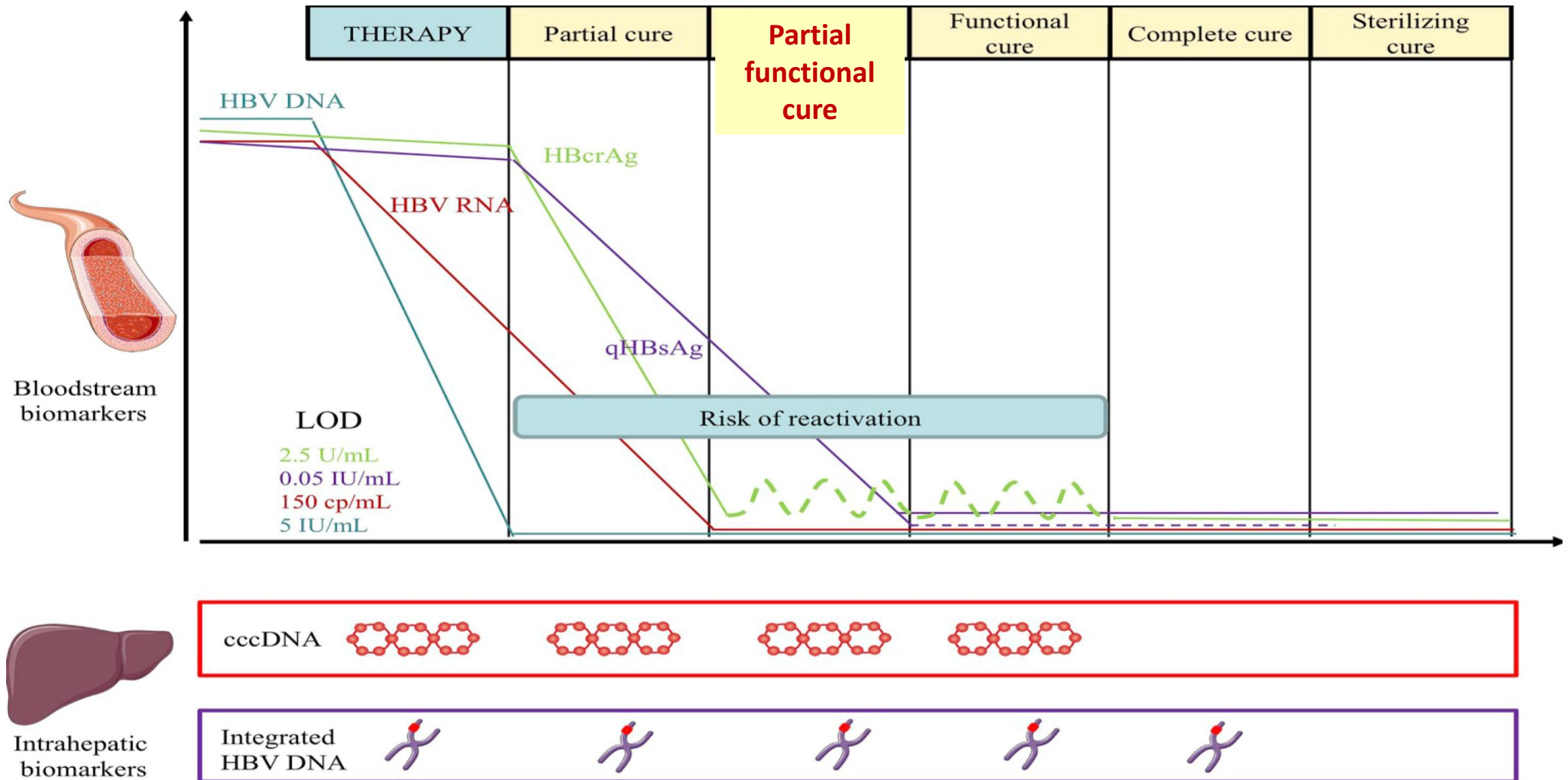
<sup>4</sup>Yang et al. *J Hepatol.* 2018 Aug;69(2):286-292.

# Serum level of antibodies against hepatitis B core protein is associated with clinical relapse after discontinuation of nucleos(t)ide analogue therapy

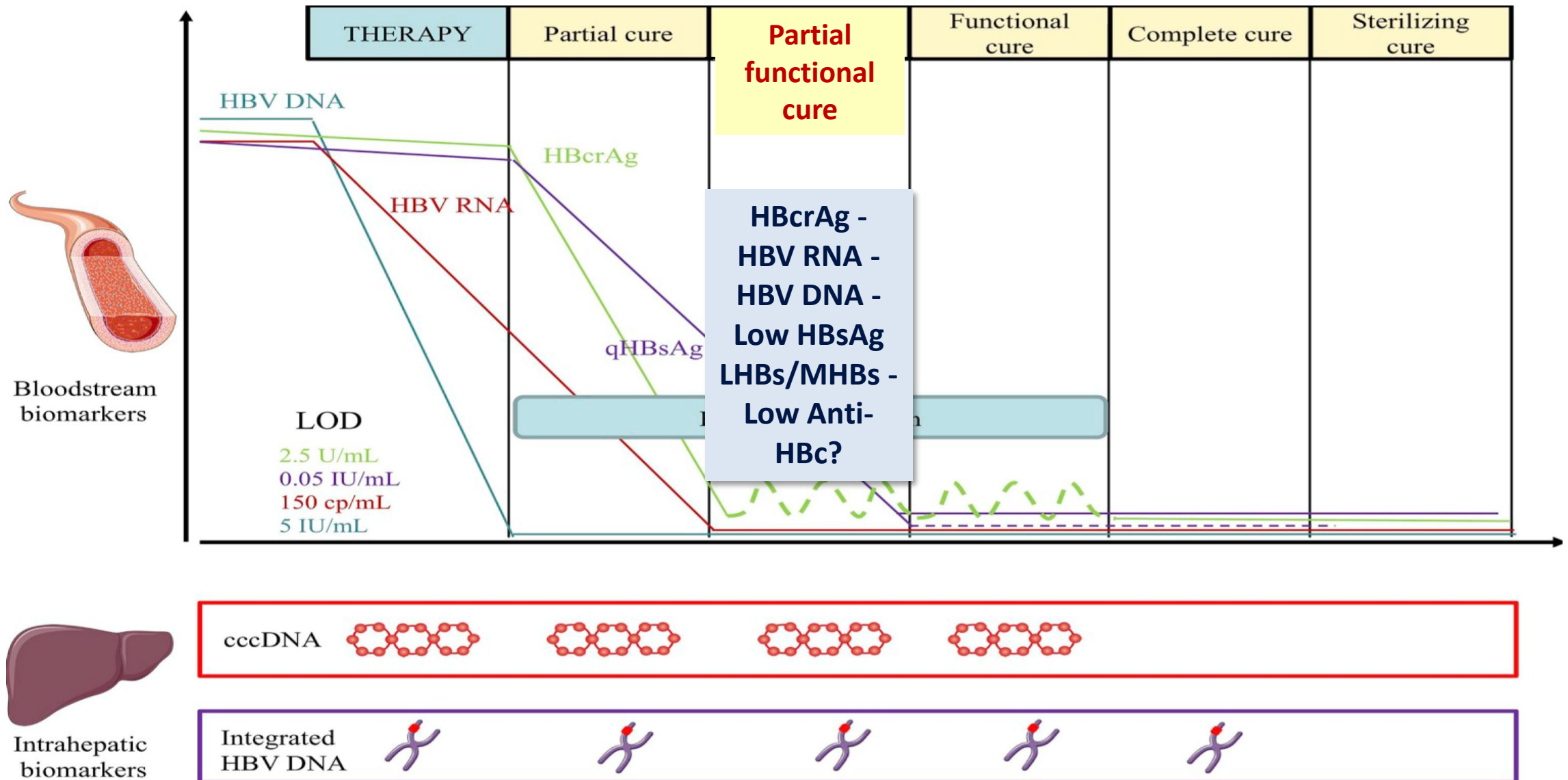




# Classification of HBV-cure with new biomarkers

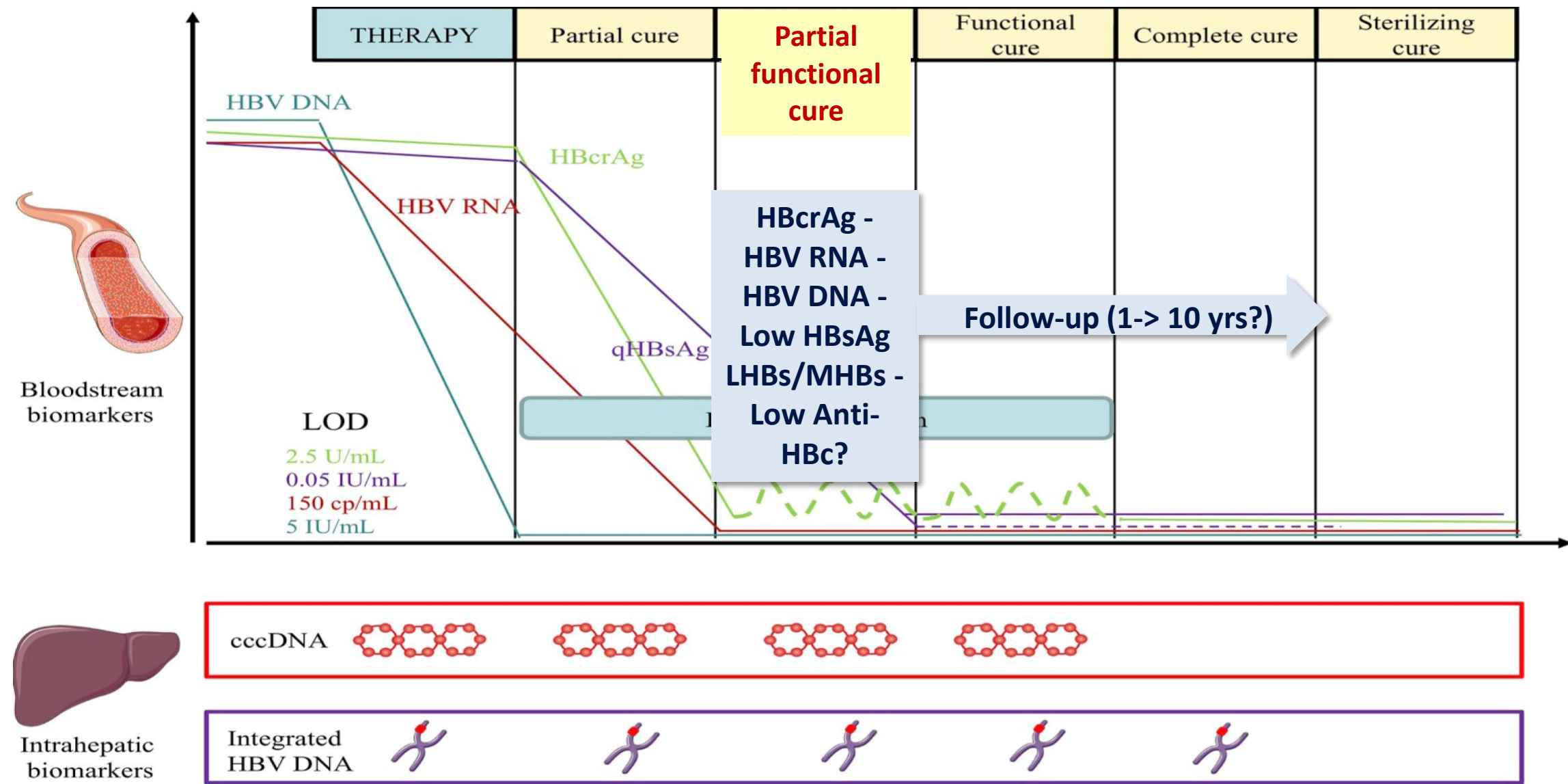


# Classification of HBV-cure with new biomarkers





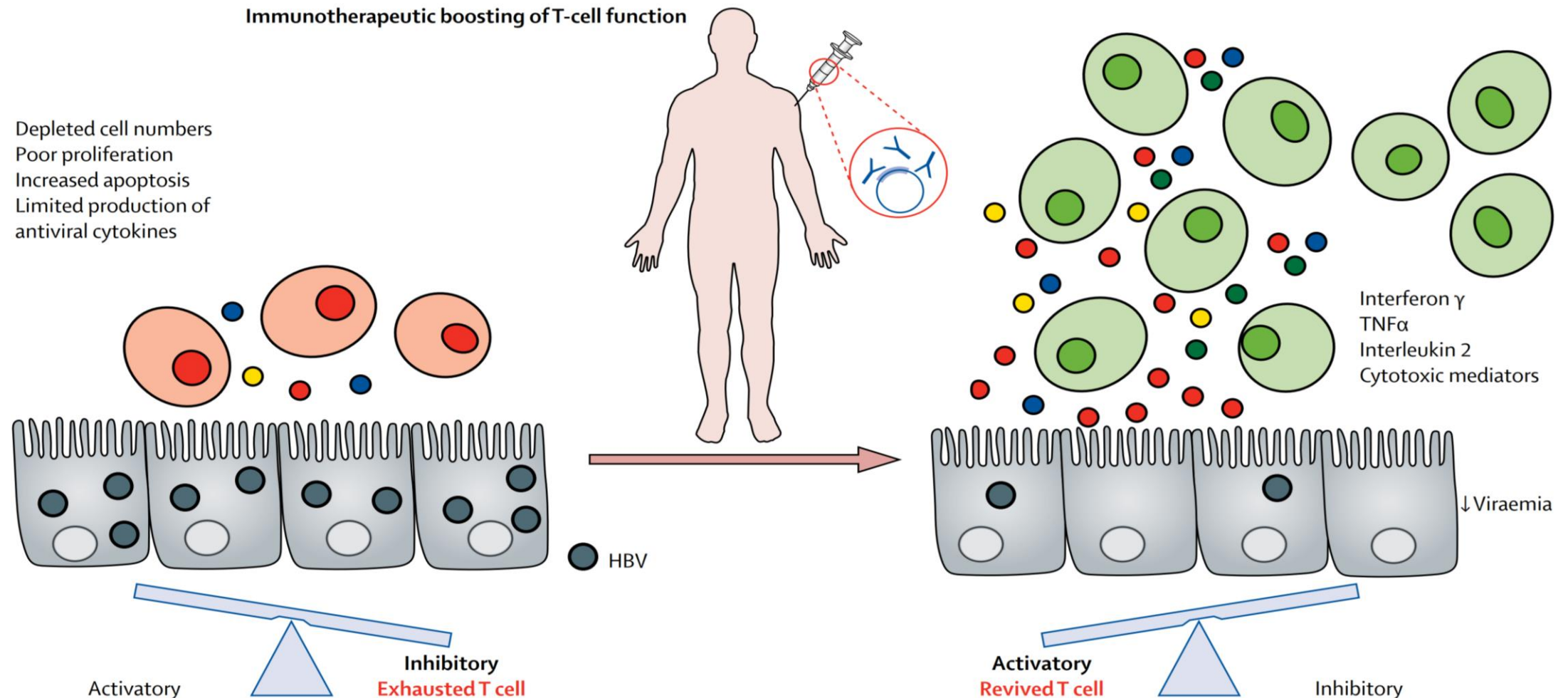
# Chance for transition into a true functional or even complete cure stage after having achieved a partial functional cure?



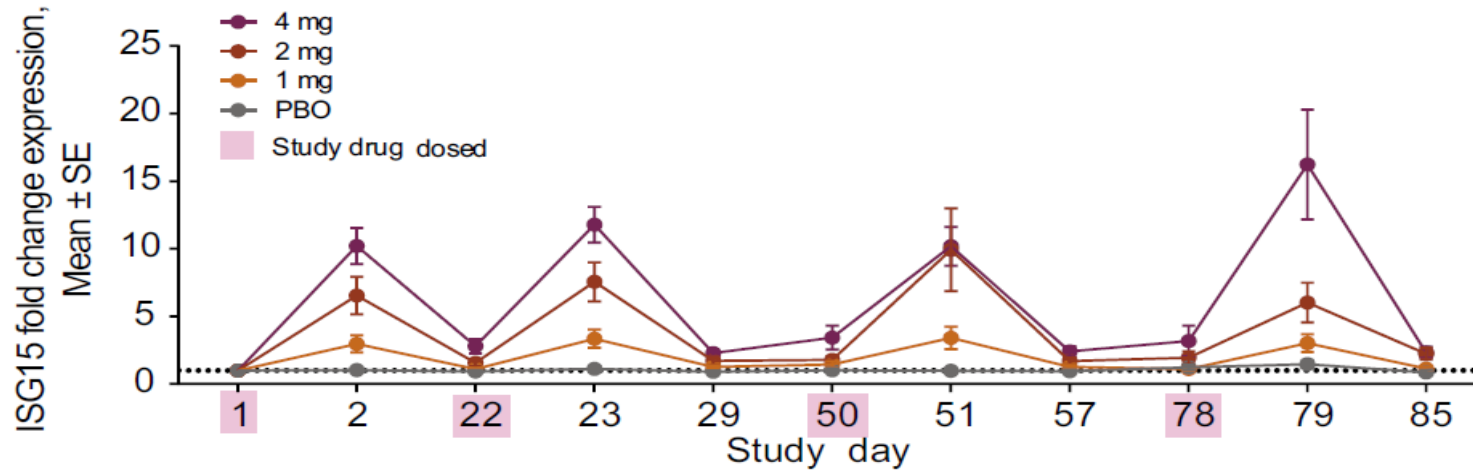


# Immunotherapeutic augmentation of therapeutic vaccination

Expand a population of functional T cells, able to clonally expand and produce antiviral cytokines

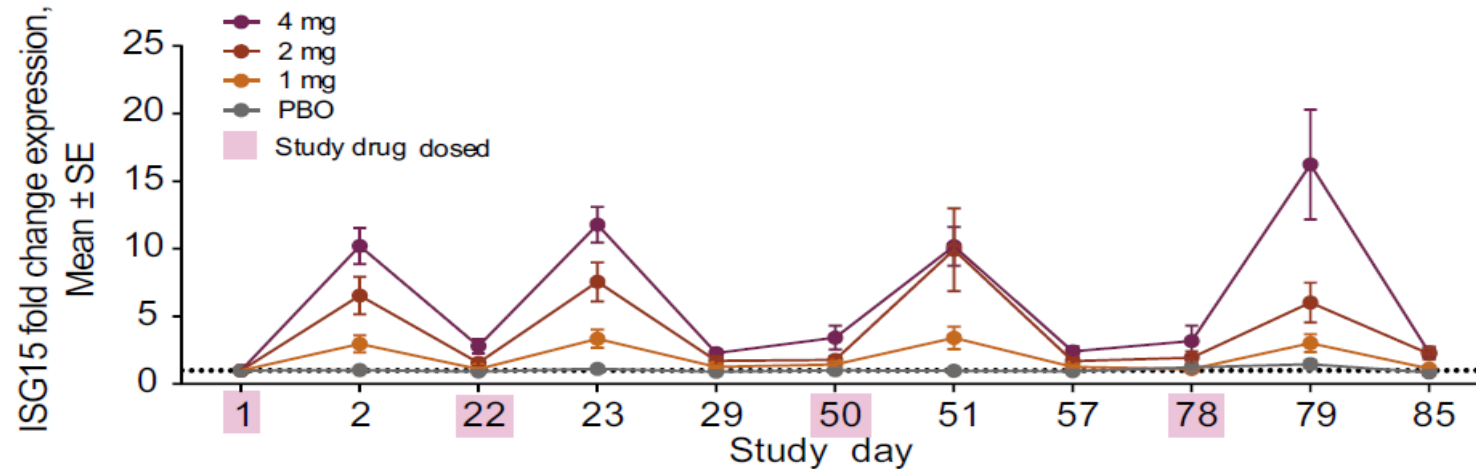


# TLR7-Agonist Vesatolimod to Treat Chronic Hepatitis B (Phase-II-Study)

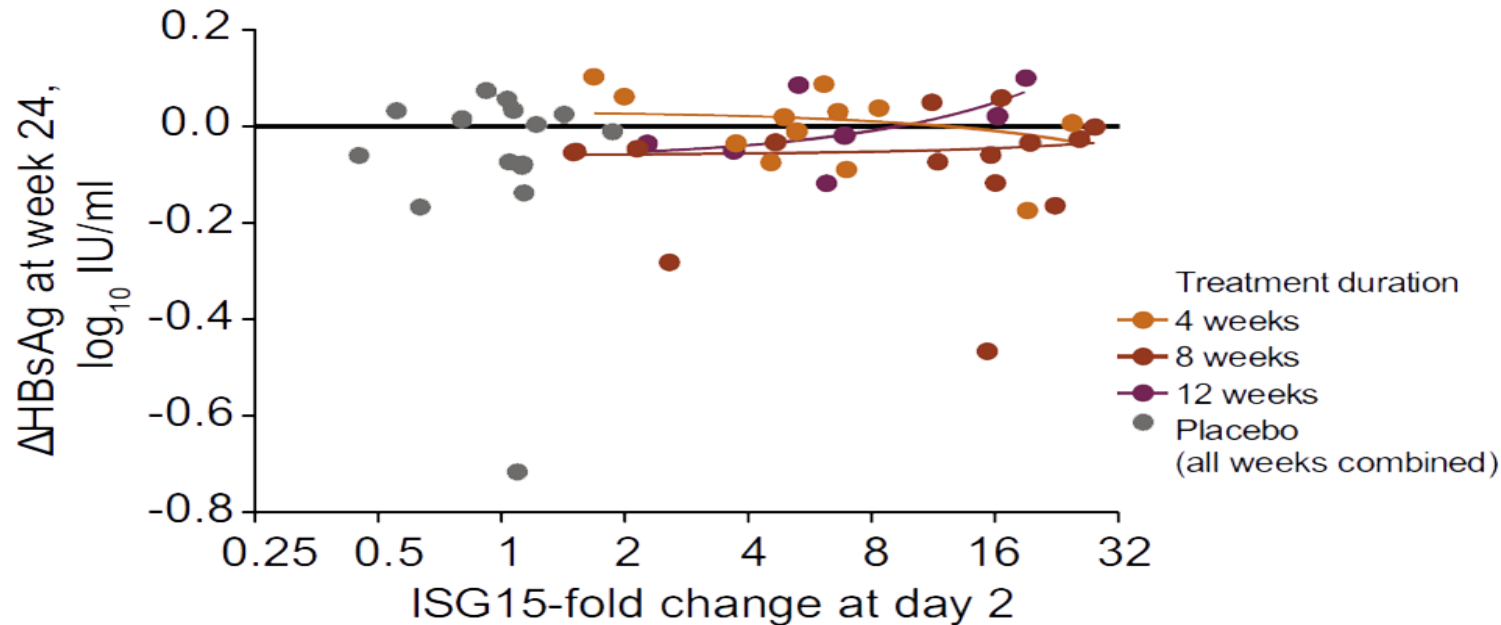


Clear-cut dose-dependent induction of ISG15 expression (ISG15-mRNA, fold changes) under TLR7-stimulation

# TLR7-Agonist Vesatolimod to Treat Chronic Hepatitis B (Phase-II-Study)

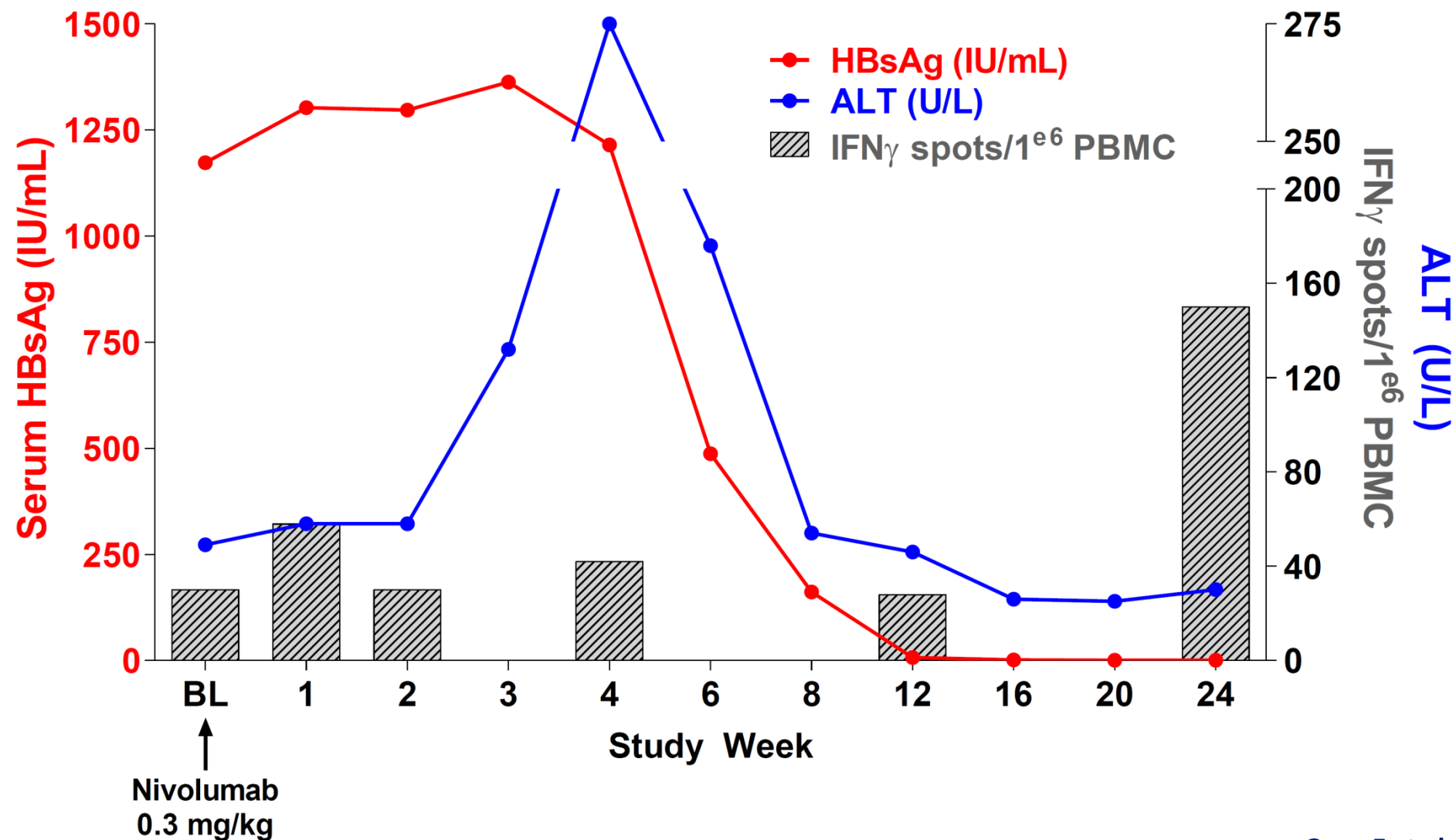


Clear-cut dose-dependent induction of ISG15 expression (ISG15-mRNA, fold changes) under TLR7-stimulation



But ....  
ISG15 induction not associated with HBsAg decline at week 24

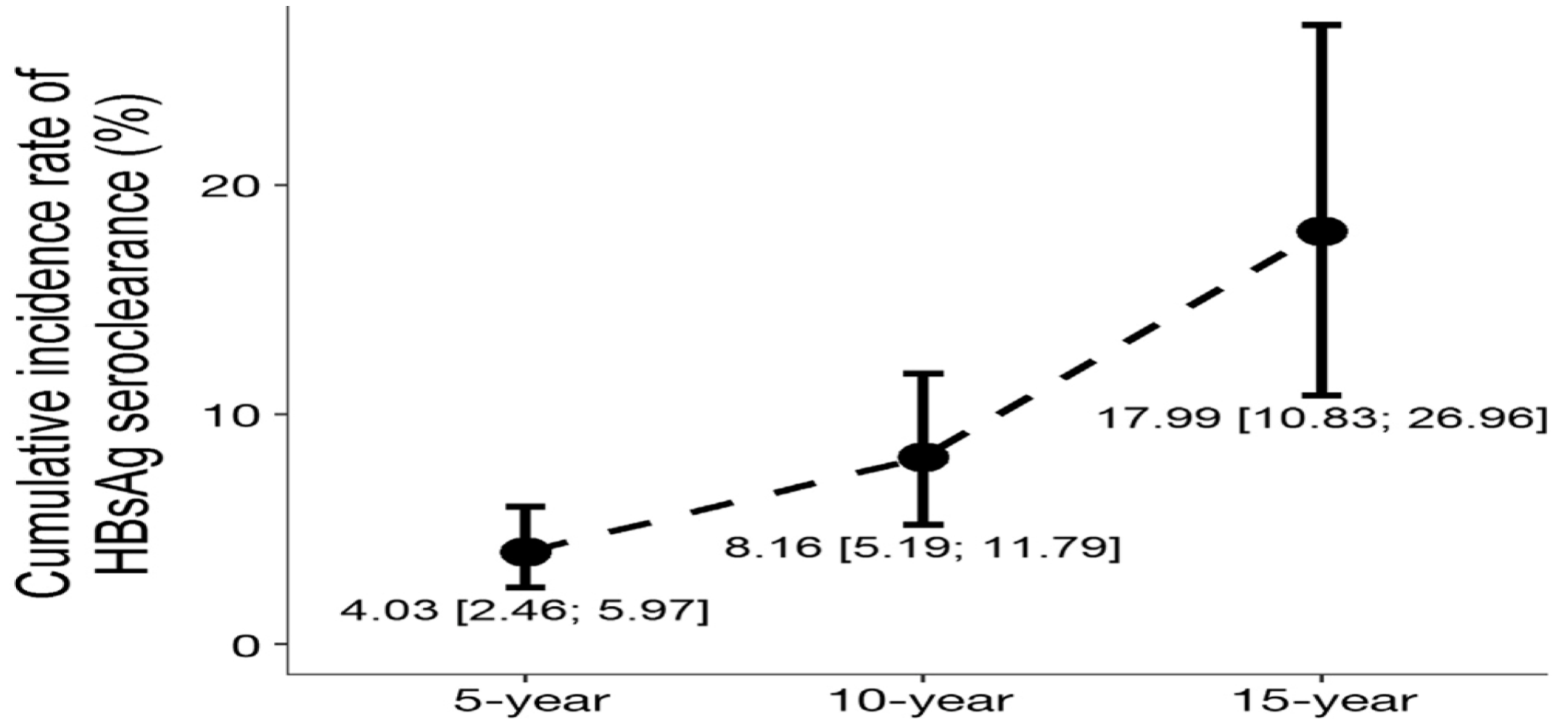
# Anti-PD1 (immune checkpoint) blockade with nivolumab w/wo therapeutic vaccine in virally suppressed patients





# Inzidenz der HBsAg Serumclearance

Yeo YH et al. Gastroenterology 2019;156(3):635-646



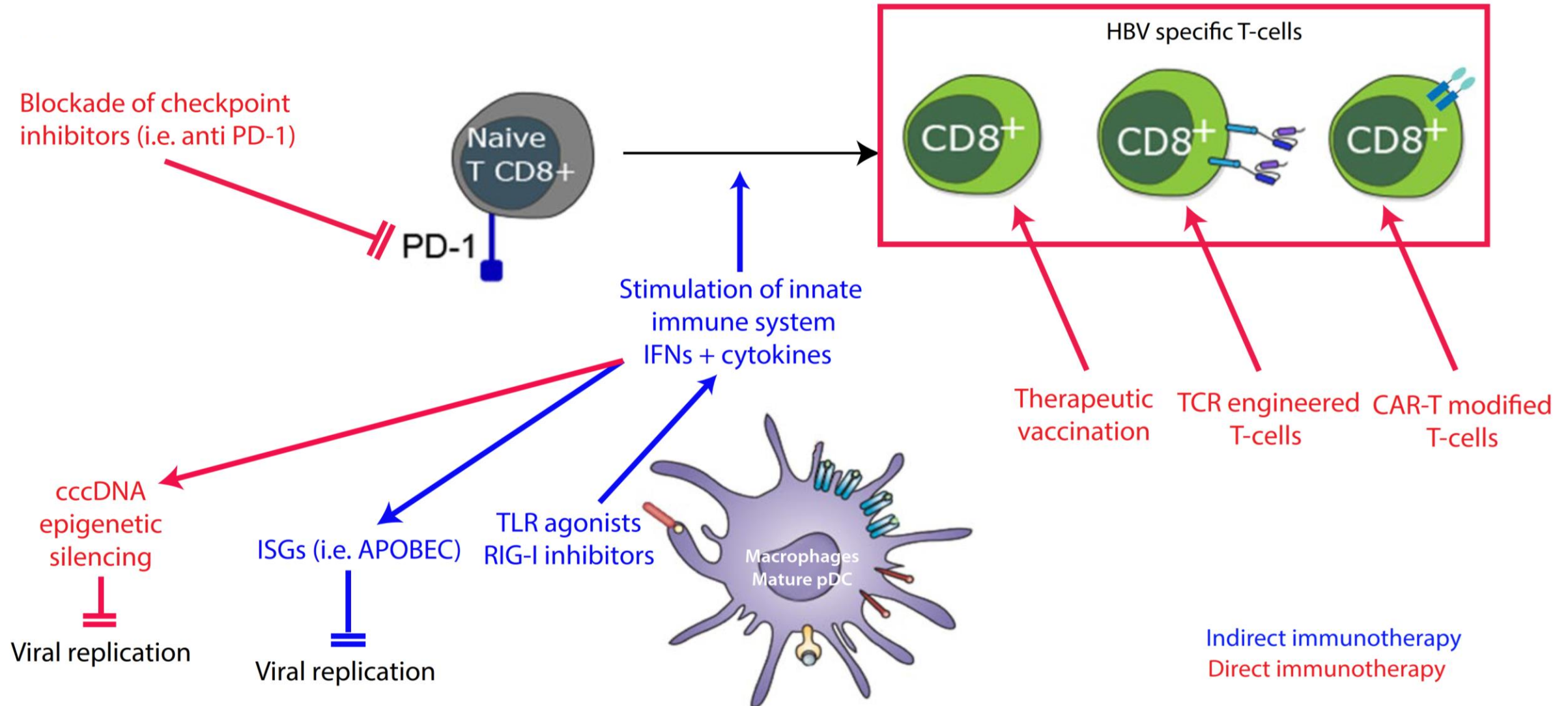


# Hepatitis B core-related antigen

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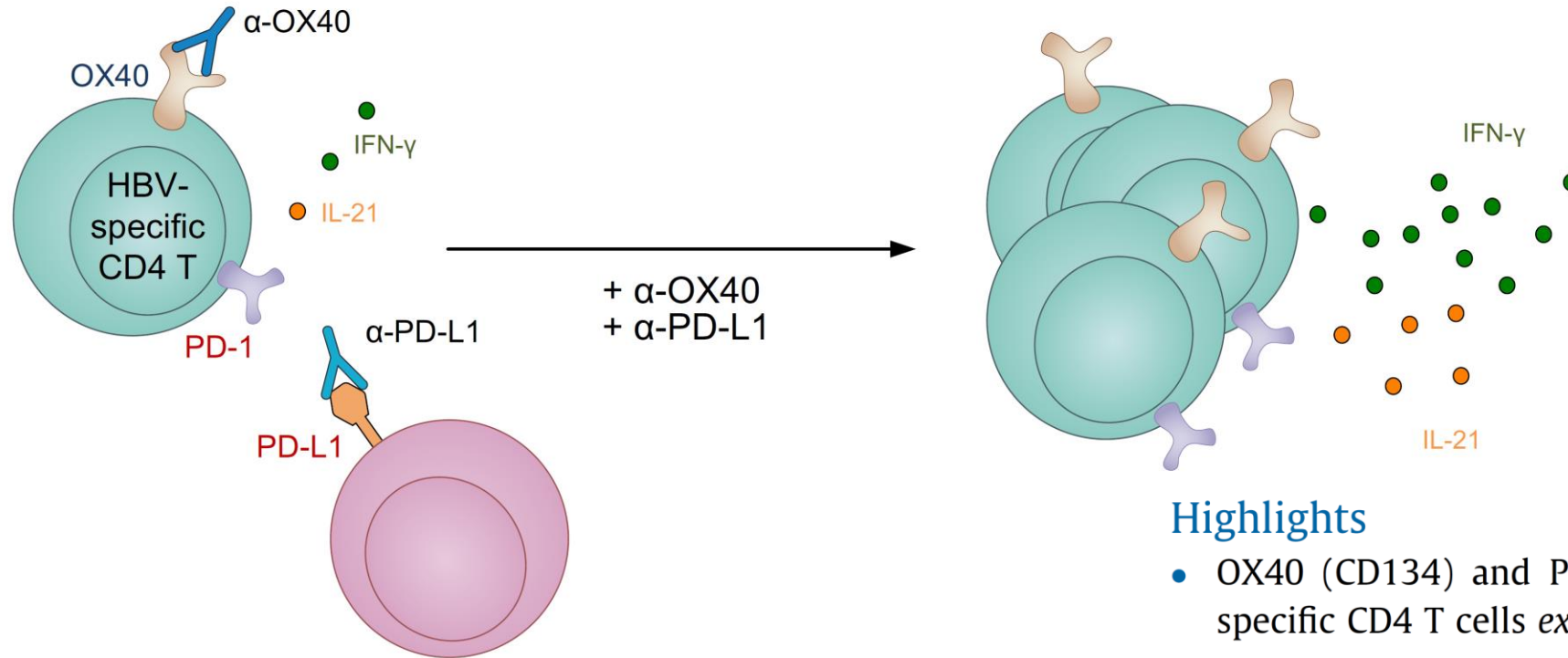
- Composite marker comprises HBcAg, HBeAg and a 22kDa truncated core-related protein (p22cr)
- Available since 2014 fully automated quantitative assay (Lumipulse)
- Not influenced by translation of integrated sequences
- Surrogate marker for cccDNA and its transcriptional activity
- Sensitivity problems (3log)

# HBV replication cycle and main targets of viral therapies



# Augmenting HBV-specific CD4 T cells - it takes two to tango

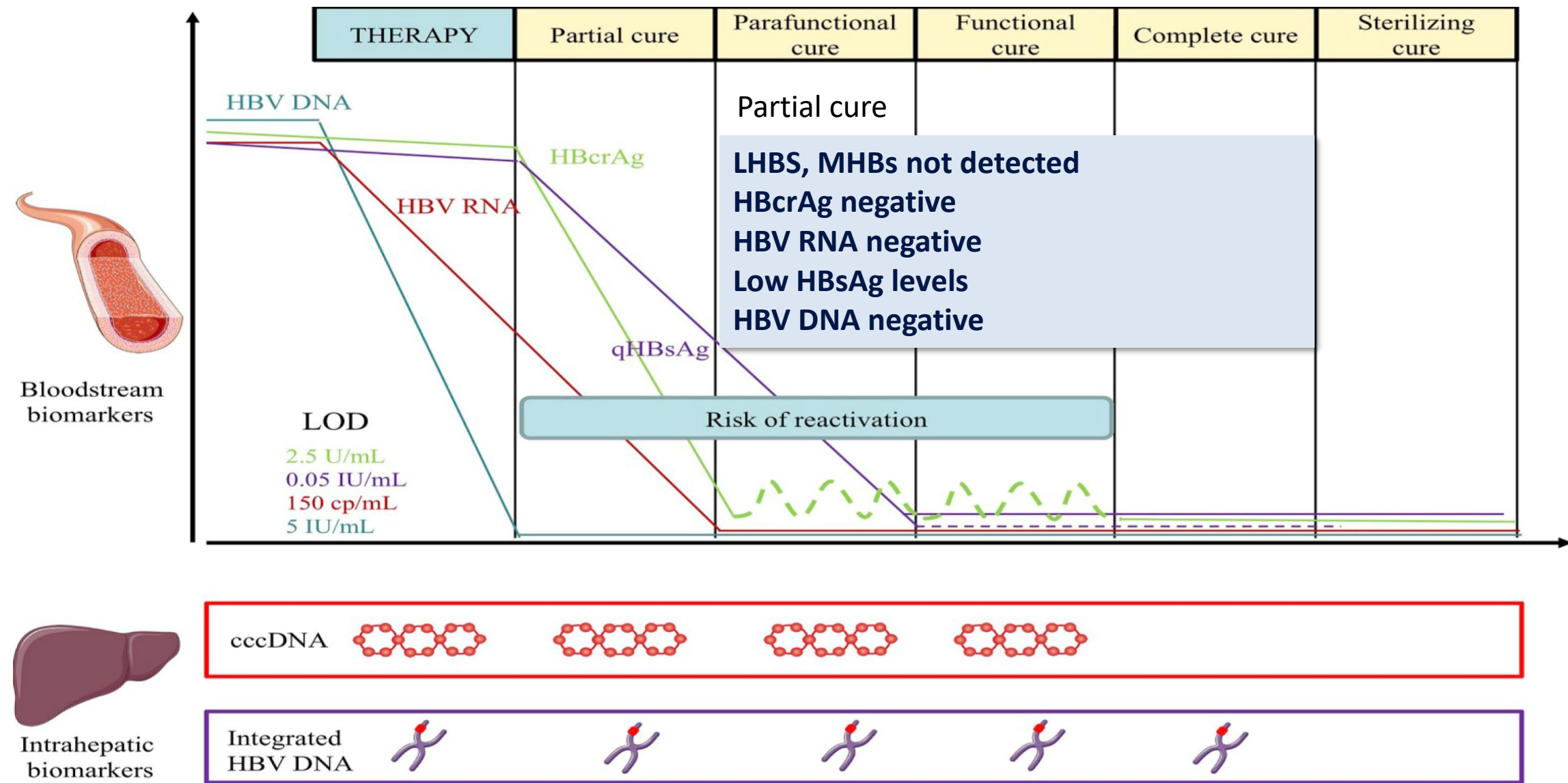
Functionally augmented HBV-specific CD4 T cells were observed only when combining OX40 stimulation and PD-L1 blockade



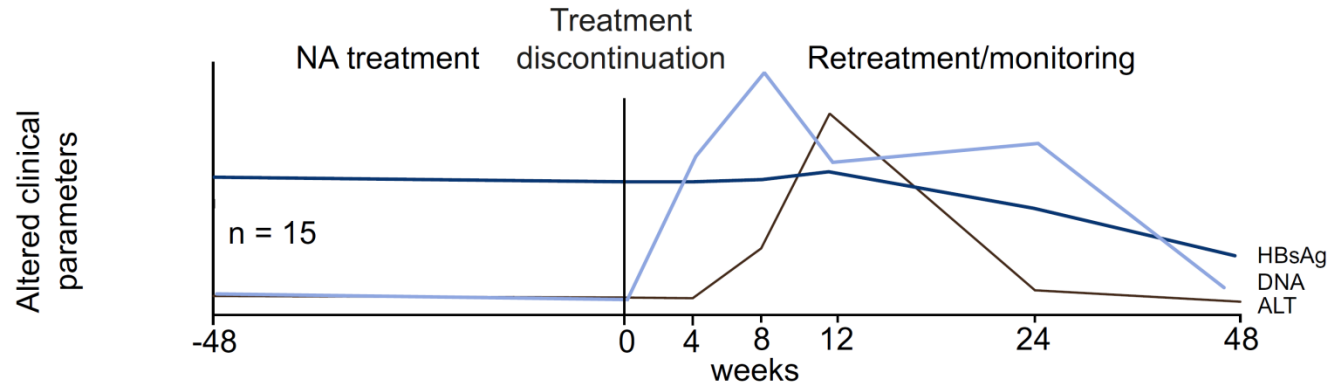
## Highlights

- OX40 (CD134) and PD-1 are strongly expressed on HBV-specific CD4 T cells *ex vivo*.
- The HBV-specific CD4 T cells predominantly target the polymerase and core proteins.
- Combined OX40 stimulation and PD-L1 blockade functionally augment HBV-specific CD4 T cells.

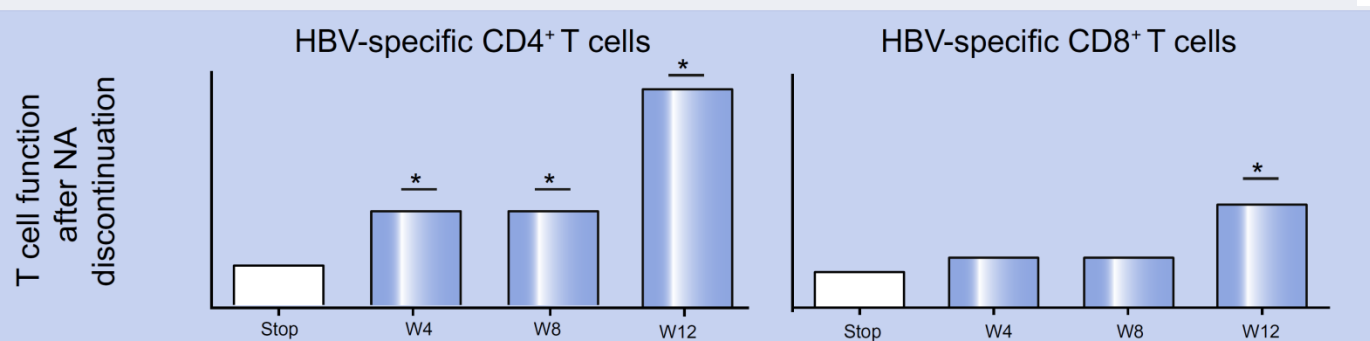
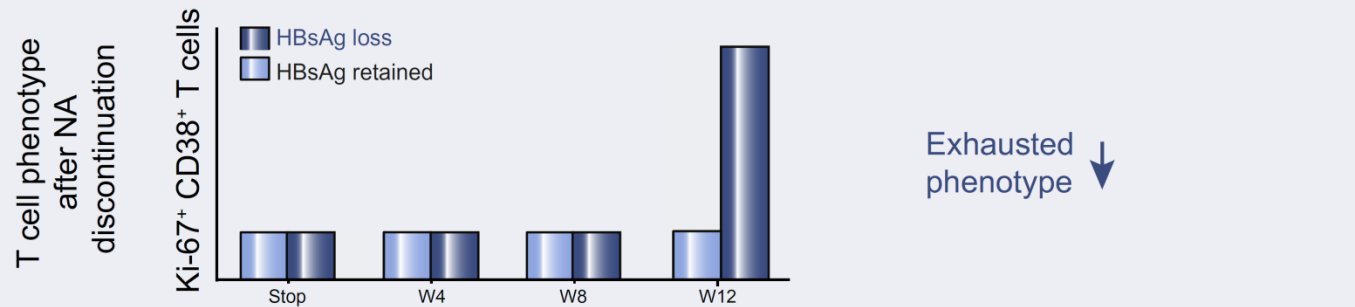
# Does HBsAg positive „cure“ of the disease and disease associated outcomes exist? How to best define HBsAg positive cure of active infection



# HBV-specific T cell responses after stopping NA

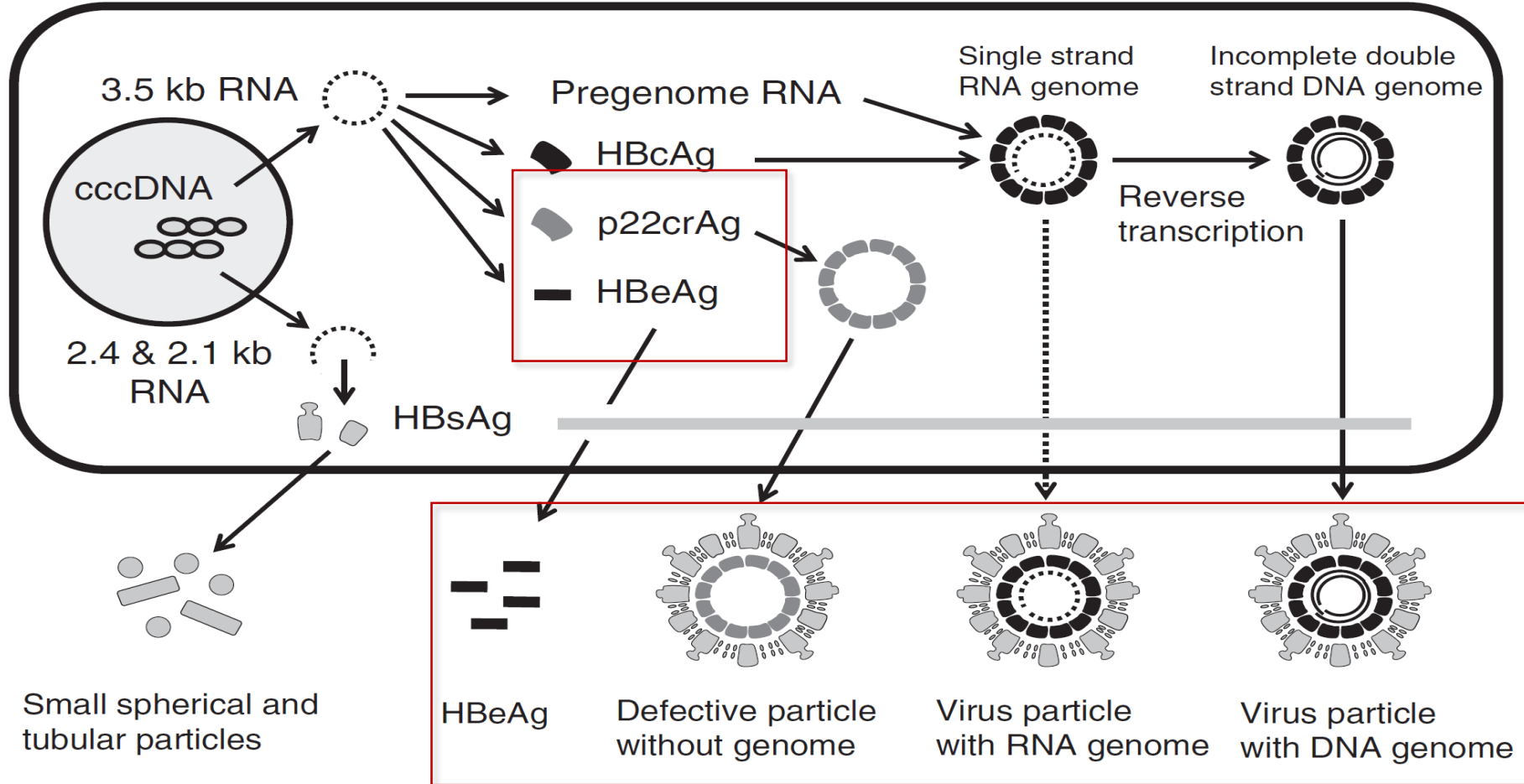


- Discontinuation of NA therapy leads to higher HBsAg loss rates in HBeAg-negative CHB patients.
- T cells from patients with subsequent HBsAg loss show a less exhausted phenotype.
- These T cells also express higher levels of activation and proliferation markers at week 12 after discontinuation of therapy.
- Relapse of active HBV replication may trigger immunological environment that enhances responsiveness of HBV-specific T cells *in vitro*.



# Secreted factors of hepatitis B virus

## Secreted particles and proteins considered by the measurements of HBcrAg versus HBsAg



**Hepatitis B Virus DNA-negative Dane particles containing a 22-kDa precore protein  
(exceeding Dane particles by ~ 100 fold)**

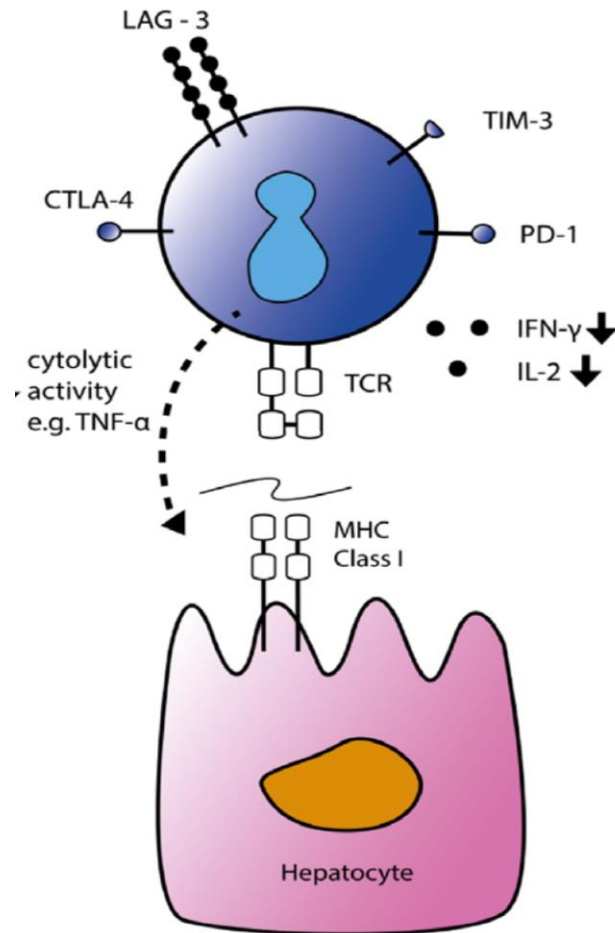
Kimura T et al. JBC 2005; 280: 21713; Luckenbaugh L et al. J Viral Hepat 2015; 22: 561

**Figure modified according to Tanaka E and Matsumoto A. Hepatol Res 2014; 44: 1-8**

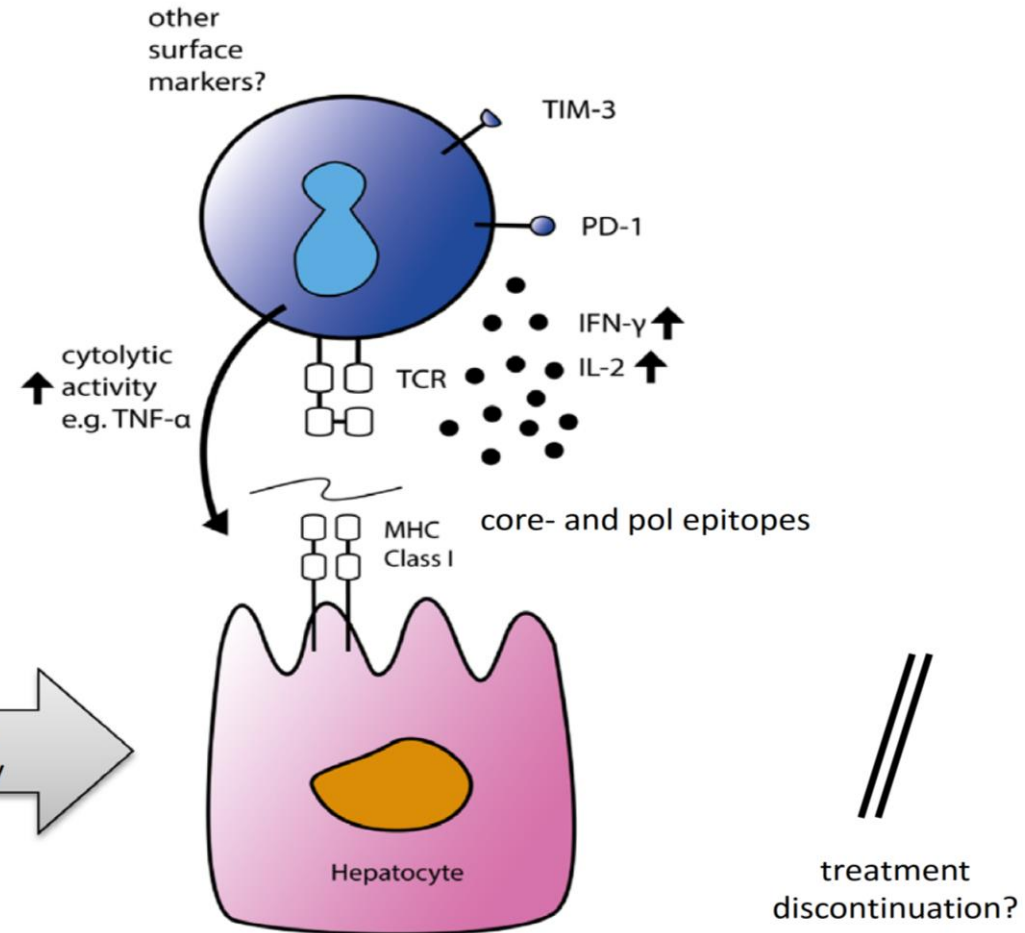


# Possible restoration of HBV specific CD8+ T-cell function

Chronic HBeAg negative  
HBV infection



Possible restoration of  
HBV-specific CD8<sup>+</sup> T cell  
responses

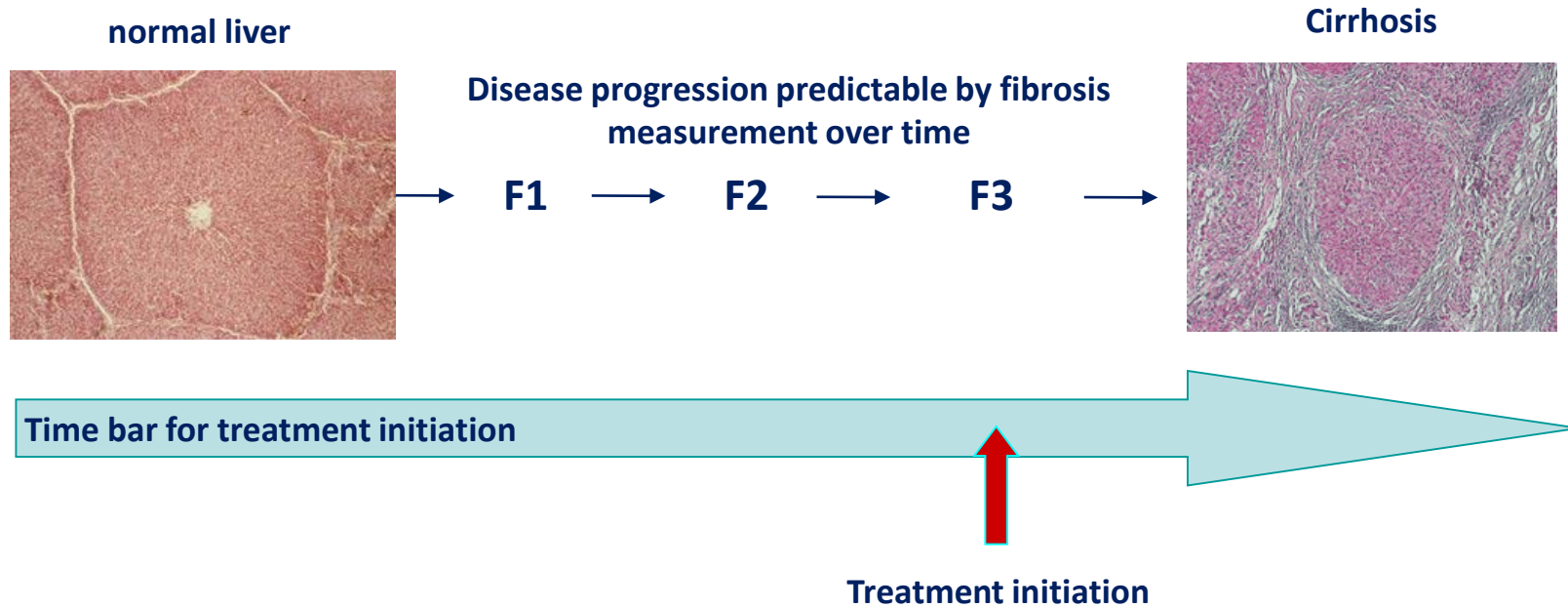


long-term  
NA therapy

treatment  
discontinuation?

# The HBV journey

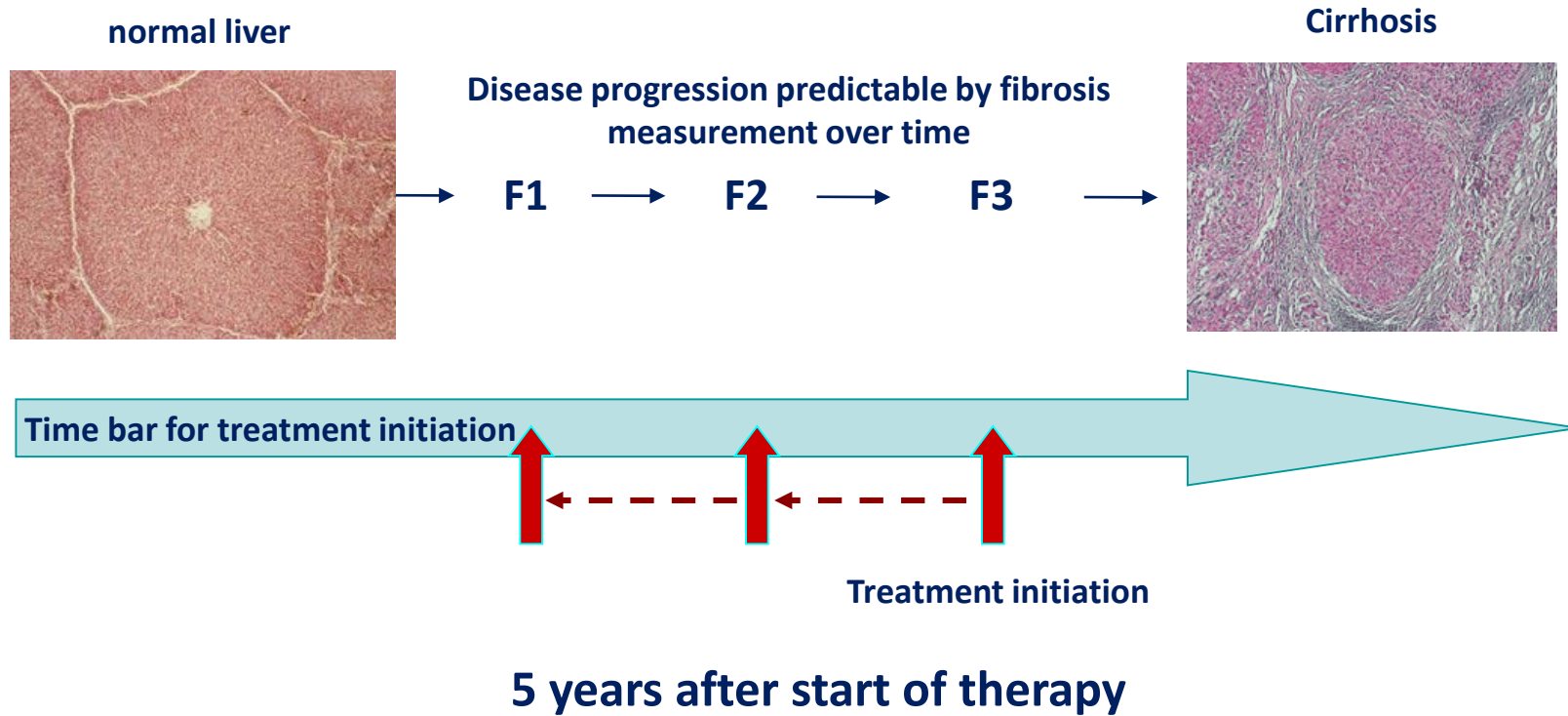
HBV infection = A fibrotic liver disease





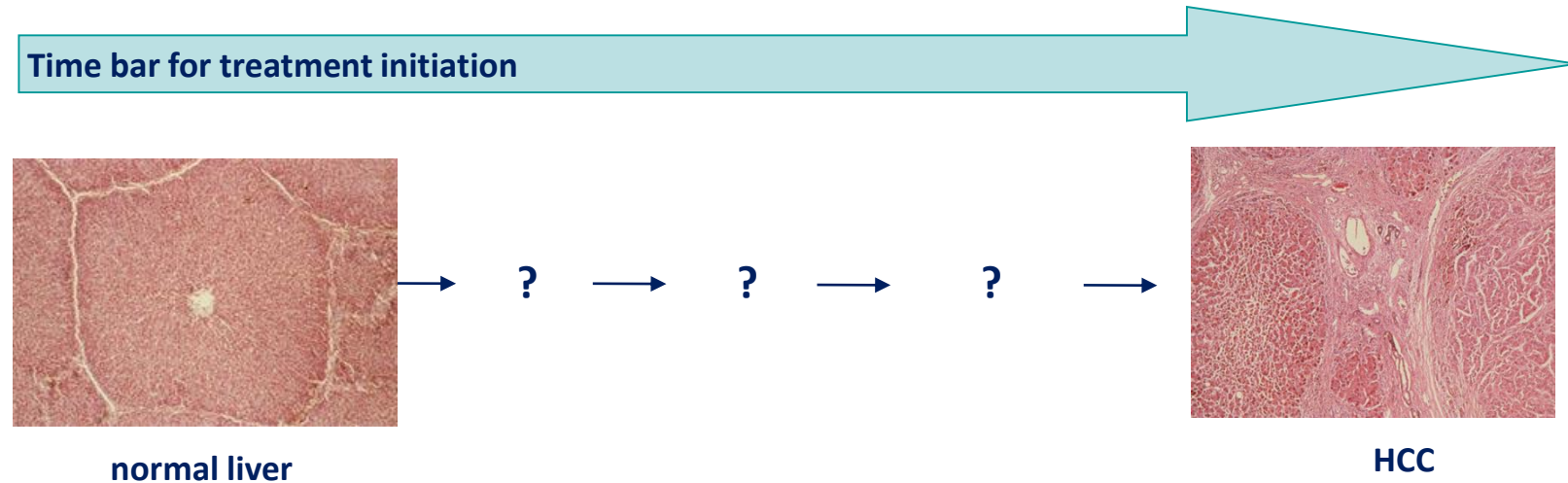
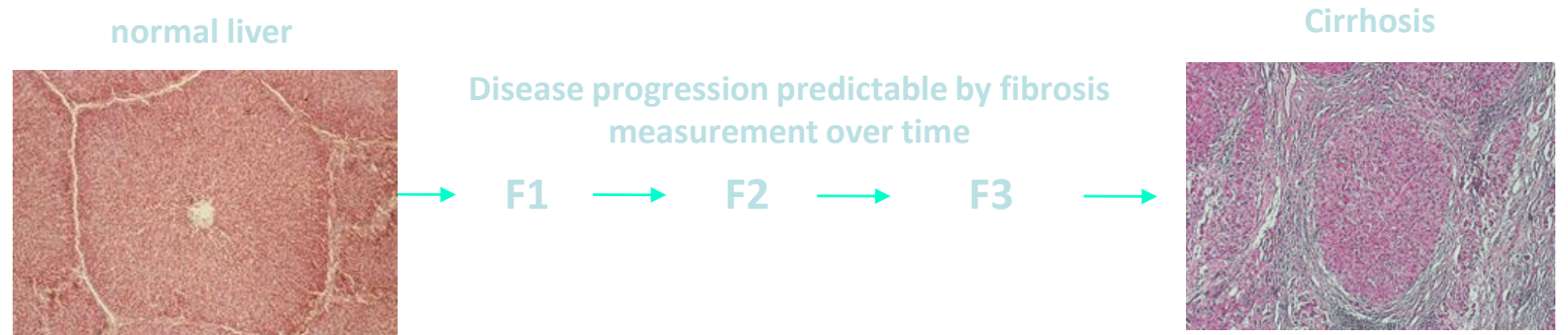
# The HBV journey

HBV infection = A fibrotic liver disease



# The HBV journey

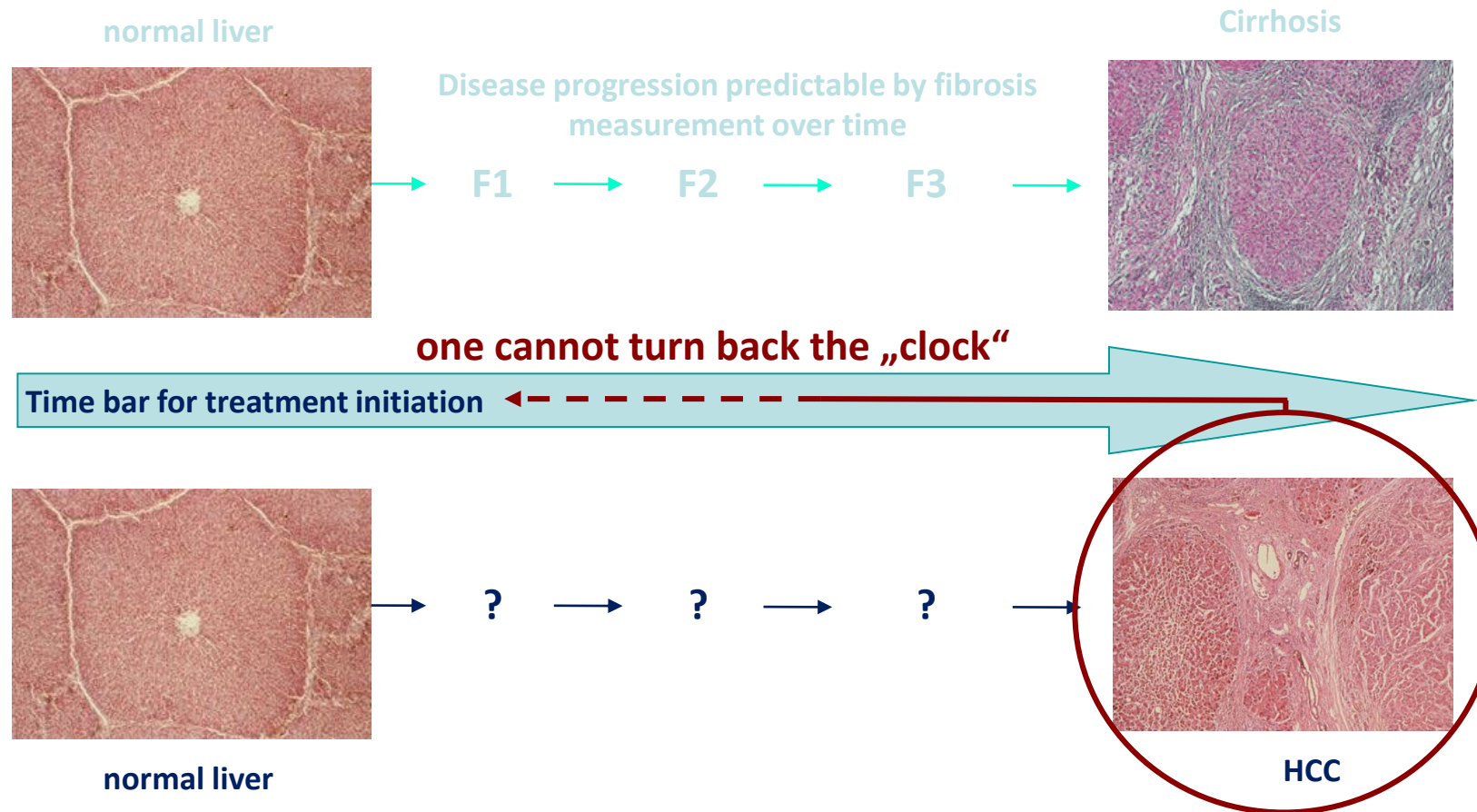
HBV infection = A fibrotic liver disease



HBV infection = An oncogenic disease

# The HBV journey

HBV infection = A chronic liver disease



HBV infection = An oncogenic disease

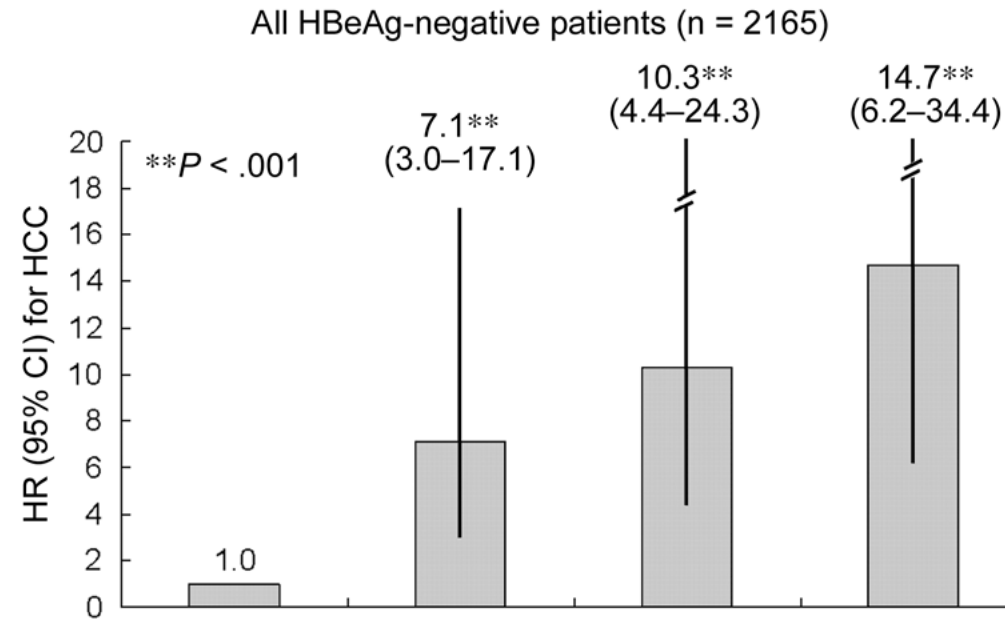
# Risikostratifizierung des HCC bei HBeAg-negativer HBV Infektion durch Kombination viraler Biomarker

Tseng TC et al. *JID* 2013;208; 584-593

| Risk group         | Minimal<br>(Reference) |                 | Medium      |                 | Medium high     |                    | High           |
|--------------------|------------------------|-----------------|-------------|-----------------|-----------------|--------------------|----------------|
| HBV DNA<br>(IU/mL) | <2000                  | 2000-<br>19 999 | <2000       | 2000-<br>19 999 | 2000-<br>19 999 | 20 000-<br>199 999 | $\geq$ 200 000 |
| HBsAg (IU/mL)      | <1000                  | <100            | $\geq$ 1000 | 100-<br>999     | $\geq$ 1000     |                    |                |
| Patient number (%) | 662 (30.6)             |                 | 674 (31.1)  |                 | 501 (23.1)      |                    | 328 (15.2)     |

# Risikostratifizierung des HCC bei HBeAg-negativer HBV Infektion durch Kombination viraler Biomarker

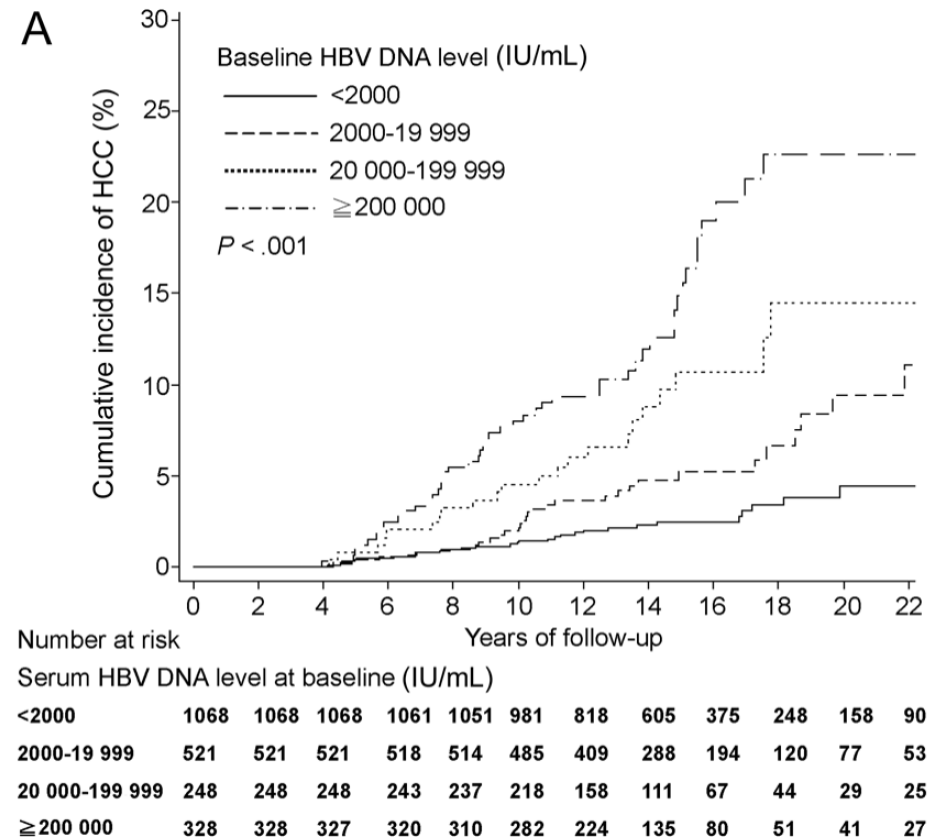
Tseng TC et al. *JID* 2013;208; 584-593



| Risk group         | Minimal<br>(Reference) |                 | Medium     |                 | Medium high     |                    | High       |
|--------------------|------------------------|-----------------|------------|-----------------|-----------------|--------------------|------------|
| HBV DNA<br>(IU/mL) | <2000                  | 2000-<br>19 999 | <2000      | 2000-<br>19 999 | 2000-<br>19 999 | 20 000-<br>199 999 | ≥200 000   |
| HBsAg (IU/mL)      | <1000                  | <100            | ≥1000      | 100-<br>999     | ≥1000           |                    |            |
| Patient number (%) | 662 (30.6)             |                 | 674 (31.1) |                 | 501 (23.1)      |                    | 328 (15.2) |

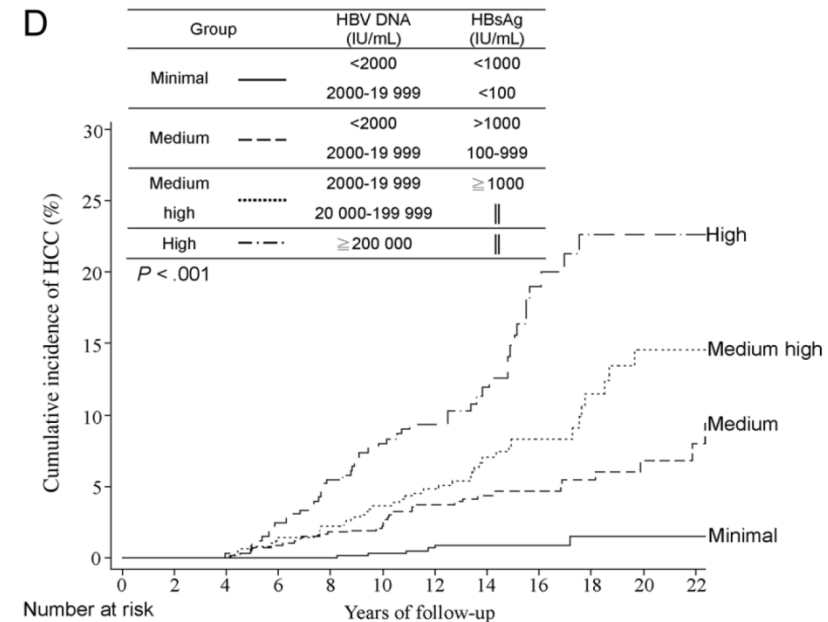
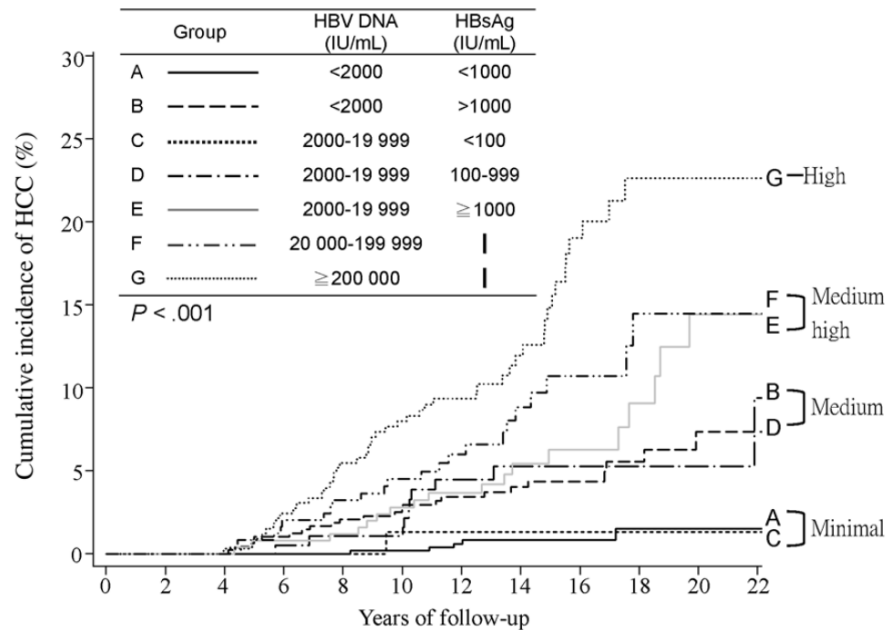
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# Risikostratifizierung des HCC bei HBeAg-negativer HBV Infektion durch Kombination viraler Biomarker

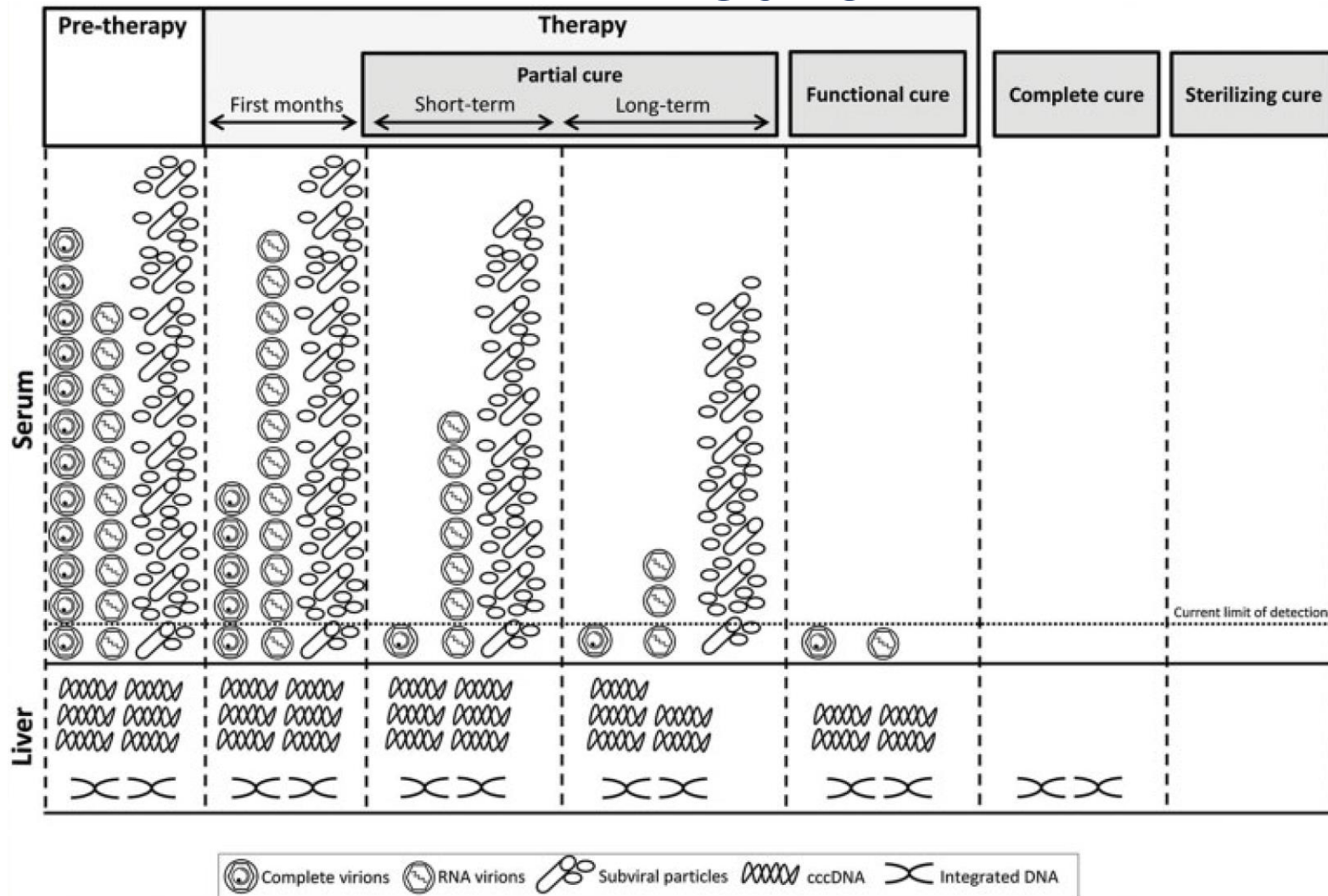
Tseng TC et al. *JID* 2013;208; 584-593



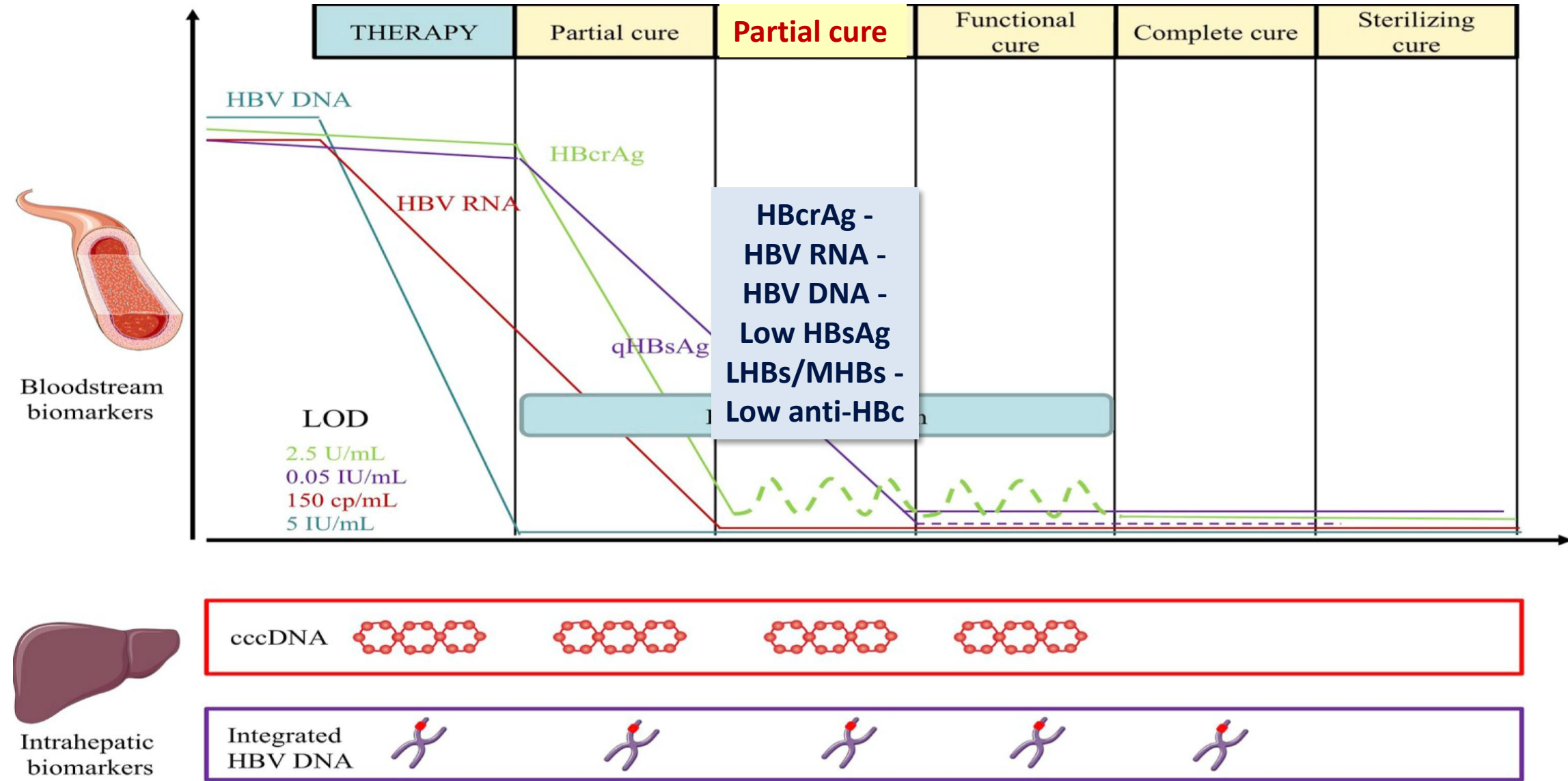
**Does HBsAg positive „cure“ of the disease and diseases associated outcomes exist? How to best define HBsAg positive cure of active infection**



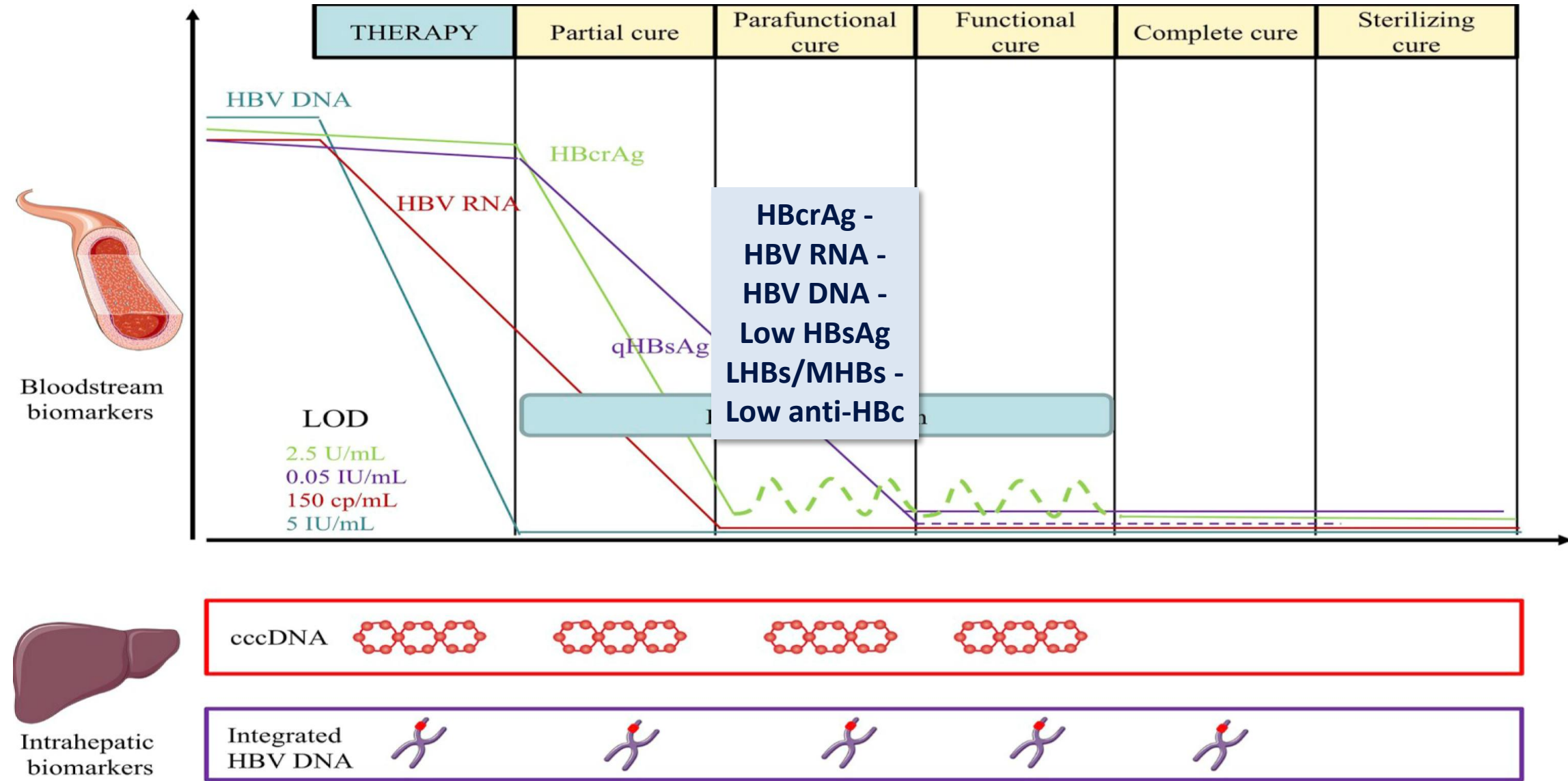
# HBV particles detected in serum and liver of HBV-infected patients at the various endpoints of HBV cure



# Chance for transition into a true functional or even complete cure stage after having achieved a partial cure?



# Chance for transition into a true functional or even complete cure stage after having achieved a partial cure?



**HBsAg Pre-S truncation**

**ATTACHMENT**

**Uncoating**

**Nuclear Transport**

**GOLGI**

**ER**

**HBsAg proteins:**

- LHBsAg
- MHBsAg
- SHBsAg

**HBcAg**

**p22crAg**

**HBeAg**

**Immature Nucleocapsid**

**Reverse Transcription**

**RC-DNA**

**Mature Nucleocapsid**

**DSL-DNA**

**HBV Integration**

**Host Genome**

**Pre-S1 mRNA**

**Pre-S2/S mRNA**

**Precore mRNA**

**Pregenomic RNA**

**HBx mRNA**

**cccDNA**

**SMC5/6**

**Intracellular Conversion Pathway**

**Major DNA Integration Pathway**

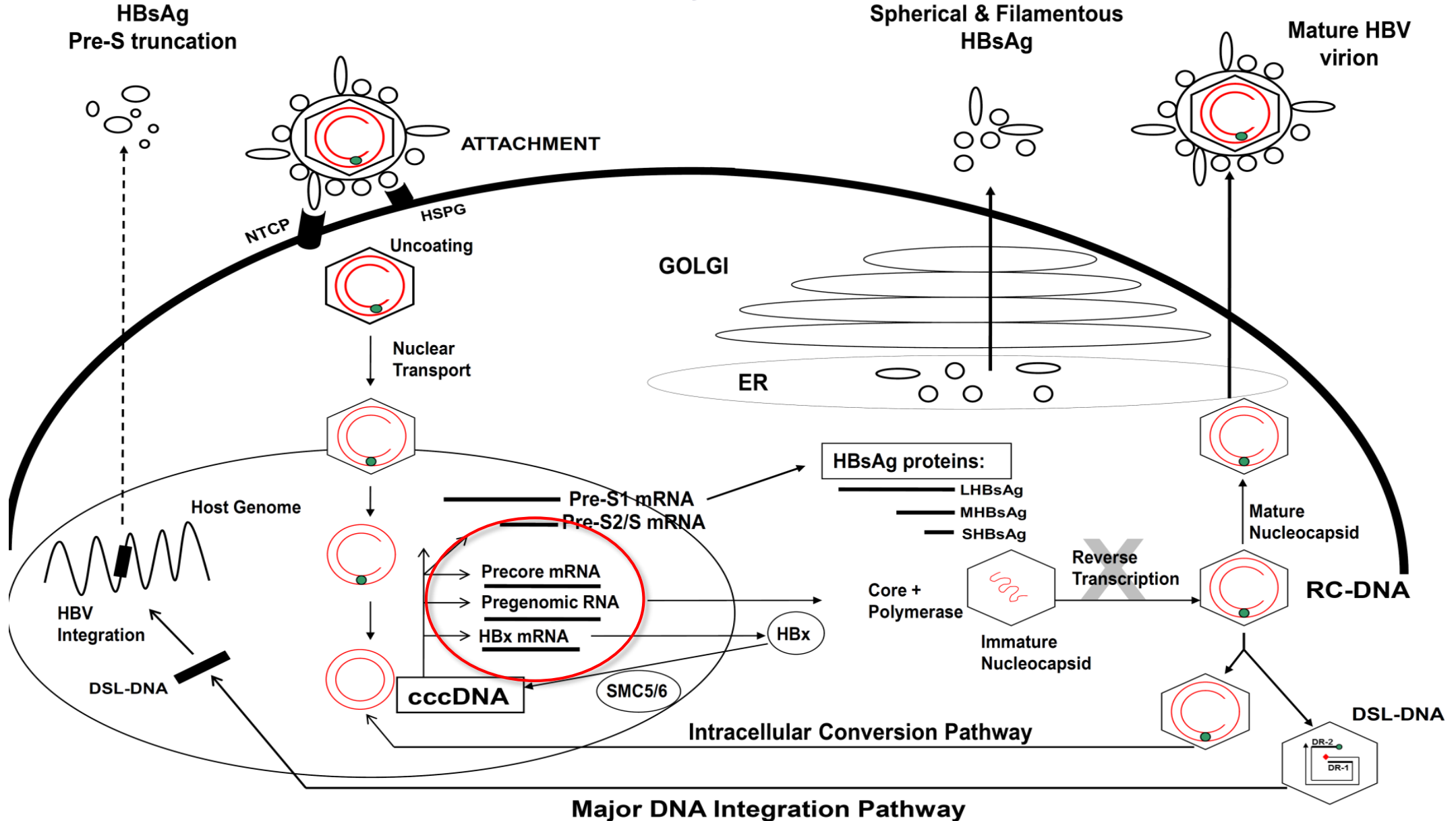
**Defective particle without genome**

**Enveloped pgRNA virion**

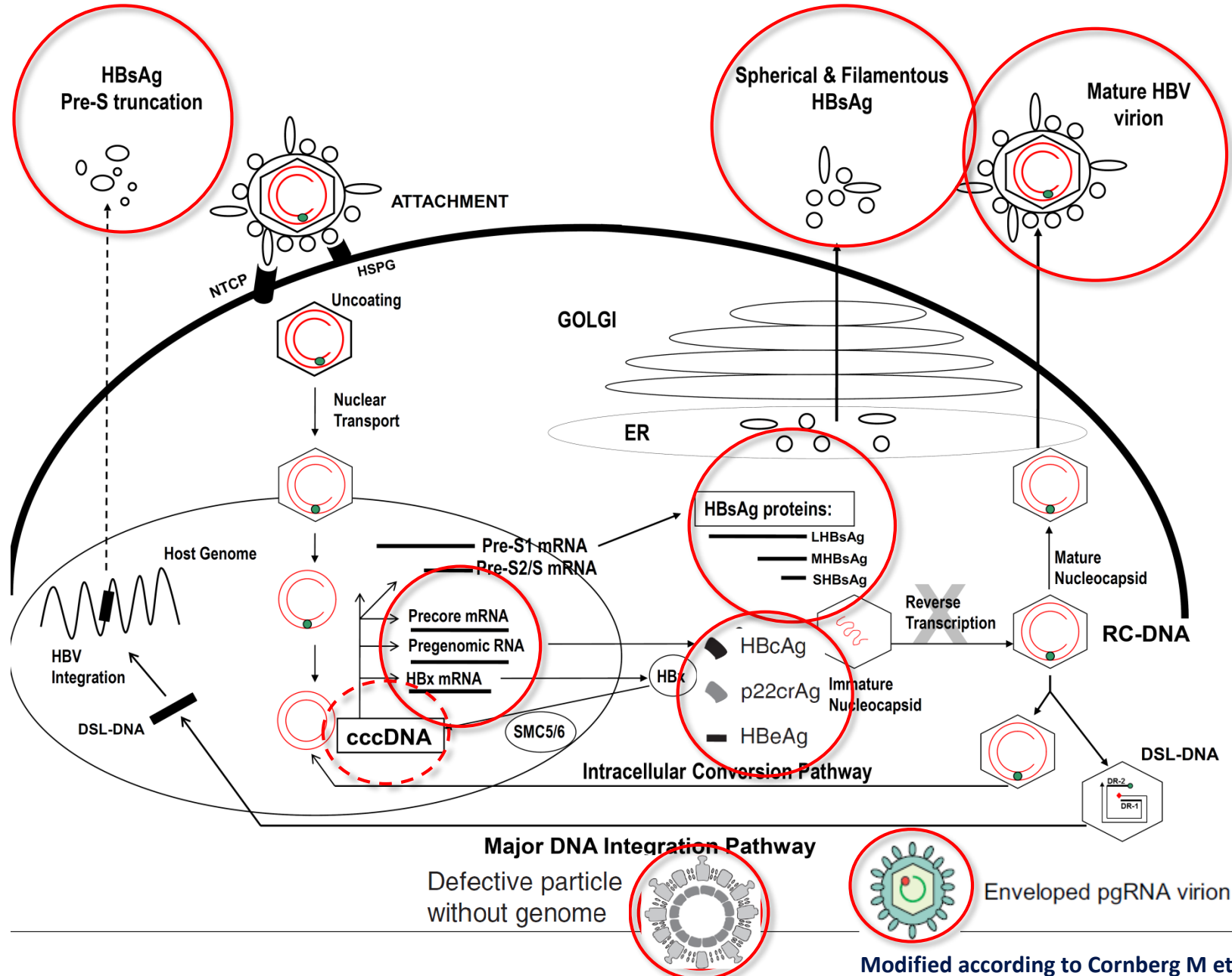
**Modified according to Cornberg M et al.**



# HBV life cycle – HBV RNA



# Potential biomarkers for assessing endpoints



# Clinical utility of quantitative anti-HBc levels

- Anti-HBc levels are higher in HBsAg-positive patients as compared to those being anti-HBs-positive (but anti-HBc affinity was lower in chronic infection than during recovery)<sup>1</sup>
- Low anti-HBc levels are associated with HBsAg seroclearance (cut-off < 3log)<sup>2</sup>
- Serum levels of anti-HBc correlate with cccDNA positivity in OBI<sup>3</sup>
- Anti-HBc levels (cut-off  $\geq 6.41$  IU/ml) were significantly associated with high risk of HBV reactivation during immunosuppressive therapy<sup>4</sup>

<sup>1</sup>Han et al. *J Clin Virol.* 2011;52(4):295-9

<sup>2</sup>Hu et al. *Clin Gastroenterol Hepatol.* 2018 [Epub ahead of print]

<sup>3</sup>Caviglia et al. *J Hepatol.* 2018;69(2):301-307

<sup>4</sup>Yang et al. *J Hepatol.* 2018 Aug;69(2):286-292.

# Why aiming for HBsAg loss

- The loss of HBsAg is regarded as the optimal treatment endpoint, termed 'functional cure', but it is only rarely achieved with our current antiviral armamentarium.
- Spontaneous HBsAg seroreversion with reactivation of the inflammatory liver process after HBsAg loss is rare and may occur in patients with a significant impairment of their immune function
- The main advantage of HBsAg loss is that it allows a safe discontinuation of antiviral therapy.



# Serum level of antibodies against hepatitis B core protein is associated with clinical relapse after discontinuation of nucleos(t)ide analogue therapy

- Clinical relapse occurred in 39 patients (in 46% of patients at year 4 after discontinuation). High level of anti-HBc at the end of treatment (hazard ratio [HR], 0.31 per log IU/mL;  $P = .002$ ) and low level of HB [surface antigen](#) (HBsAg) at the end of treatment (HR, 1.71 per log IU/mL;  $P = .032$ ) were associated with a reduced risk of clinical relapse after adjusting for age, start of nucleos(t)ide analogue therapy, HBeAg-status, and consolidation therapy duration. At year 4, 21% of patients with anti-HBc levels at the end of treatment  $\geq 1000$  IU/mL developed a clinical relapse compared to 85% of patients with levels  $< 100$  IU/mL ( $P < .001$ ).