

RHU « CirB-RNA »
ANR-17-RHUS-0003

International Workshop on Viral Biomarkers

September 7, 2023
Institut Lumière, Lyon



The molecular biology of HBV RNAs: insights on new biomarkers

Massimo Levrero (Lyon, France)

RHU « CirB-RNA »
ANR-17-RHUS-0003

International Workshop on Viral Biomarkers

September 7, 2023
Institut Lumière, Lyon



The molecular biology of HBV RNAs: insights on new biomarkers

Massimo Levrero (Lyon, France)

the CirB-RNA WP2 members



RHU « CirB-RNA »
ANR-17-RHUS-0003

International Workshop on Viral Biomarkers

September 7, 2023
Institut Lumière, Lyon



The molecular biology of HBV RNAs: insights on new biomarkers

Massimo Levrero (Lyon, France)

the CirB-RNA WP2 and WP3 Roche members



Marynthia Heil



Christian
Voitenleitner



Aaron
Hamilton



Debra Liggett



Beth Scott



Thomas
Meister



Paul Dawson

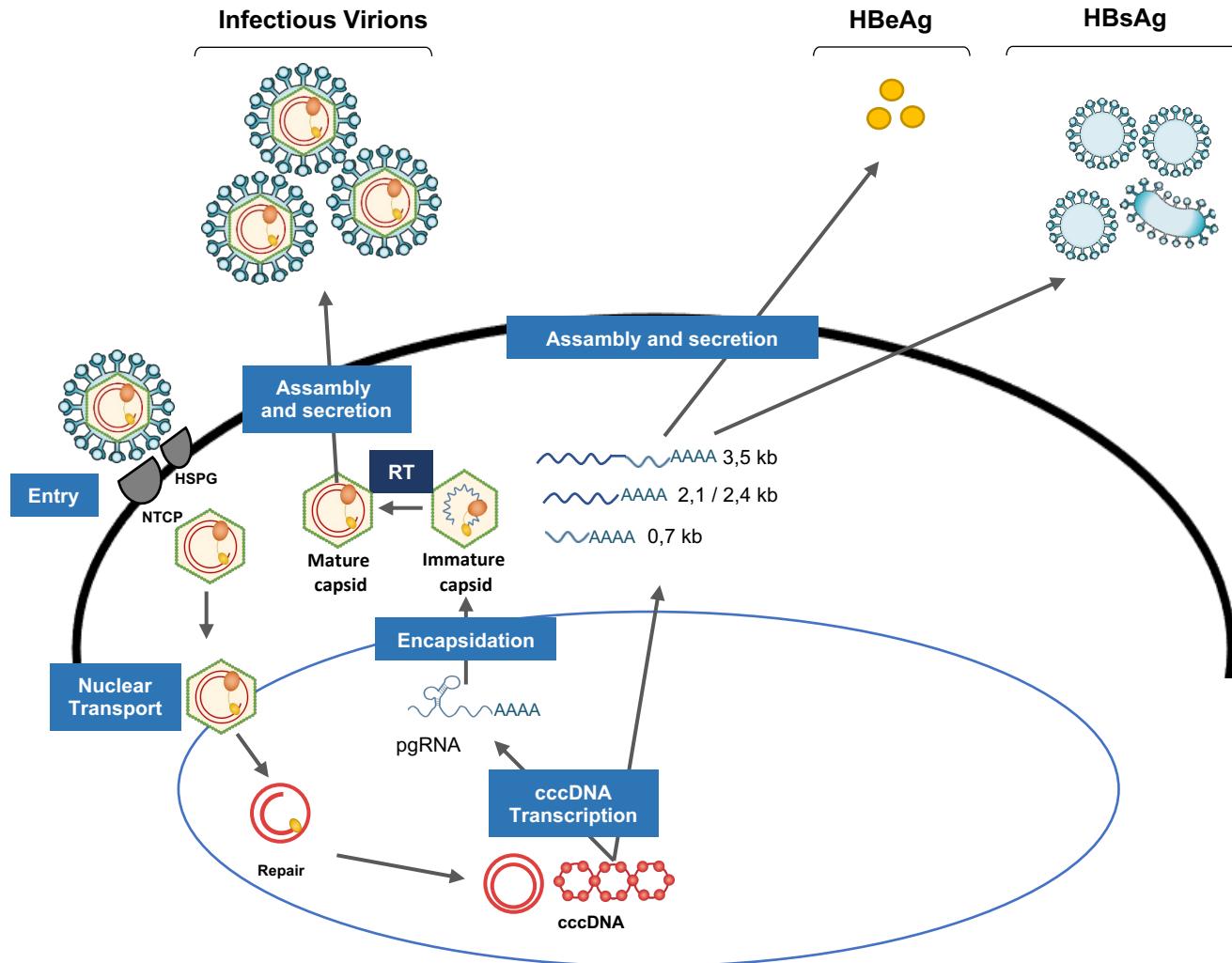


Michael Lobritz



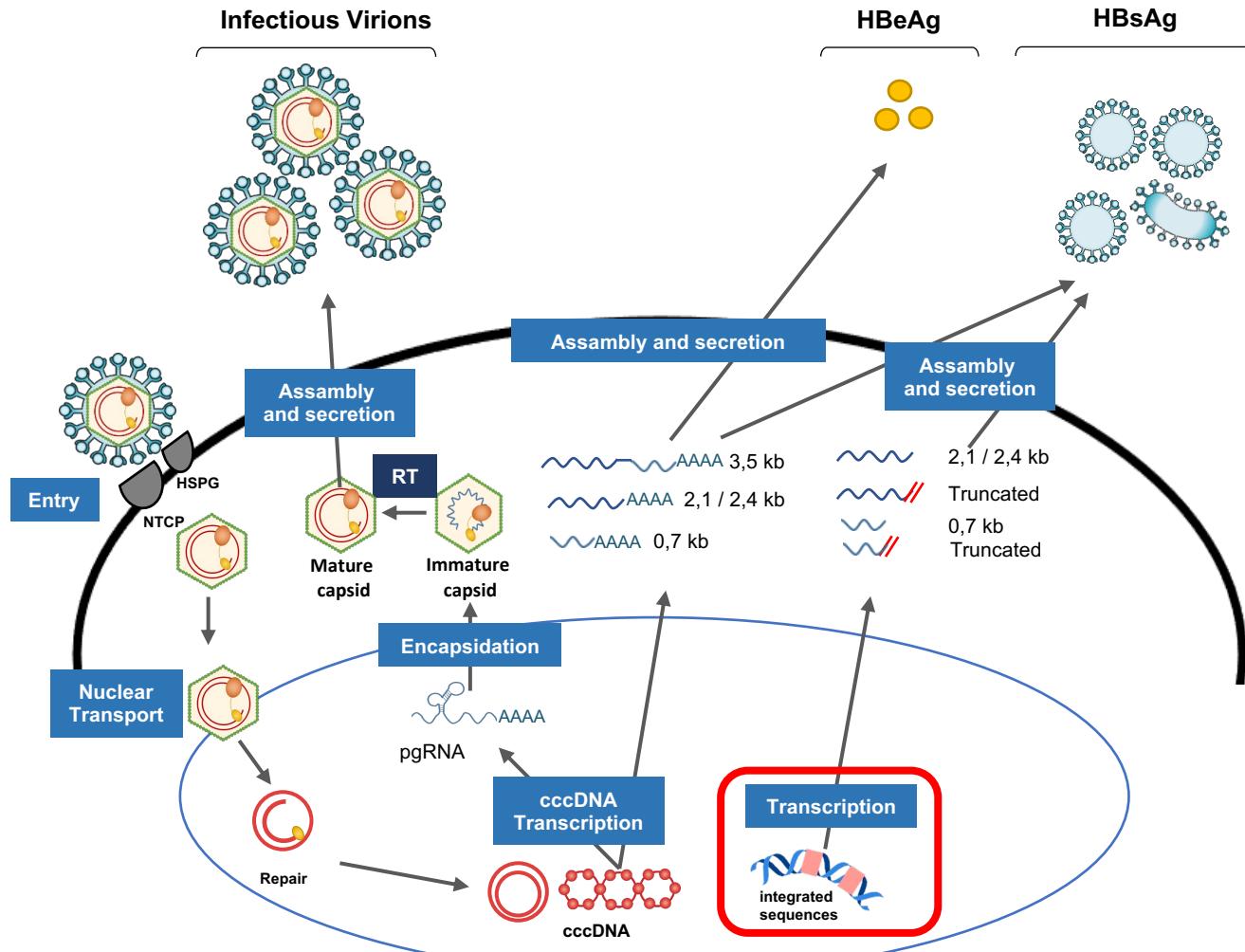
Alan Blair

The molecular biology of HBV RNAs .1



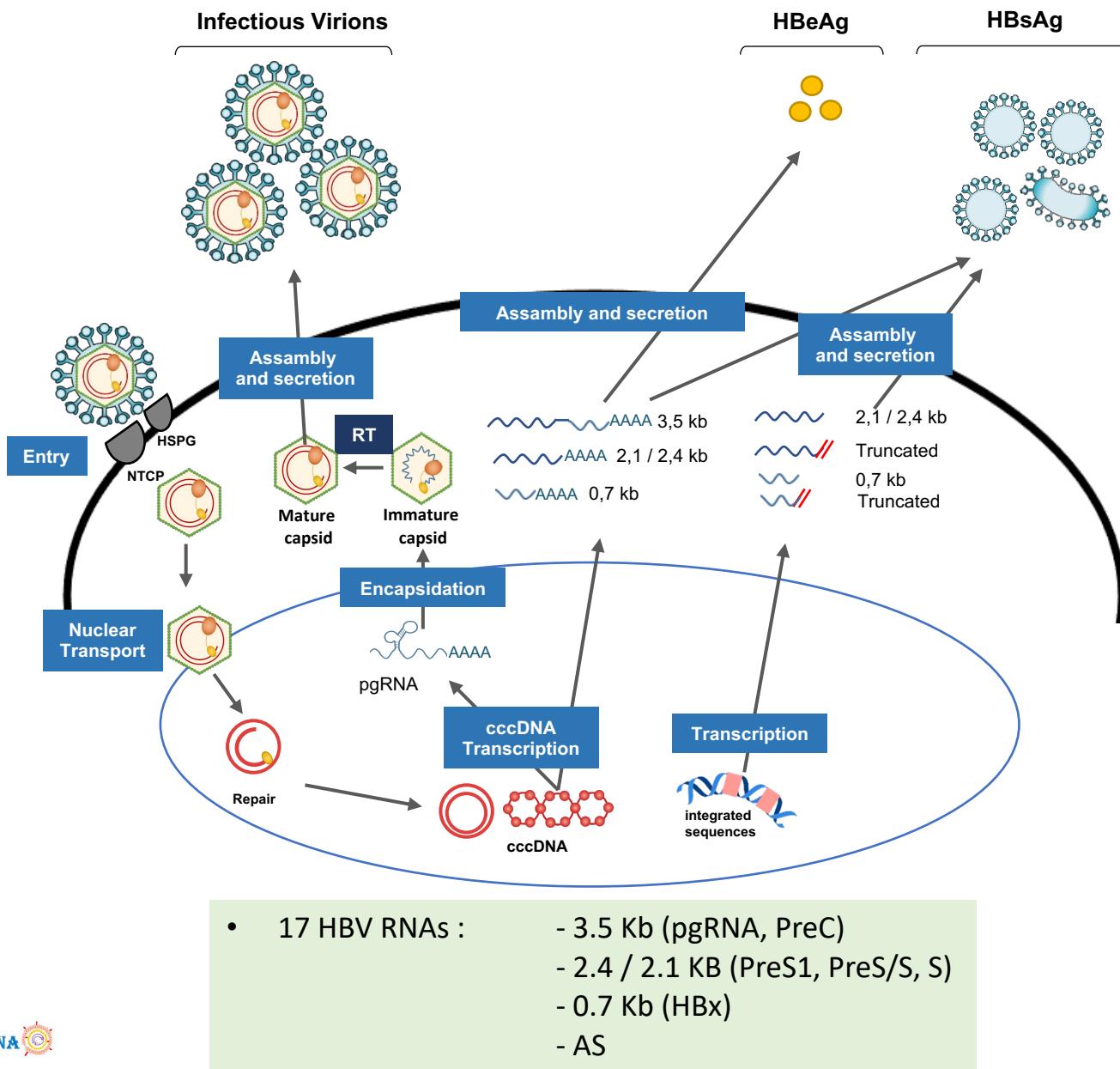
- 5 HBV RNAs :
 - 3.5 Kb (pgRNA, PreC)
 - 2.4 / 2.1 KB (PreS1, PreS/S)
 - 0.7 Kb (HBx)

The molecular biology of HBV RNAs .1

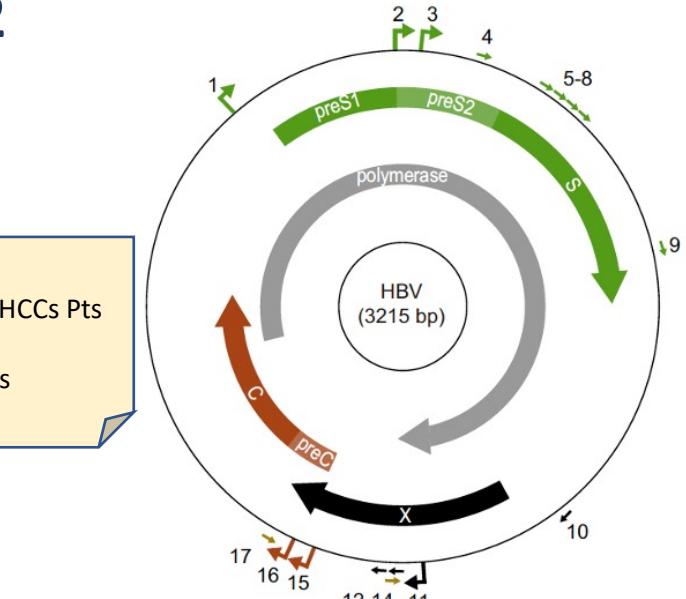


- 5 HBV RNAs :
 - 3.5 Kb (pgRNA, PreC)
 - 2.4 / 2.1 KB (PreS1, PreS/S)
 - 0.7 Kb (HBx)

The molecular biology of HBV RNAs .2



Liver:
- 16 T/NT HCCs Pts
Blood
- 8 CHB Pts



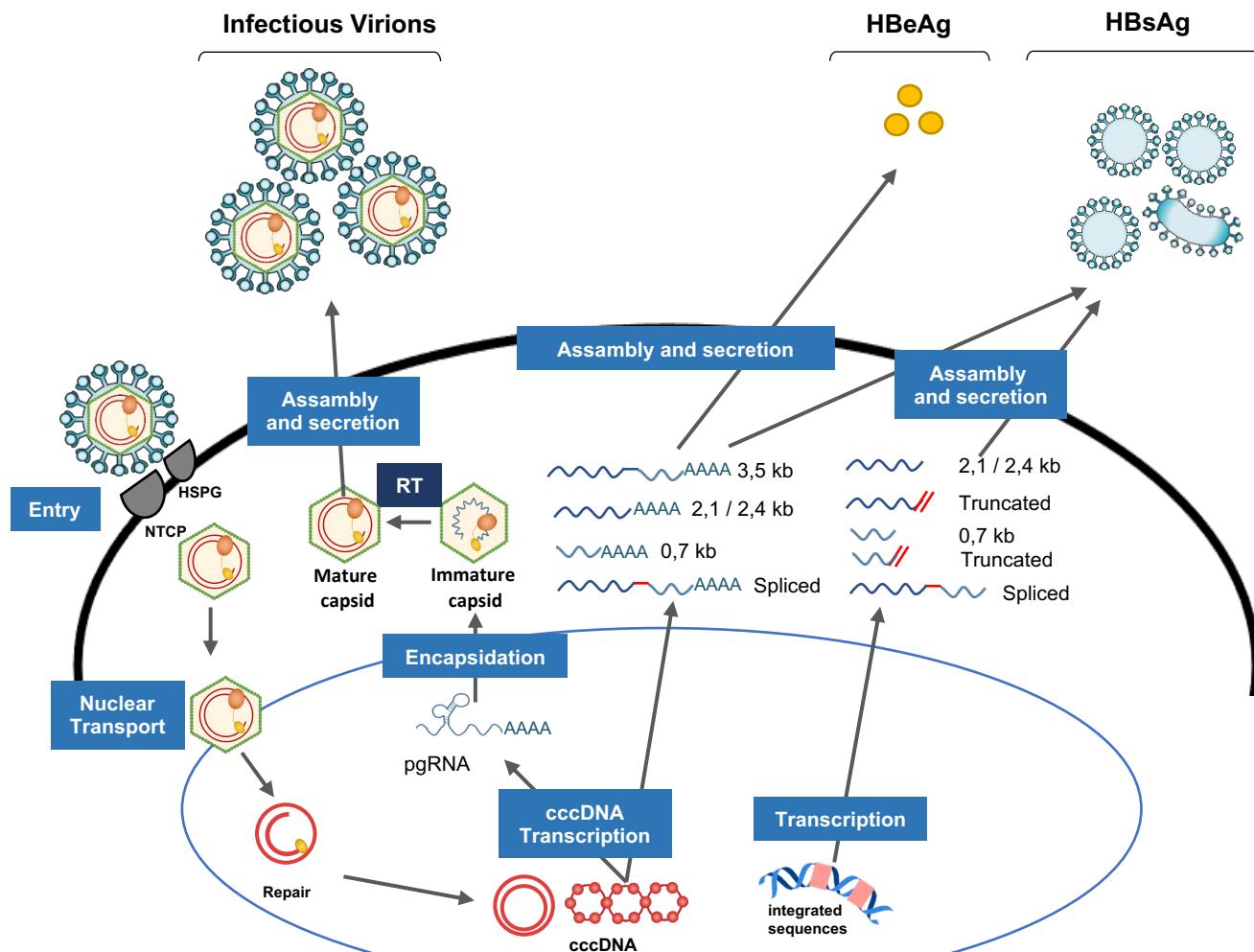
Major peaks (>1,000 raw tags)

#	Gene	Start	End	Total count	Strand	Model
1	PreS1	2802	2807	12,748	+	Y
2	PreS/S	3183	19	201,883	+	Y
3	S	25	121	19,004	+	Y
11	X	1523	1524	1,013	+	Y
15	PreC/C	1745	1746	1,106	+	N
16	pgRNA	1816	1819	112,452	+	Y

Minor peaks (100 ~ 1,000 raw tags)

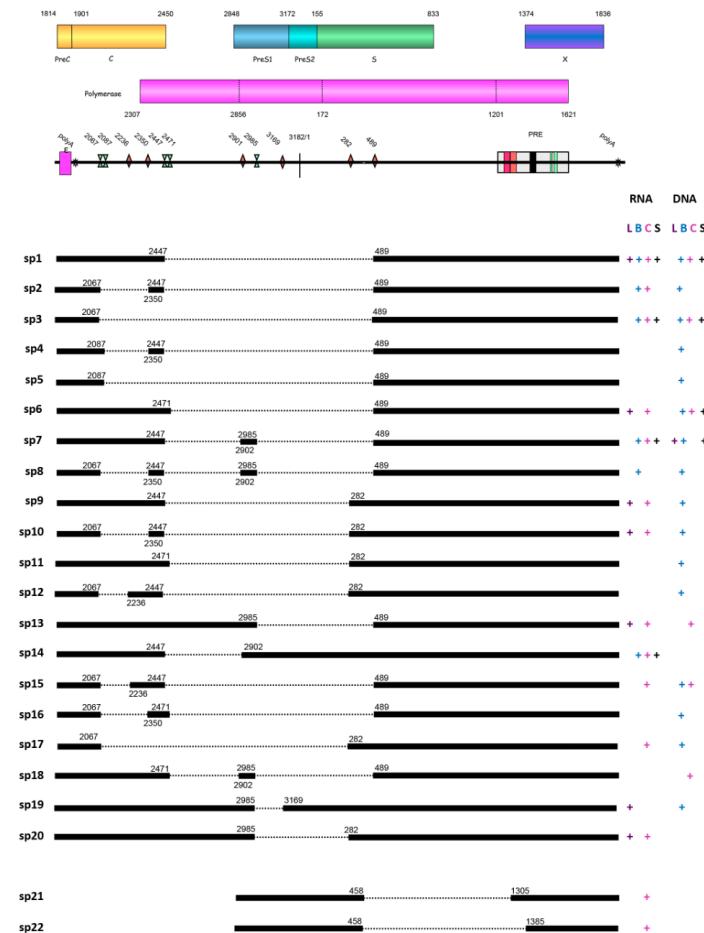
#	Gene	Start	End	Total count	Strand	Model
4	S	132	135	248	+	N
5	S	203	207	527	+	Y
6	S	258	262	282	+	N
7	S	284	285	528	+	Y
8	S	302	303	118	+	N
9	S	759	760	729	+	Y
10	X	1210	1211	124	+	Y
12	AS	1557	1610	157	-	Y
13	X	1572	1573	117	+	N
14	X	1655	1656	316	+	N
17	AS	1822	1834	207	-	N

The molecular biology of HBV RNAs .3



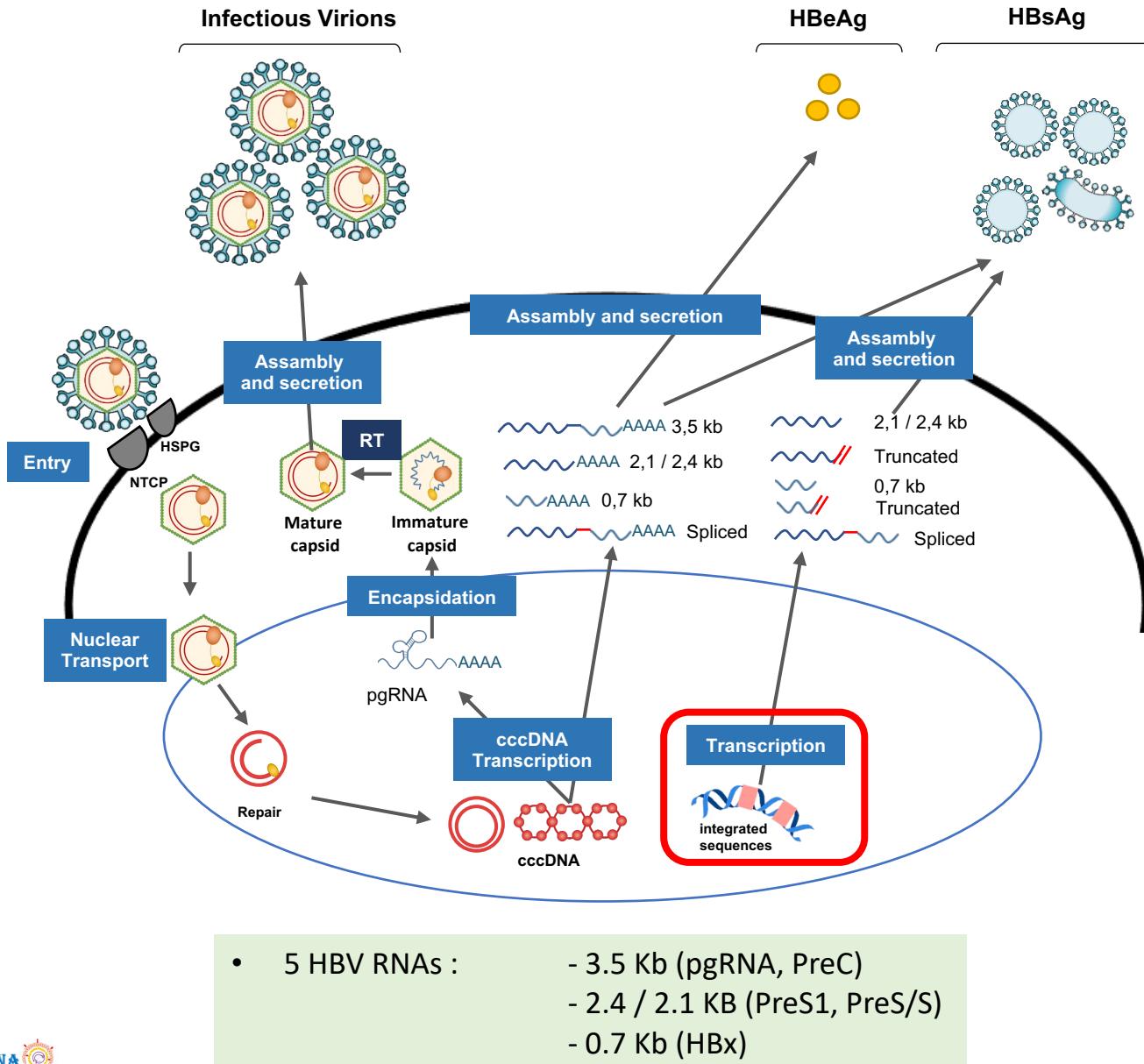
- 5 HBV RNAs :
 - 3.5 Kb (pgRNA, PreC)
 - 2.4 / 2.1 KB (PreS1, PreS/S)
 - 0.7 Kb (HBx)
- 22 spliced species

spliced RNAs



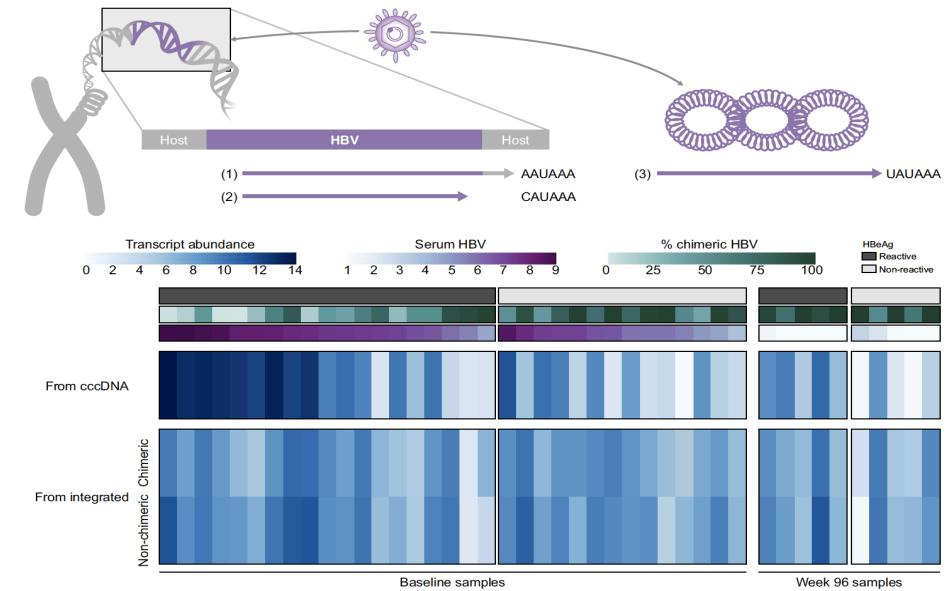
Kremsdorff et al. 2021

The molecular biology of HBV RNAs .4



Iso-Seq PacBio RNA sequencing

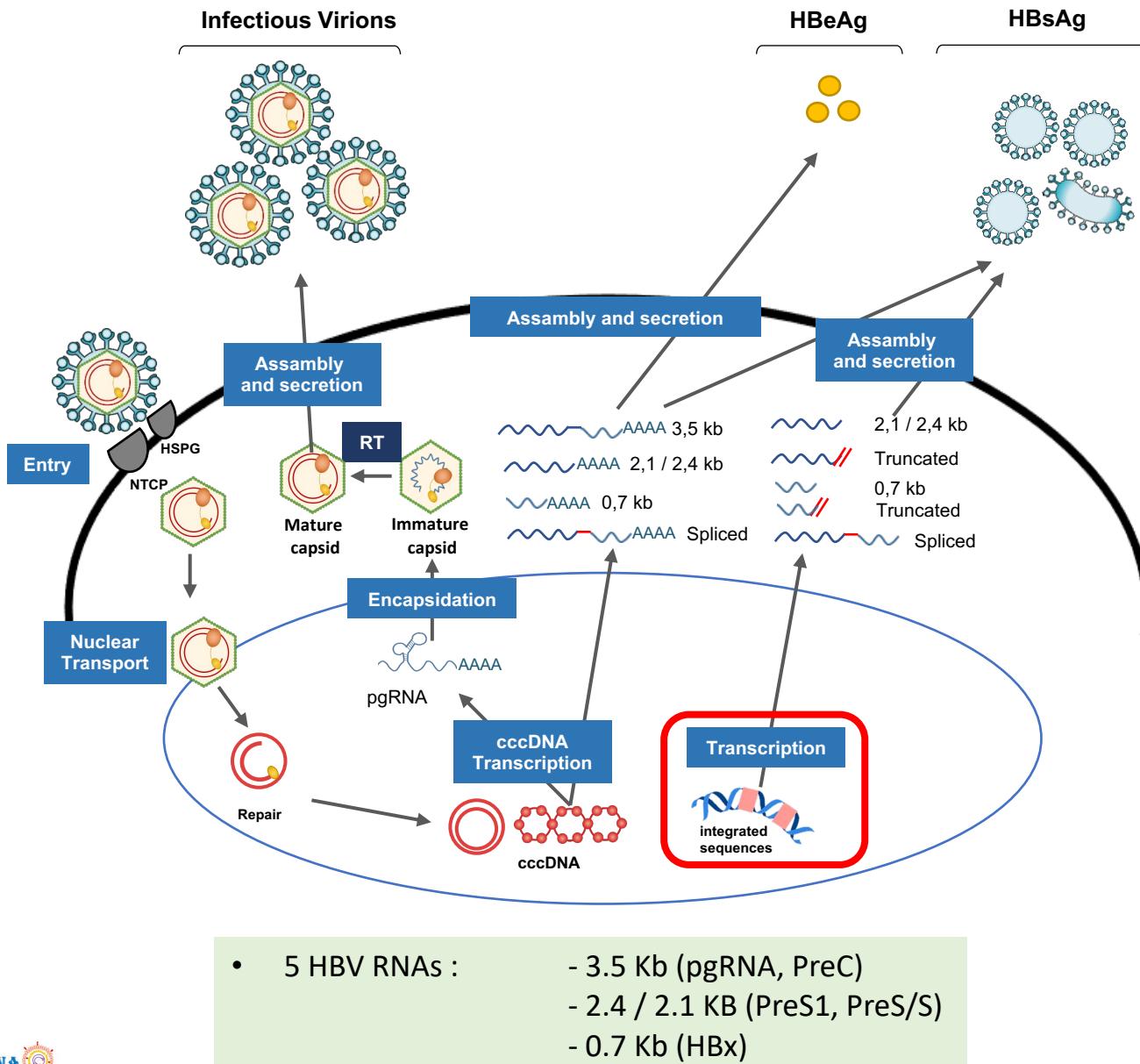
42 CHB patients from Gilead's study GS-US-174-0149



- The amount of transcription from integrations similar between HBeAg(+) and HBeAg(-) pts
- Lower transcription from cccDNA in HBeAg(-) pts
- Ratio of integrated HBV transcripts to cccDNA transcripts much higher in HBeAg(-) pts

Van Buuren, JHEP Reports 2022

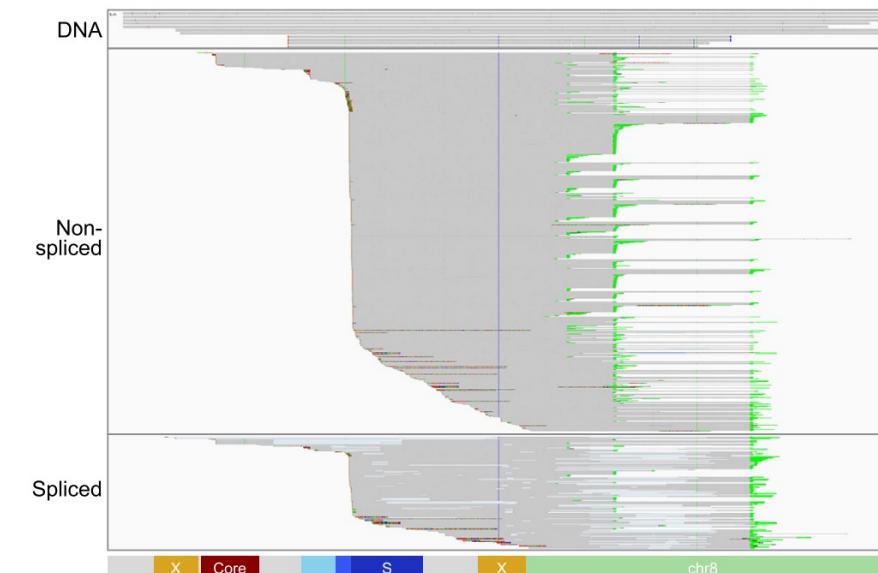
The molecular biology of HBV RNAs .4



Iso-Seq PacBio RNA sequencing

42 CHB patients from Gilead's study GS-US-174-0149

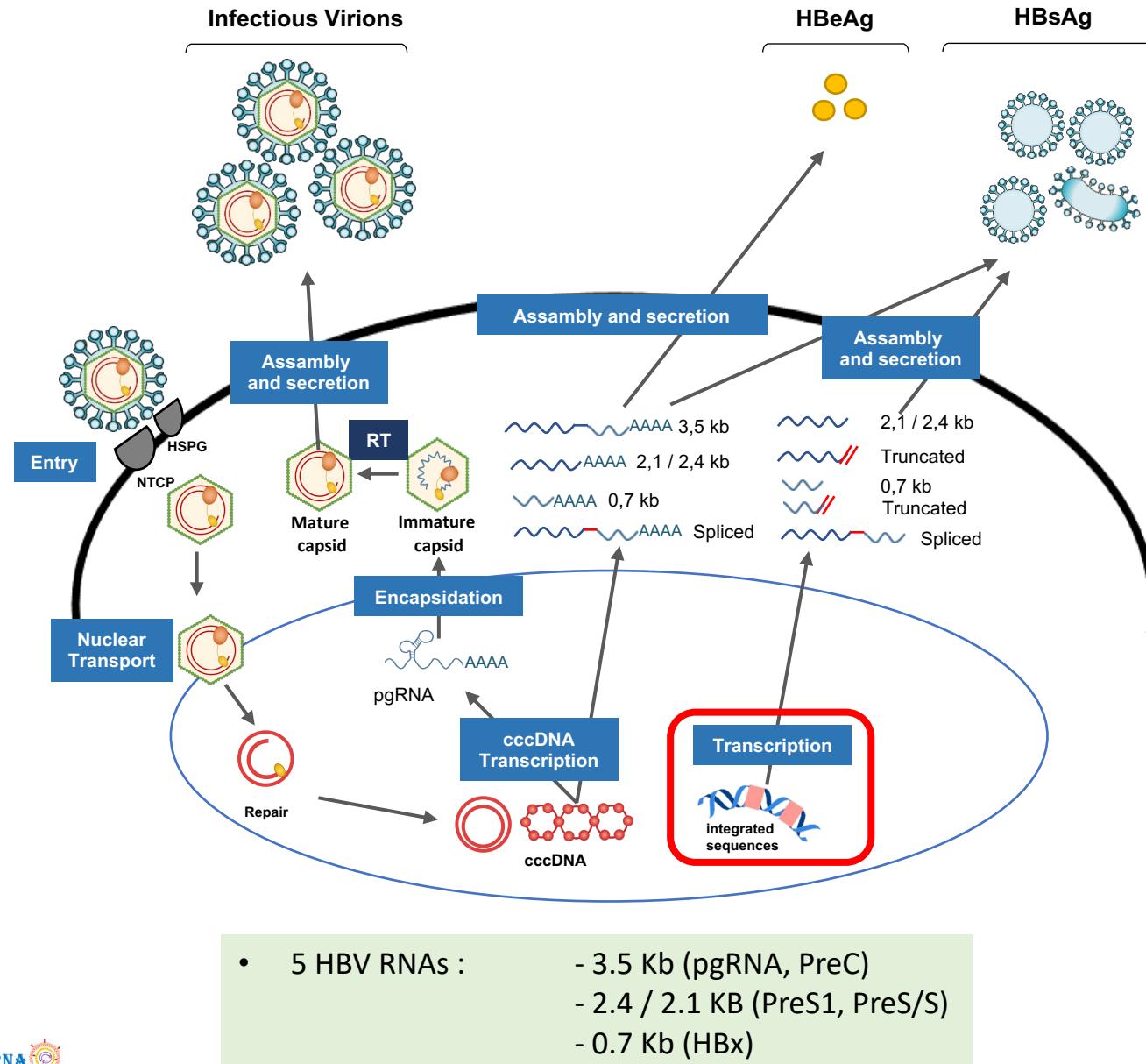
- an unusually long clonally expanded HBV sequence (4,119 bp), chimeric with chromosome 8 at the 3' end generates transcripts expressing all of Core, PreS1, and PreS2/S and utilises 3 different host poly(A) sequences.



- ratio of spliced to non-spliced HBV RNAs ranging from 2% to over 20%
- highest in pHBsAg-positive patients
- the majority of spliced HBV RNAs originated from cccDNA,

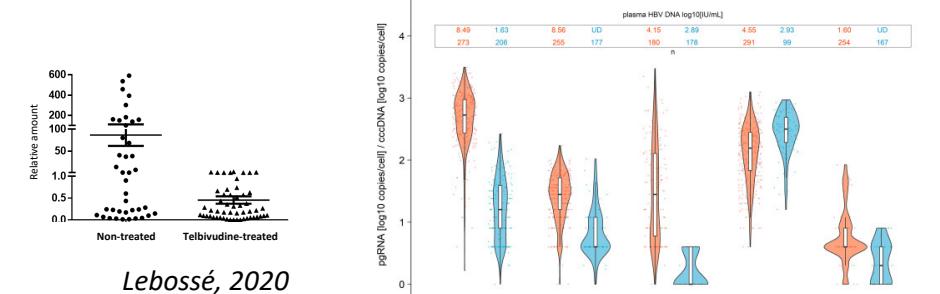
Van Buuren, JHEP Reports 2022

The molecular biology of HBV RNAs .5

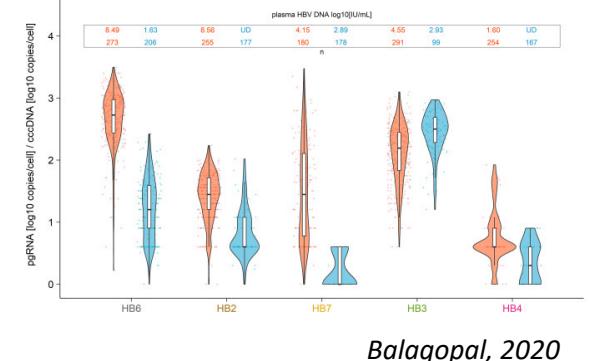


- NUCs decrease transcription from the cccDNA over time ...

3.5Kb RNA quantification

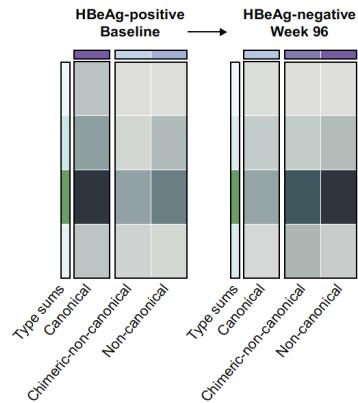


Paired biopsies during ttX

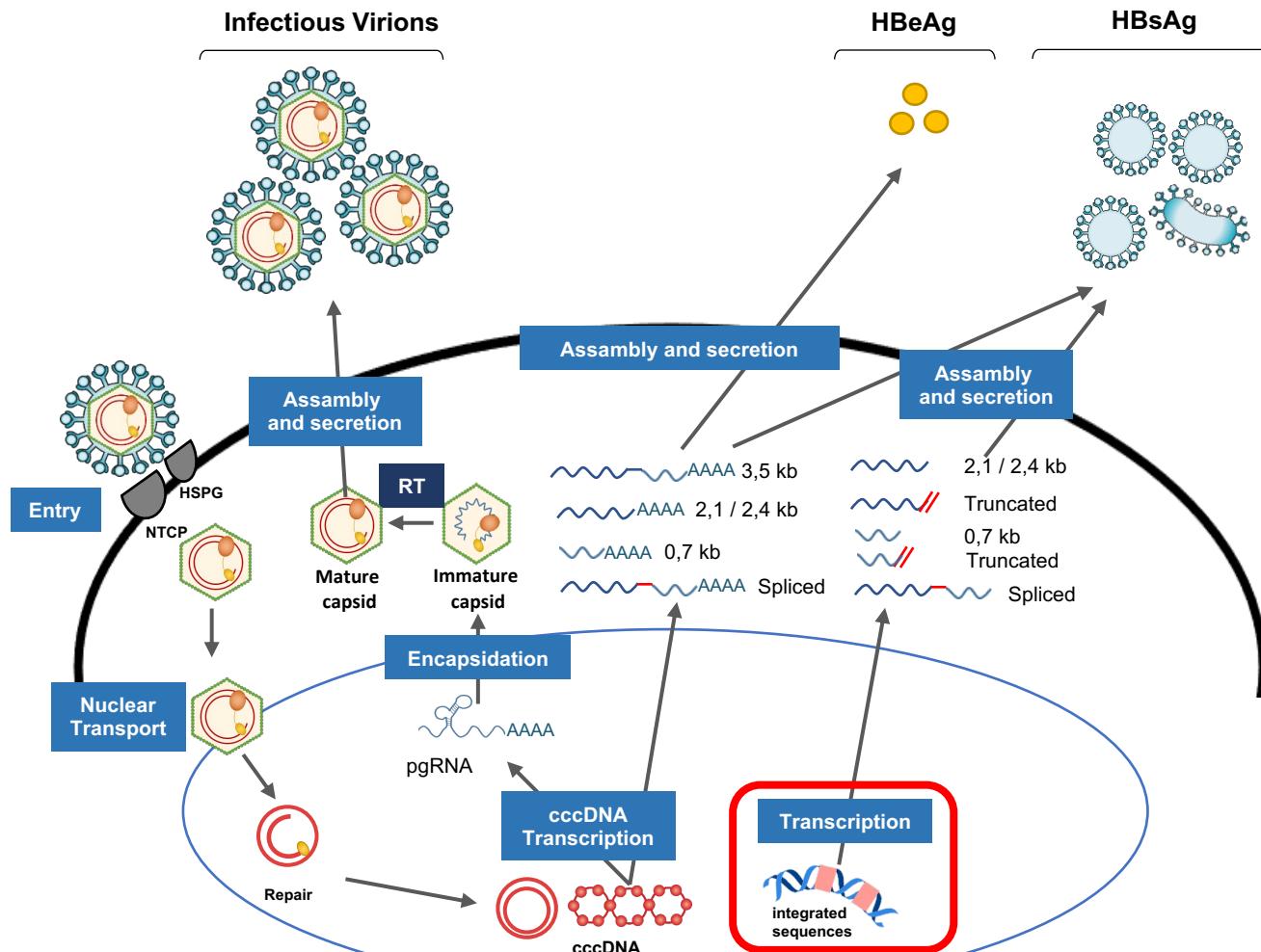


.... but do not affect transcription from HBV integrants

Iso-Seq PacBio RNA sequencing

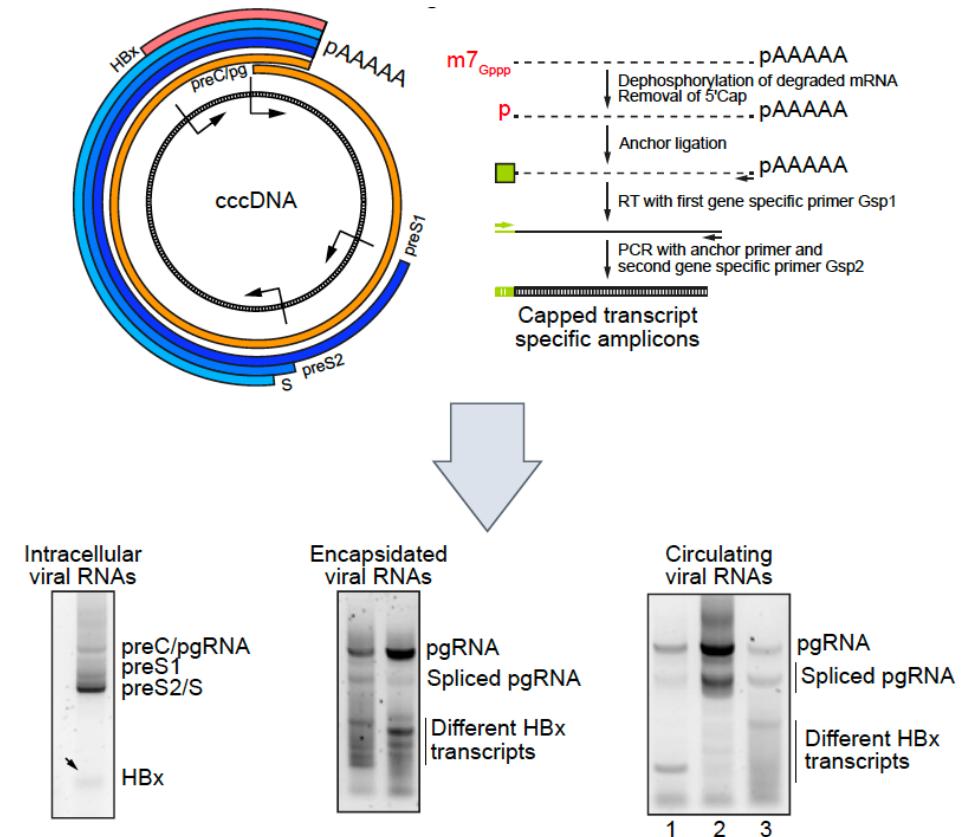


The molecular biology of HBV RNAs .6



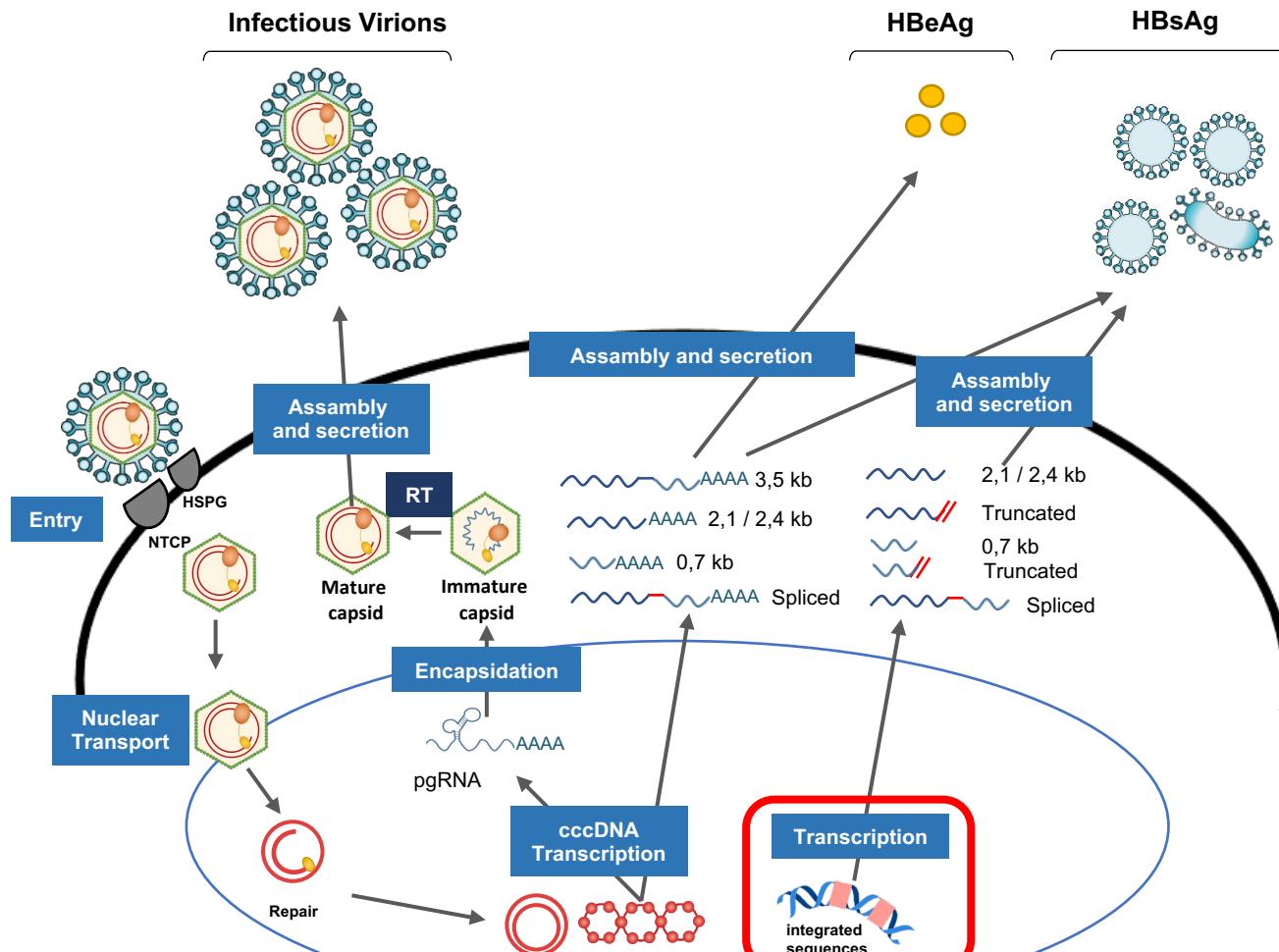
- 5 HBV RNAs :
 - 3.5 Kb (pgRNA, PreC)
 - 2.4 / 2.1 KB (PreS1, PreS/S)
 - 0.7 Kb (HBx)

- Full-length 5'RACE to identify HBV transcripts in HBV-infected hepatocytes ... and patient serum



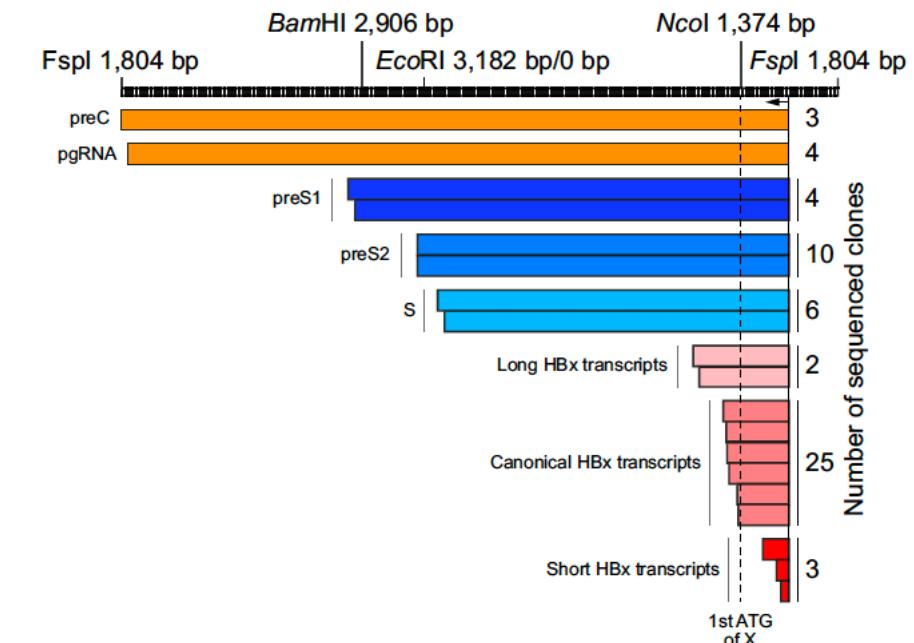
Stadelmayer B et Al, J Hepatol 2020

The molecular biology of HBV RNAs .6



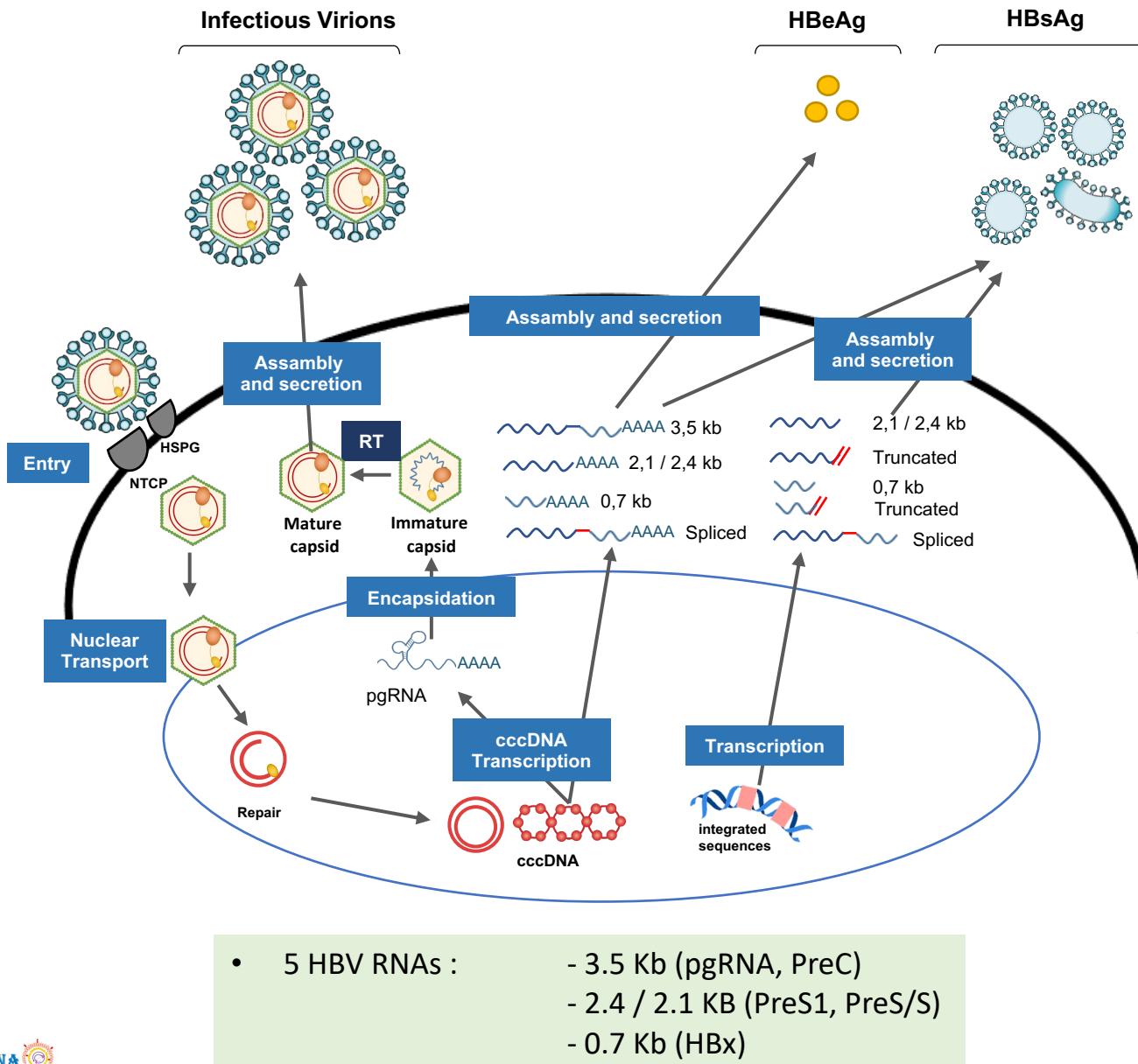
- 5 HBV RNAs :
 - 3.5 Kb (pgRNA, PreC)
 - 2.4 / 2.1 KB (PreS1, PreS/S)
 - 0.7 Kb (HBx)

- Full-length 5'RACE to identify HBV transcripts in HBV-infected hepatocytes ... and patient serum



Stadelmayer B et Al, J Hepatol 2020

The molecular biology of HBV RNAs .7



pgRNA variants

- 3' truncated transcript resulting from a cryptic polyadenylation signal (cryptic tRNA), thought to arise from integrated HBV genomes

Hilger, 1991

van Bömmel et al., 2015

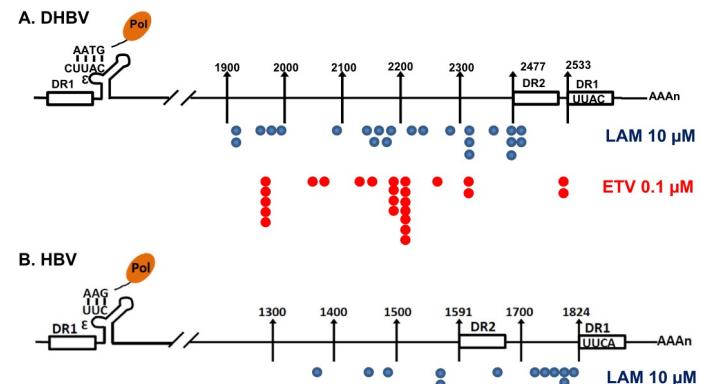
- pgRNA spliced variants

Bayliss, 2013

Betz-Stablein, 2016

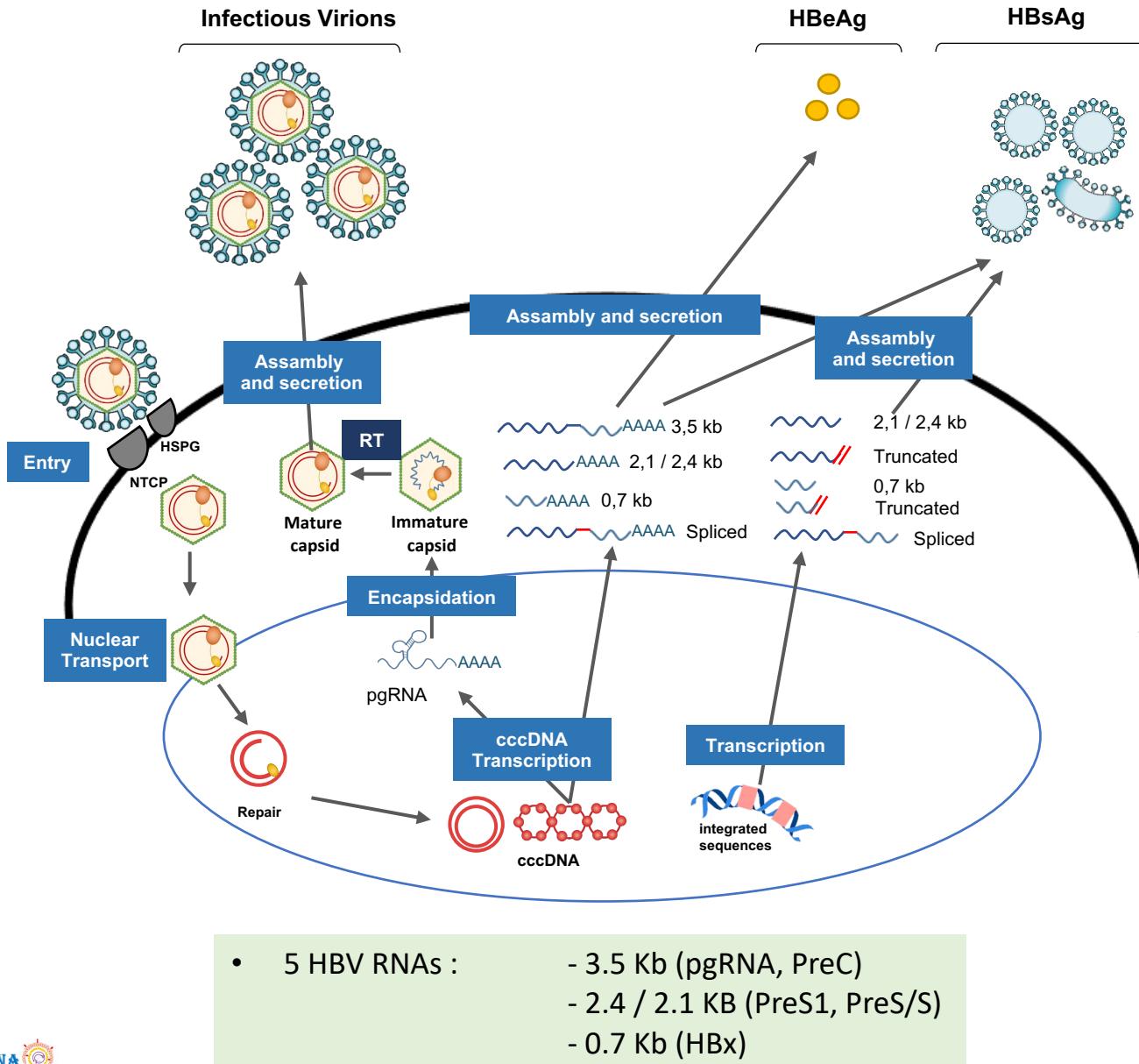
Lim, 2021

- non-polyadenylated 3' truncated variant transcripts (incomplete inhibition of the RNase H) during NA therapy



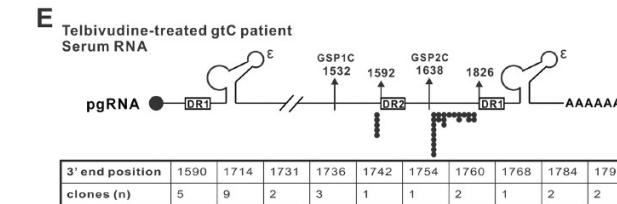
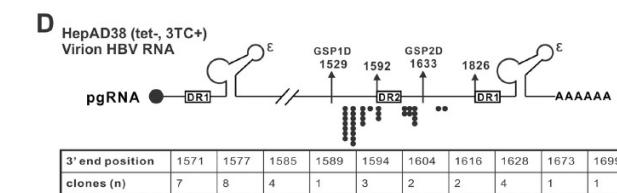
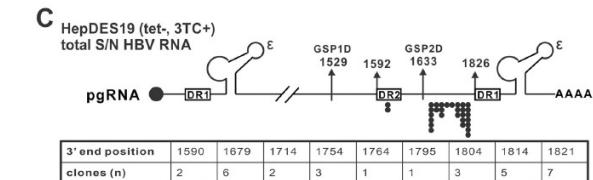
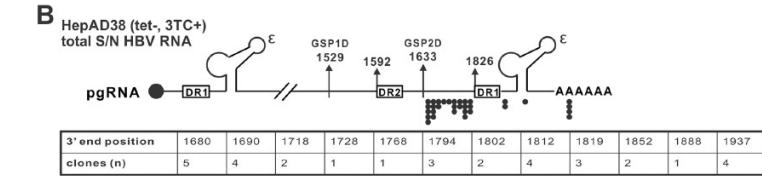
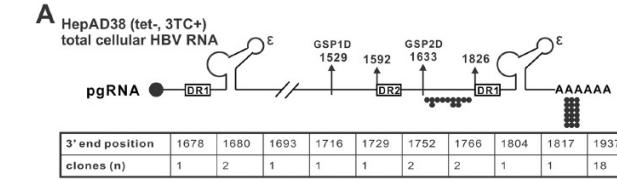
Zhang et al., 2016

The molecular biology of HBV RNAs .7



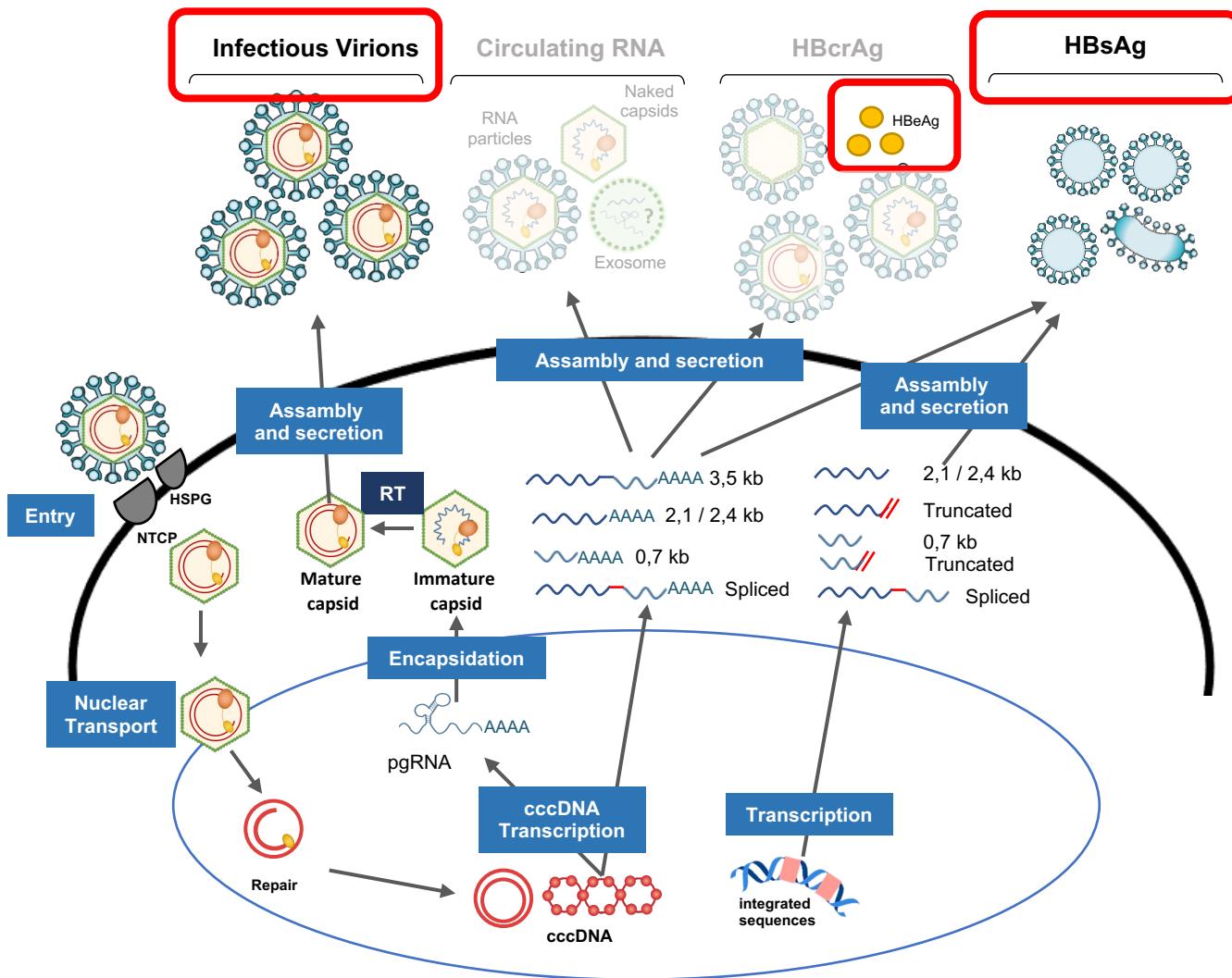
pgRNA variants

- non-polyadenylated 3' truncated variant transcripts (incomplete inhibition of the RNase H) during NA therapy



The molecular biology of HBV RNAs: insights on new biomarkers

What old HBV biomarkers cannot tell us?



rcDNA contained in virions can originate only from cccDNA, but NUC-induced serum HBV DNA suppression does not eliminate cccDNA

→ Not useful in combination therapies !

HBeAg reflects HBV replication and cccDNA transcription, but HBeAg can disappear from the serum because of the selection of HBeAg-defective viruses

→ Not sufficient !

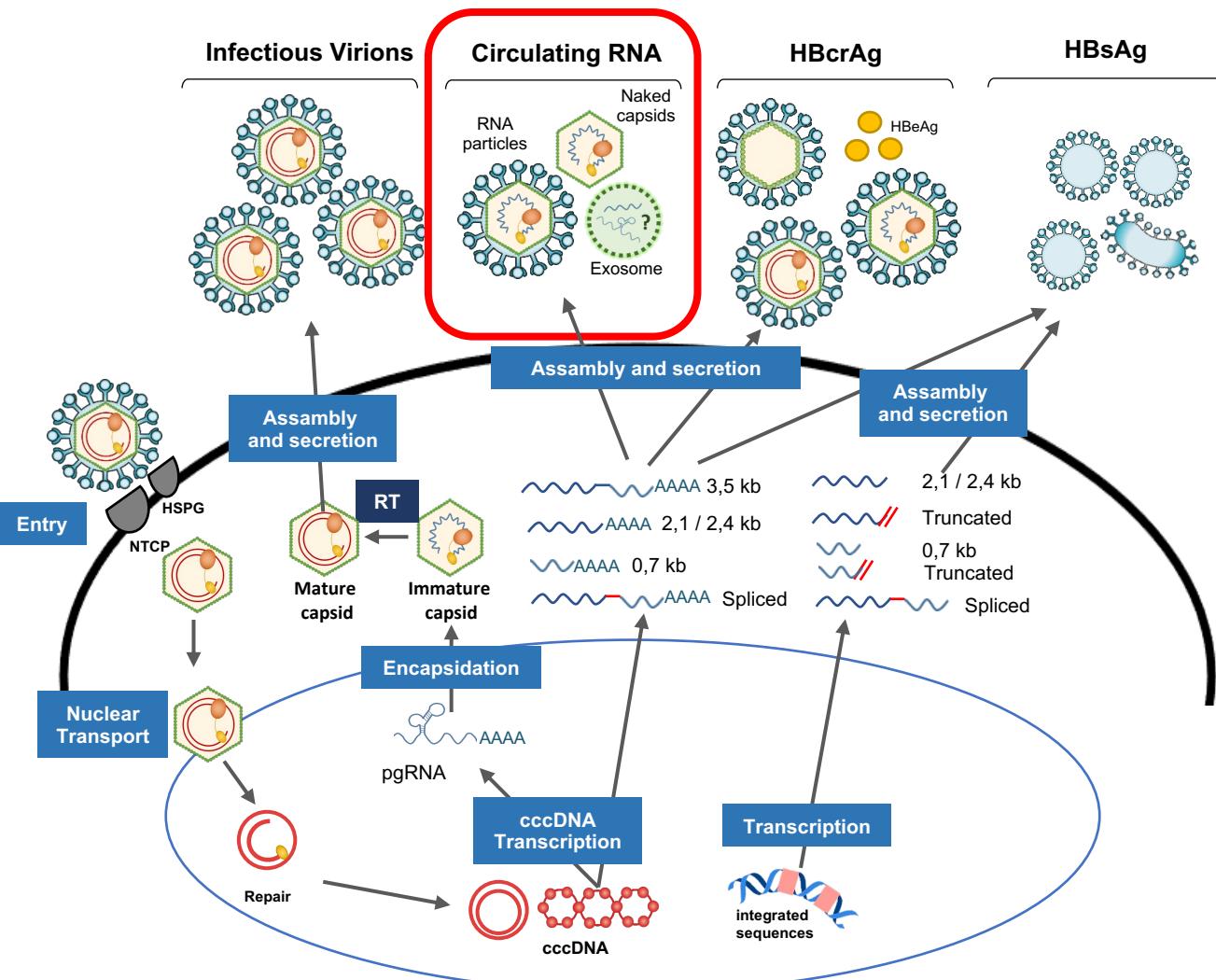
HBsAg can originate from cccDNA AND integrated HBV DNA

→ Not specific to cccDNA

→ HBsAg loss reflects both cccDNA loss and cccDNA inactivation

→ qHBs decline kinetic is too slow to be informative

The molecular biology of HBV RNAs: insights on new biomarkers

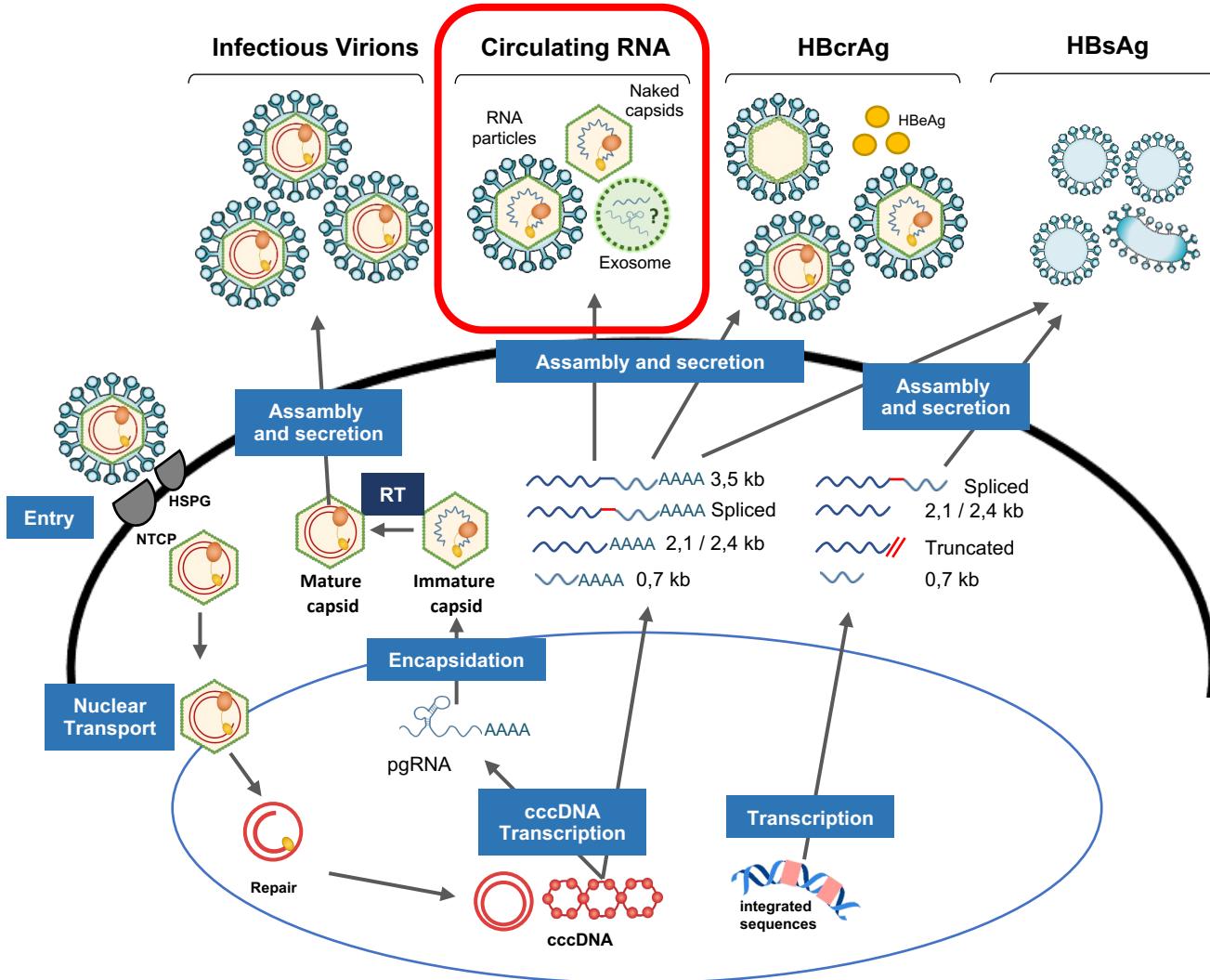


serum HBV-RNAs as a viral biomarker

- clinical question asked
 - reflecting virus / cccDNA activity
 - detecting drugs target engagement
 - predicting clinical endpoints
- biomarker biology
 - which RNA species
 - which vehicle (RNA particles, naked capsids, exosomes, EV) and their impact on detection and significance
- contribution of HBV integrations
 - chimeric transcripts
- detection and quantification strategy
 - assay design and performances

According to the different quantification strategies,
the biological/clinical significance
may be significantly different!

The molecular biology of HBV RNAs: insights on new biomarkers



serum HBV-RNAs as a viral biomarker

- pgRNA is transcribed (only) from cccDNA
- pgRNA well correlates with intrahepatic cccDNA activity in Hu-Hep mouse model

Giersch, 2017

- serum HBV RNA correlates with intrahepatic cccDNA levels/activity

Huang, 2017

Wang, 2021

Testoni, 2023

...but

- serum HBV RNA levels varies in the different CHB phases
- pgRNA is not the only circulating HBV RNA species...

Wang, 2018

Wang J, 2016

Jansen L, 2016

Prakash K, 2018

Lam AM, 2017

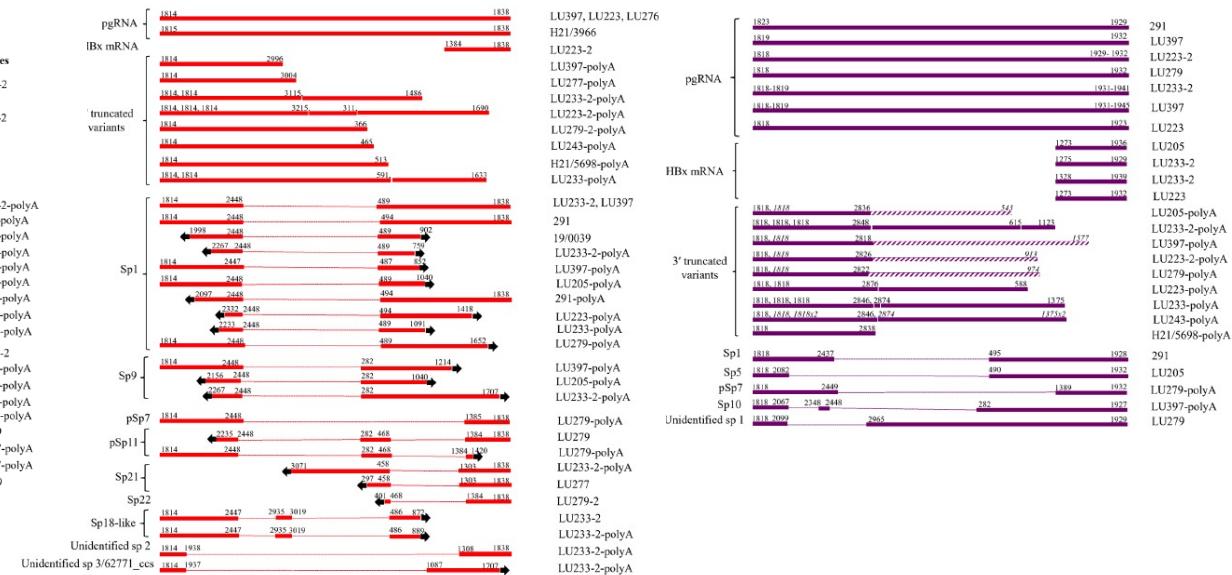
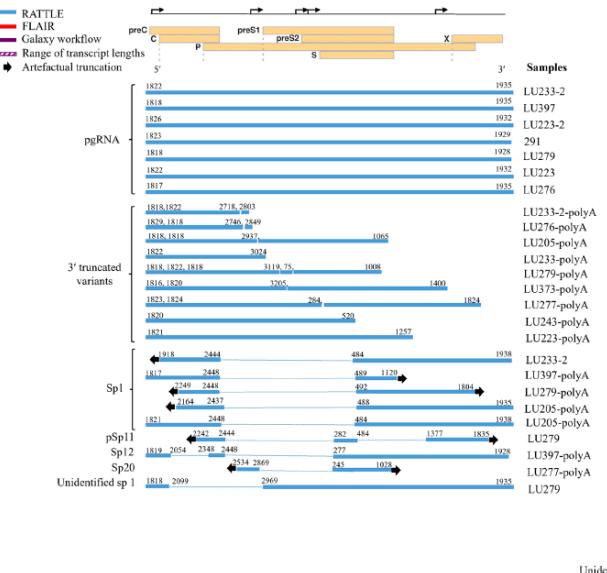
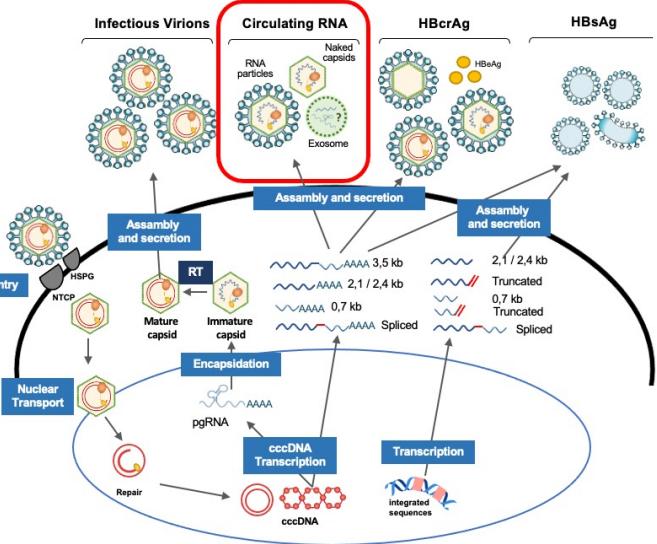
Stadelmayer B, 2020

and others...

The molecular biology of HBV RNAs: insights on new biomarkers

serum HBV-RNAs isoform composition

Nanopore long reads RNA sequencing



- HBV pregenomic RNA (pgRNA) most frequently observed transcript isoform (93.8% of patient samples)
- other detected transcripts included pgRNA spliced variants, 3' truncated variants and HBx mRNA
- spliced variants of pgRNA observed in HBV gtB, C, E, or F-infected patients
- Sp1 is the most frequently detected splice variant
- longitudinal sampling in antiviral-treated individuals showed increasing proportions of 3' truncated pgRNAs variants over time.

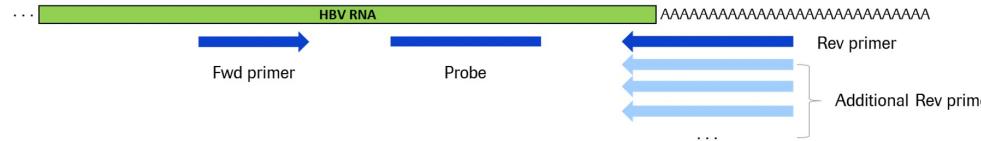
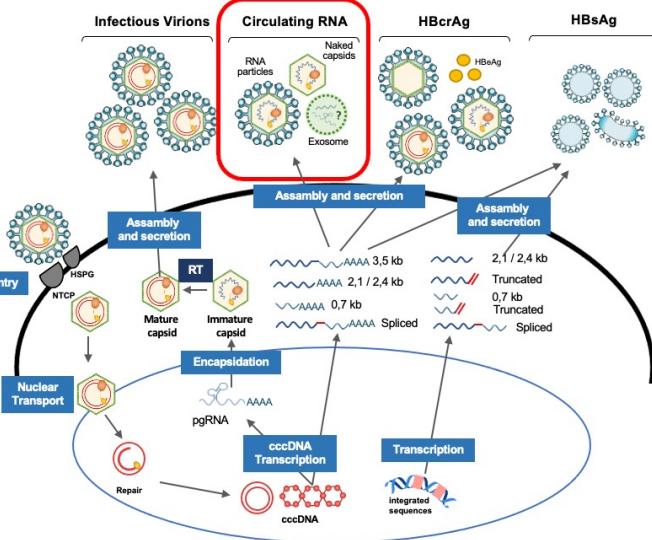
Vachon, 2018

The molecular biology of HBV RNAs: insights on new biomarkers

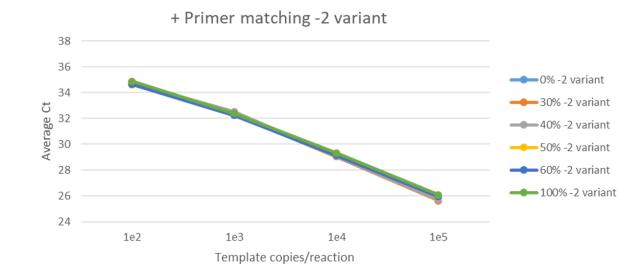
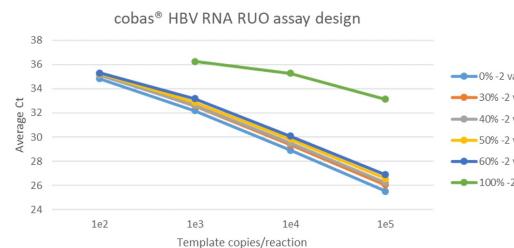
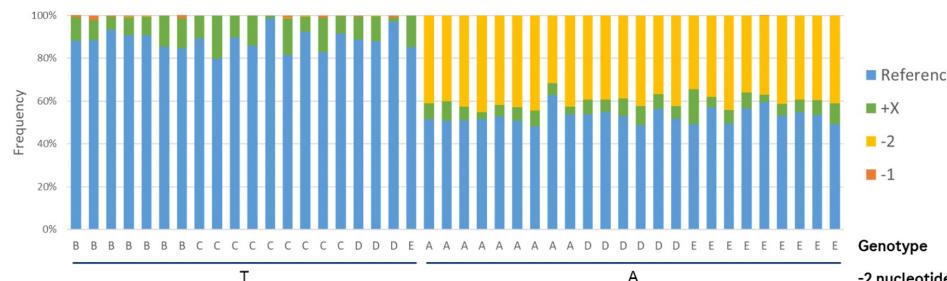
serum HBV-RNAs isoform composition

HBV capture + Illumina NGS

- Performance of the cobas® HBV RNA assay for use on the cobas® 5800/6800/8800 Systems (RUO) against genomic variants and transcript heterogeneity



Poly-A start site												
Reference	G	A	A	T	T	T	G	G	A	G	C	T
-2	G	A	A	T	T	T	G	G	A	G	C	T
-1	G	A	A	T	T	T	G	G	A	G	C	T
+1	G	A	A	T	T	G	G	A	G	C	T	T
+4	G	A	A	T	T	T	G	G	A	G	C	T
+5	G	A	A	T	T	T	G	G	A	G	C	T
+9	G	A	A	T	T	T	G	G	A	G	C	T



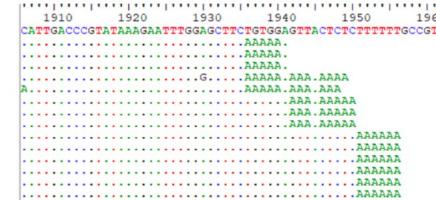
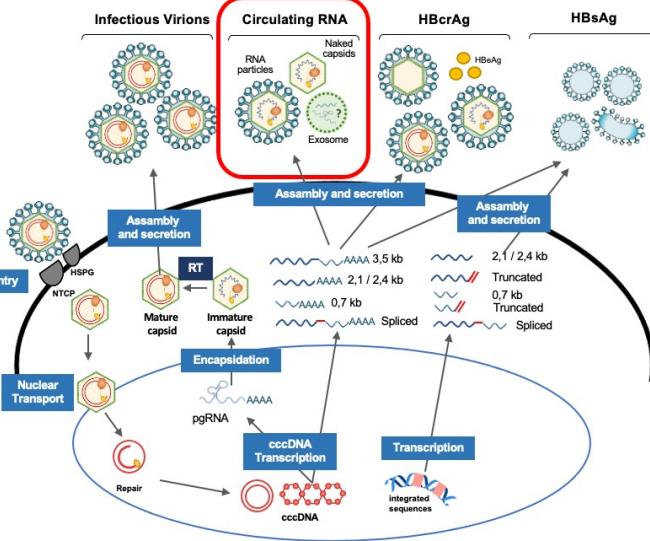
Meister, EASL 2023

The molecular biology of HBV RNAs: insights on new biomarkers

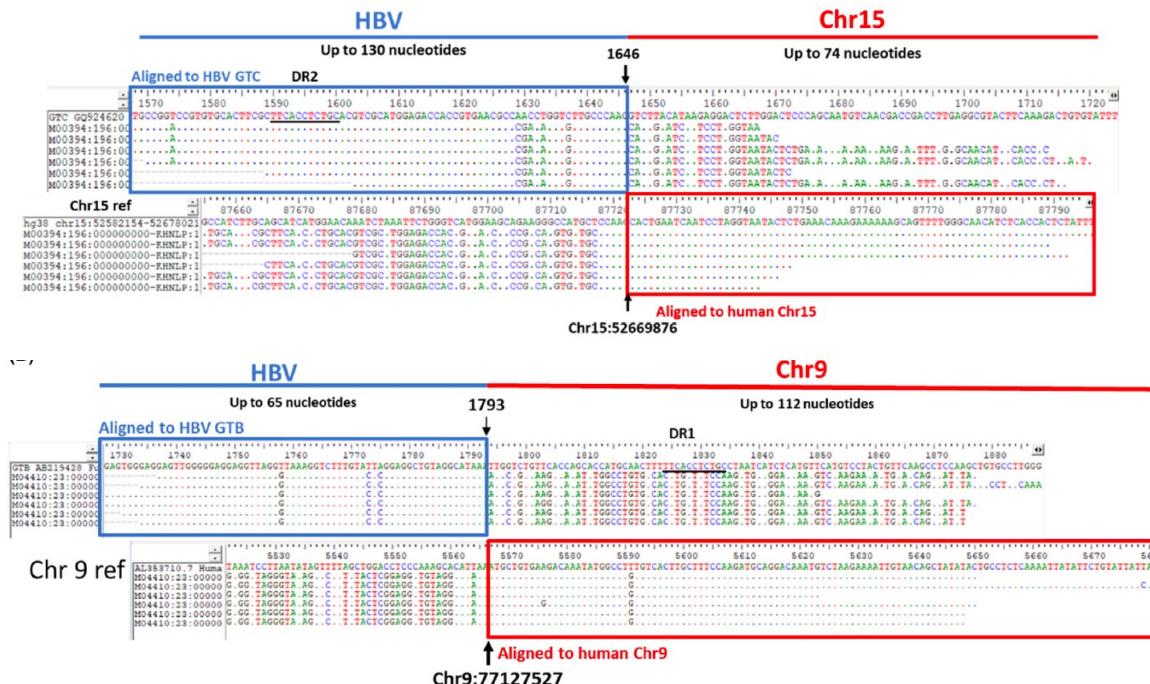
serum HBV-RNAs isoform composition

total RNA amplification coupled to NGS

- splice variants and poly-A tail variability



- 6 HBV-Human chimeric junction observed in 2 RNA samples from patient plasma



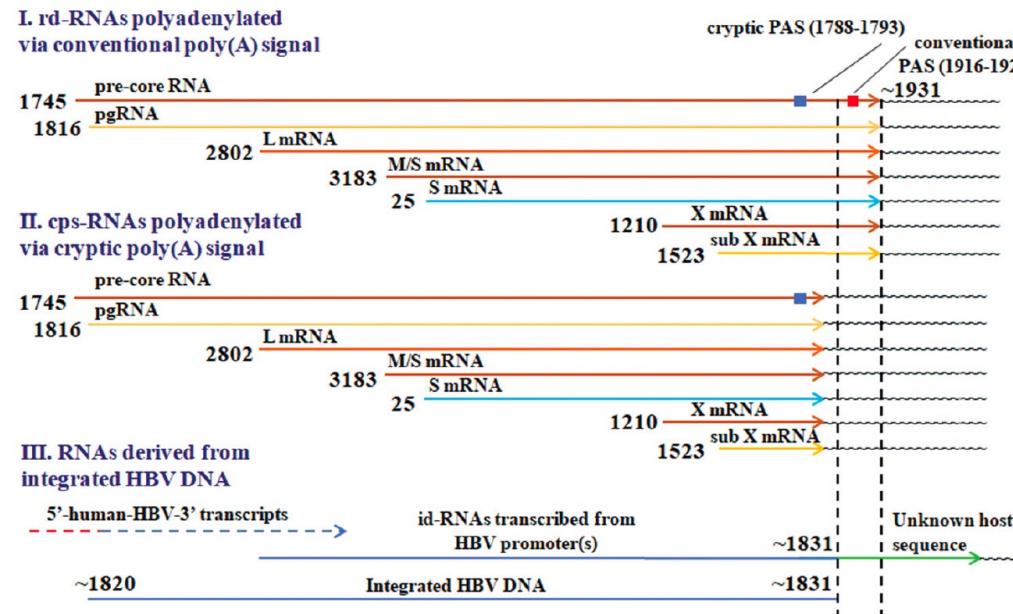
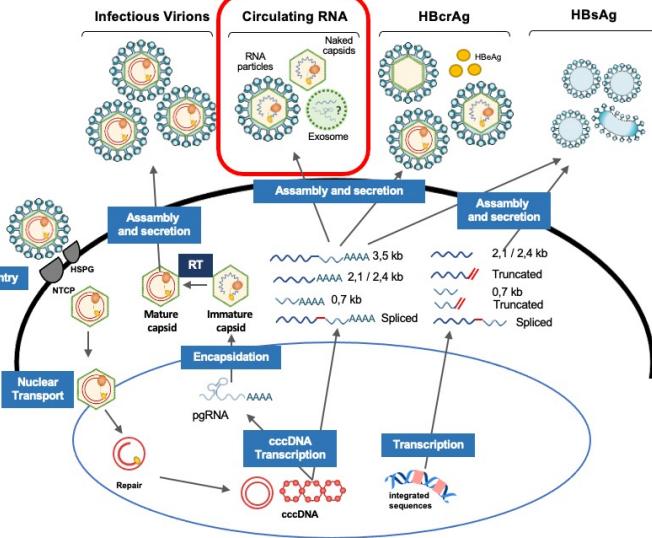
Chang, 2022

The molecular biology of HBV RNAs: insights on new biomarkers

serum HBV-RNAs isoform composition

HBV RNA amplification coupled to NGS

- both HBV replication-derived RNAs (rd-RNAs) and few (6) RNAs transcribed from integrated HBV DNA, including 5'-HBV-human-3' RNAs (integrand-derived RNAs [id-RNAs]) and 5'-human-HBV-3' transcripts,
- spliced HBV RNAs abundant in 50% of analyzed samples
- most serum rd-RNAs were polyadenylated via the conventional poly-A signal
- pregenomic RNA (pgRNA) as the major component
- the vast majority of rd-RNAs and pgRNA are associated with HBV virions but not with unenveloped capsids, exosomes, classic microvesicles, or apoptotic vesicles and bodies

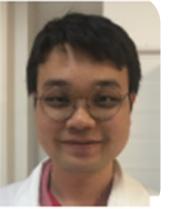


Gudima, 2023

The molecular biology of HBV RNAs: insights on new biomarkers



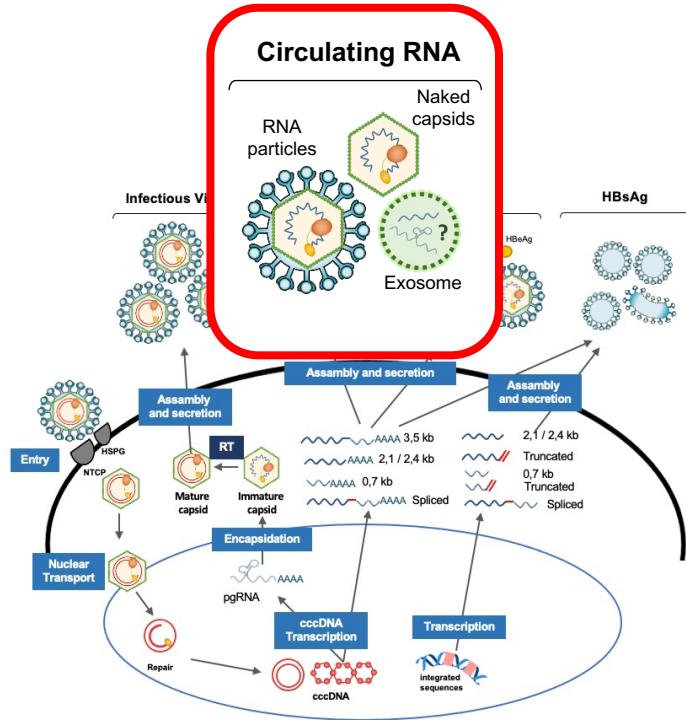
Delphine Bousquet



DooHyun Kim



Hyoseon Tak



Characterization of circulating HBV RNAs in CHB patients
and mechanism of cellular export

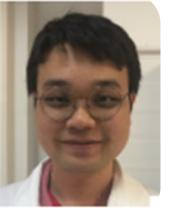
RNA Species	Reference
pgRNA	<ul style="list-style-type: none">Wang J, 2016Jansen L, 2016Prakash K, 2018Lam AM, 2017Stadelmayer B, 2020
Spliced Variants	<ul style="list-style-type: none">Lam AM, 2017Wang J, 2017Stadelmayer B, 2020
3' Truncated	<ul style="list-style-type: none">Hacker HJ, 2004Wang J, 2017
HBx RNA	<ul style="list-style-type: none">Niu C, 2017Stadelmayer B, 2020

Where ?	Reference
Virion « Like » Particles	<ul style="list-style-type: none">Wang J, 2016Bai L, 2018
Naked Capsids	<ul style="list-style-type: none">Bai L, 2018
Capsid Antibody Complexes	<ul style="list-style-type: none">Bai L, 2018
No Free	<ul style="list-style-type: none">Jansen L, 2016

The molecular biology of HBV RNAs: insights on new biomarkers



Delphine Bousquet

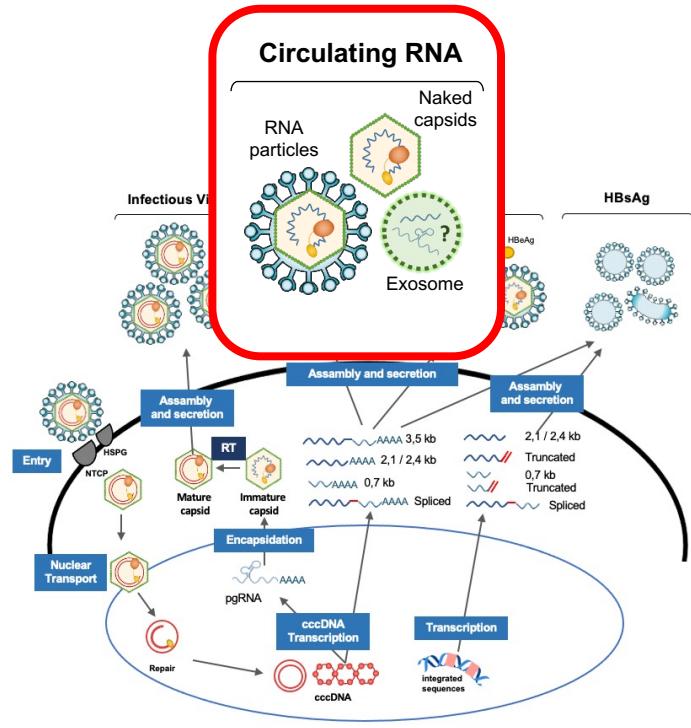


DooHyun Kim



Hyoseon Tak

Characterization of circulating HBV RNAs in CHB patients
and mechanism of cellular export



Species?

pgRNA



Spliced pgRNA



HBs RNA

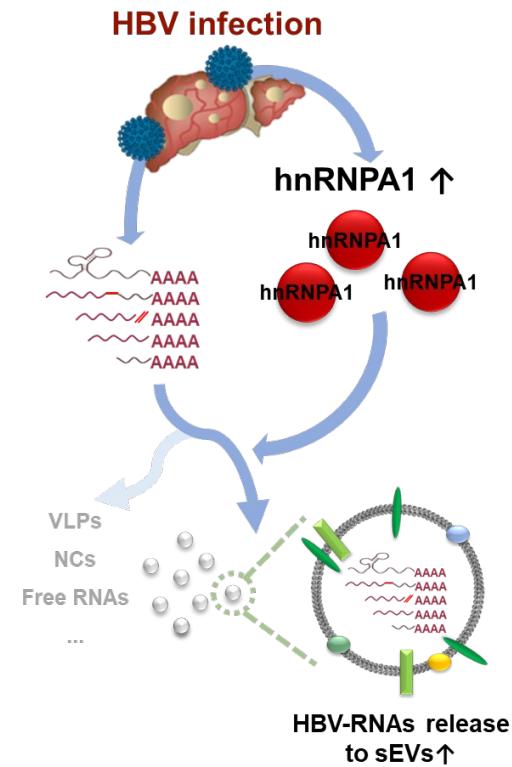


HBx RNA



Differ according to CHB stage

mechanism?



Distribution?



VLPs

sEVs

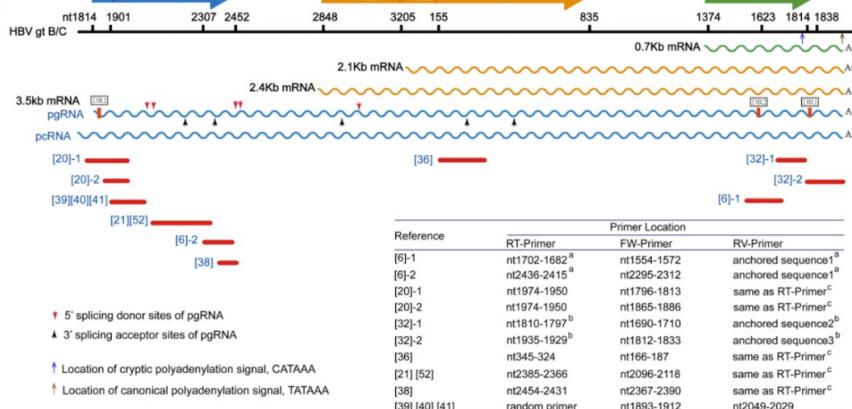
Vehicles?

- High viral load & HBsAg: **VLPs, EVs, and NCs**
- Low viral load & HBsAg: **EVs**

The molecular biology of HBV RNAs: insights on new biomarkers

Multiple assays : different performances ? comparable results?

1. Academic / home made assays



Adapted from Liu, Hepatology 2019

2. Industrial assays

Abbott ROU investigational assay (IA)

HEPATOTOLOGY



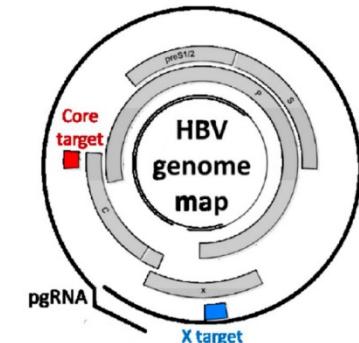
HEPATOTOLOGY, VOL. 00, NO. 00, 2018

doi: 10.1002/hep.30062. [Epub ahead of print]

Hepatitis B Virus Serum DNA and RNA Levels in Nucleos(t)ide Analog-Treated or Untreated Patients During Chronic and Acute Infection

Emily K. Butler,¹ Jeffrey Gersch,¹ Anne McNamara,¹ Ka-Cheung Luk,¹ Vera Holzmayer,¹ Maria de Medina,² Eugene Schiff,¹ Mary Kuhns,¹ and Gavin A. Cloherty¹

calibration: WHO HBV DNA



Roche @Cobas 6800/8800 investigational assay (IA)



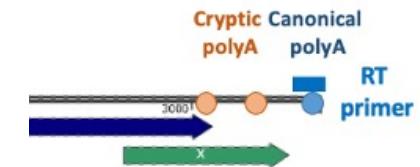
Journal of Clinical Virology

Volumes 150–151, June 2022, 105150



Performance of the cobas® HBV RNA automated investigational assay for the detection and quantification of circulating HBV RNA in chronic HBV patients

Caroline Scholtès,^{a,b,c} Aaron T. Hamilton,^d, Marie-Laure Plissonnier,^a, Caroline Charre,^{a,b}, Beth Scott,^d, Ling Wang,^d, Françoise Berby,^e, Janine French,^f, Barbara Testoni,^a, Alan Blair,^d, Miroslava Subic,^f, Matthias Hoppler,^e, Andreas Lankenau,^e, Andreas Grubenmann,^e, Massimo Leviero,^{a,b,f,g}, Marynthia L. Heil,^d, Fabien Zoulim,^{a,b}



calibration: « armored RNAs »

The molecular biology of HBV RNAs: insights on new biomarkers

Multiple assays : different performances ? comparable results?

1. Academic / home made assays



Alexia Paturel

Francesca Casuscelli

Development and characterization of a stable clonal cell line secreting HBV-RNAs as a potential RNA standard for (all) HBV RNA assays

Roche @Cobas 6800/8800 investigational assay (IA)

Journal of Clinical Virology

Volumes 150–151, June 2022, 105150

Performance of the cobas® HBV RNA automated investigational assay for the detection and quantification of circulating HBV RNA in chronic HBV patients

Caroline Scholtès ^{a b c}, Aaron T. Hamilton ^d, Marie-Laure Plissonnier ^a, Caroline Charre ^{a b}, Beth Scott ^d, Ling Wang ^d, Françoise Berby ^e, Janine French ^f, Barbara Testoni ^a, Alan Blair ^d, Miroslava Subic ^f, Matthias Hoppler ^e, Andreas Lankenau ^e, Andreas Grubenmann ^e, Massimo Leviero ^{a b f g}, Marynthia L. Heil ^d, Fabien Zoulim ^{a b}

calibration: « armored RNAs »

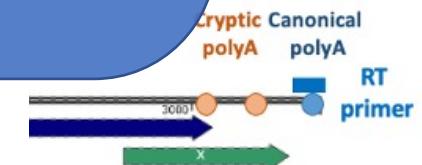
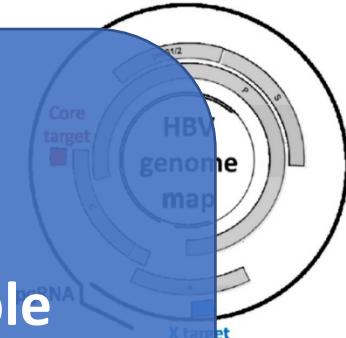
2. Industrial assays

Abbott ROU investigational assay (IA)

HEPATOTOLOGY
HEPATOTOLOGY, VOL. 00, NO. 00, 2018
doi: 10.1002/hep.30262. [Epub ahead of print]

Hepatitis B Virus Serum DNA and RNA Levels in Nucleos(t)ide Analog-Treated or Untreated Patients During Chronic and Acute Infection

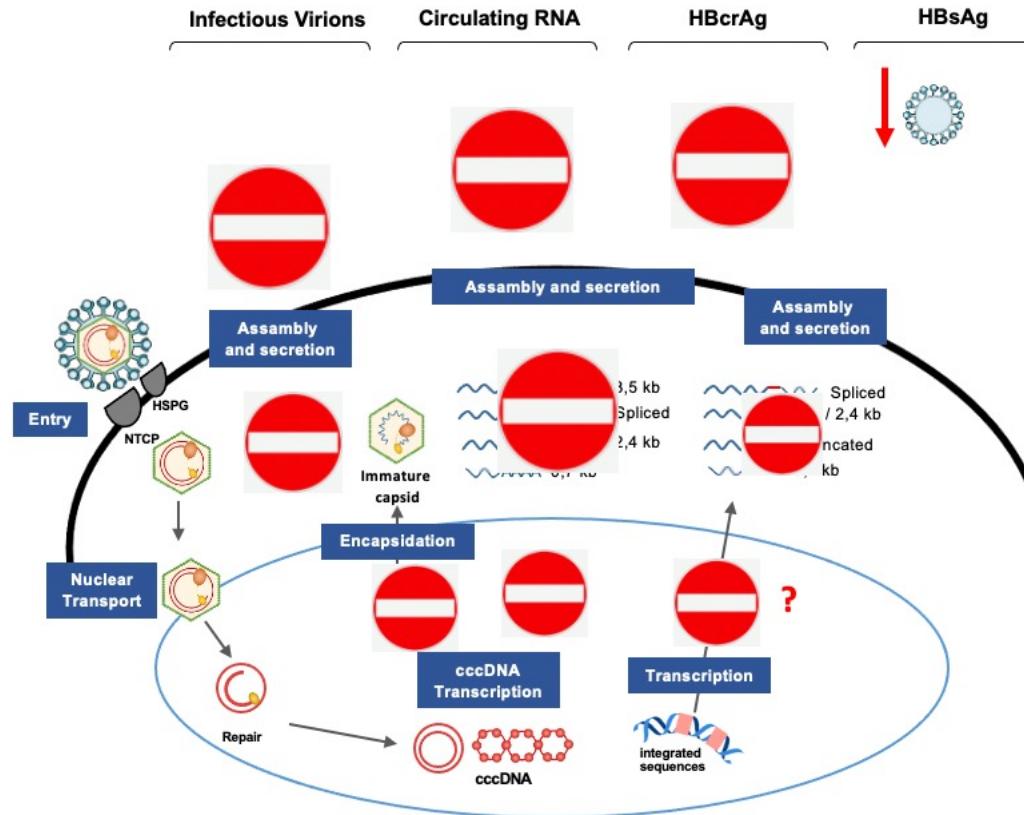
Mary Kalish,^a and Gavin A. Collier,^b



The molecular biology of HBV RNAs: insights on new biomarkers

Serum HBV RNAs in CHB treated patients: target engagement vs endpoints prediction

IFN α (IFN λ)

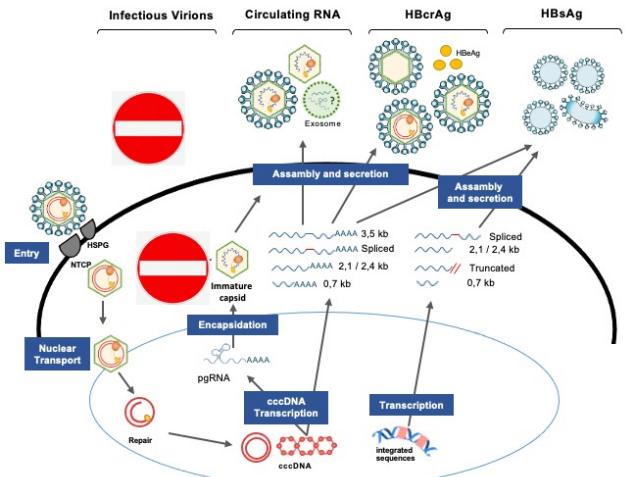


- direct targeting cccDNA transcription
- direct effect on capsid (and HBV replication)
- cccDNA destabilization not proven in patients
- contribution of immuno-modulation
- variable from patient to patient

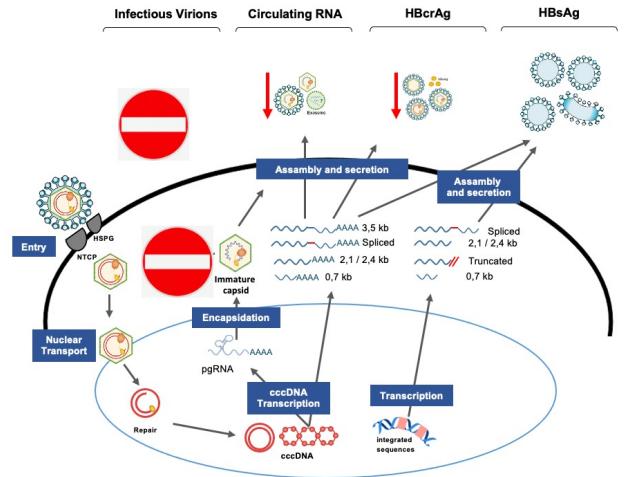
The molecular biology of HBV RNAs: insights on new biomarkers

Serum HBV RNAs in CHB treated patients: target engagement vs endpoints prediction

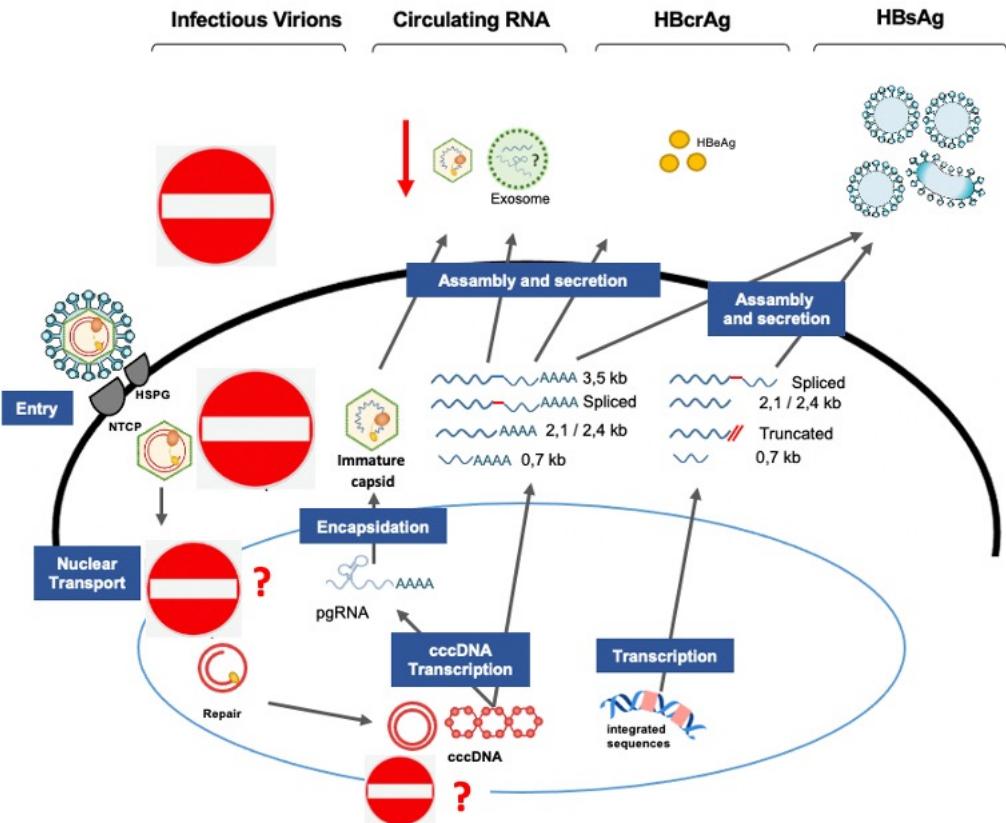
NUCs (early)



NUCs (late)



CAMs

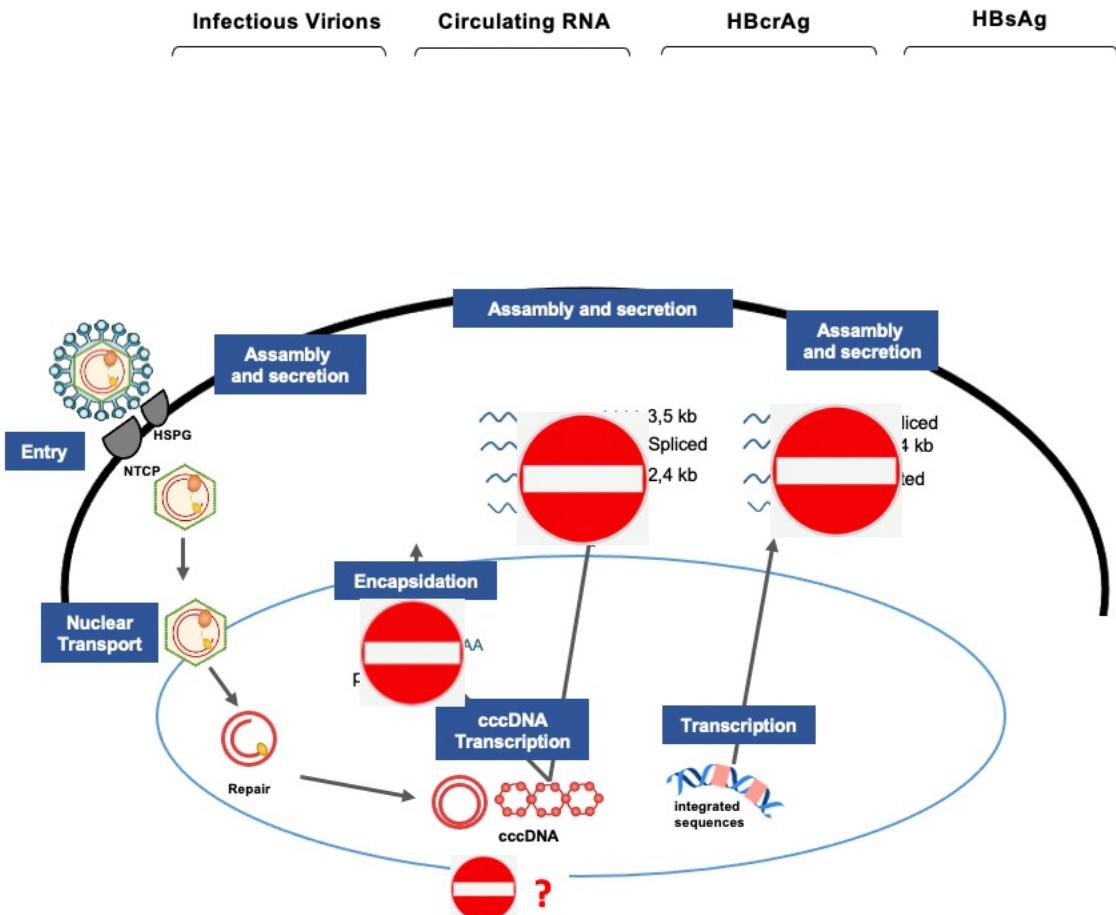


- mainly target engagement
- reduction of cccDNA pool over time
- direct effect on cccDNA activity not established

The molecular biology of HBV RNAs: insights on new biomarkers

Serum HBV RNAs in CHB treated patients: target engagement vs endpoints prediction

siRNAs (ASO, RNA destabilizers)

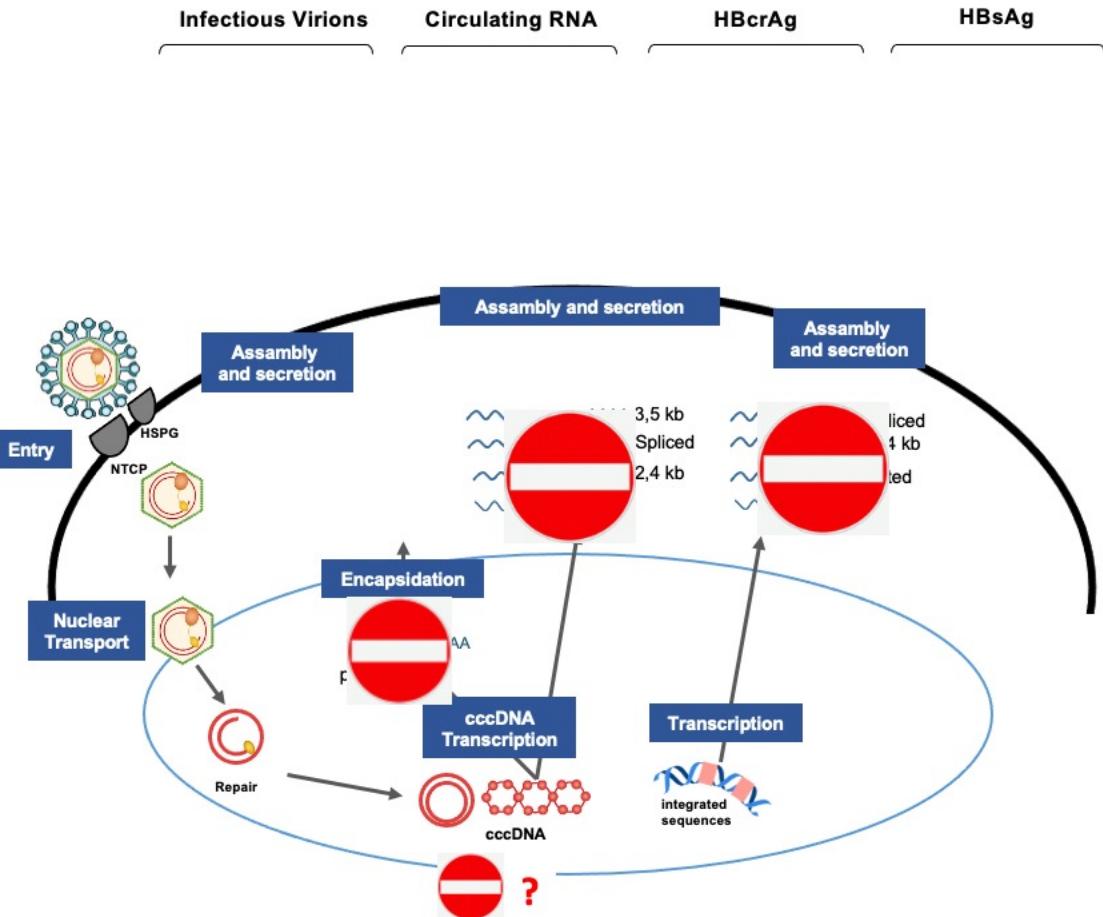


- **target engagement**
- effect on cccDNA pool size not established
- effect on cccDNA activity not established

The molecular biology of HBV RNAs: insights on new biomarkers

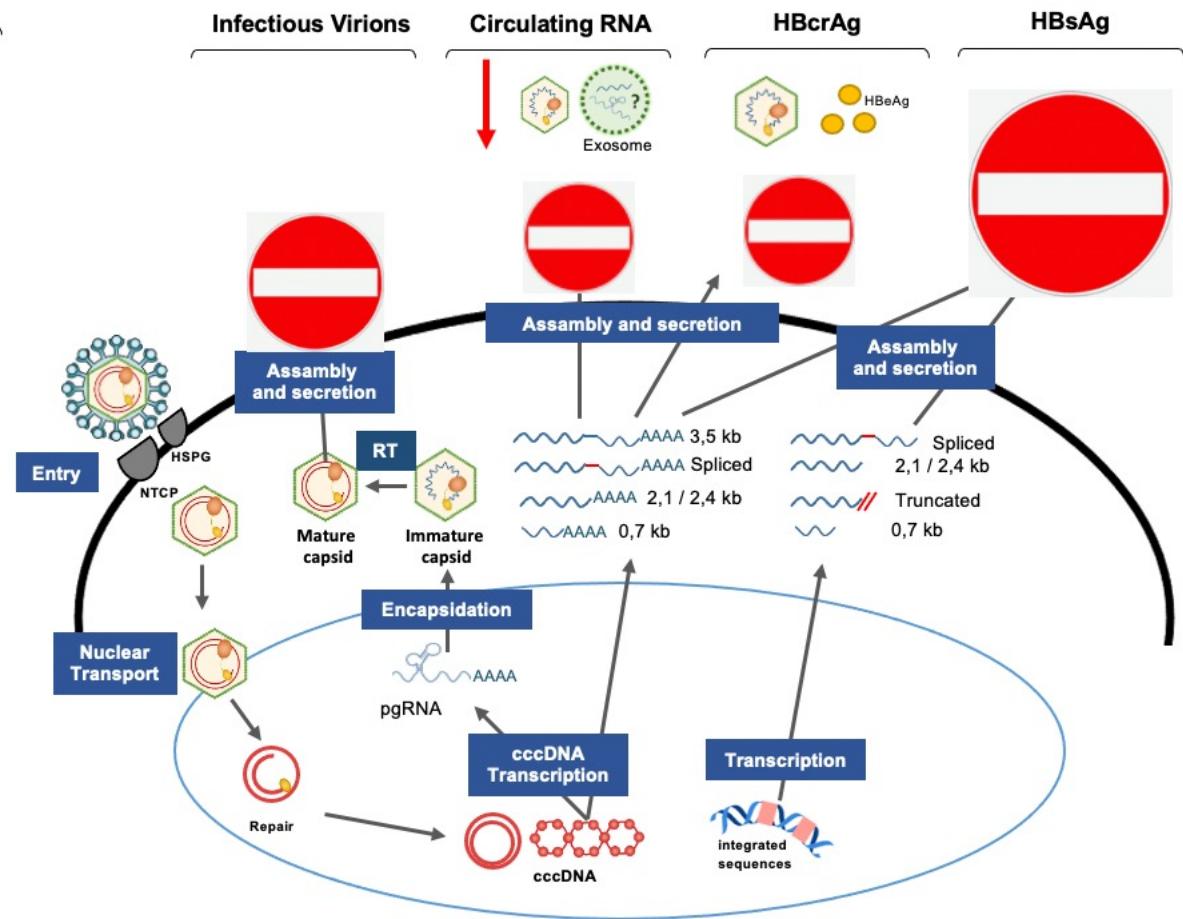
Serum HBV RNAs in CHB treated patients: target engagement vs endpoints prediction

siRNAs (ASO, RNA destabilizers)



- target engagement
- effect on cccDNA pool size not established
- effect on cccDNA activity not established

Inhibitors of HBsAg release

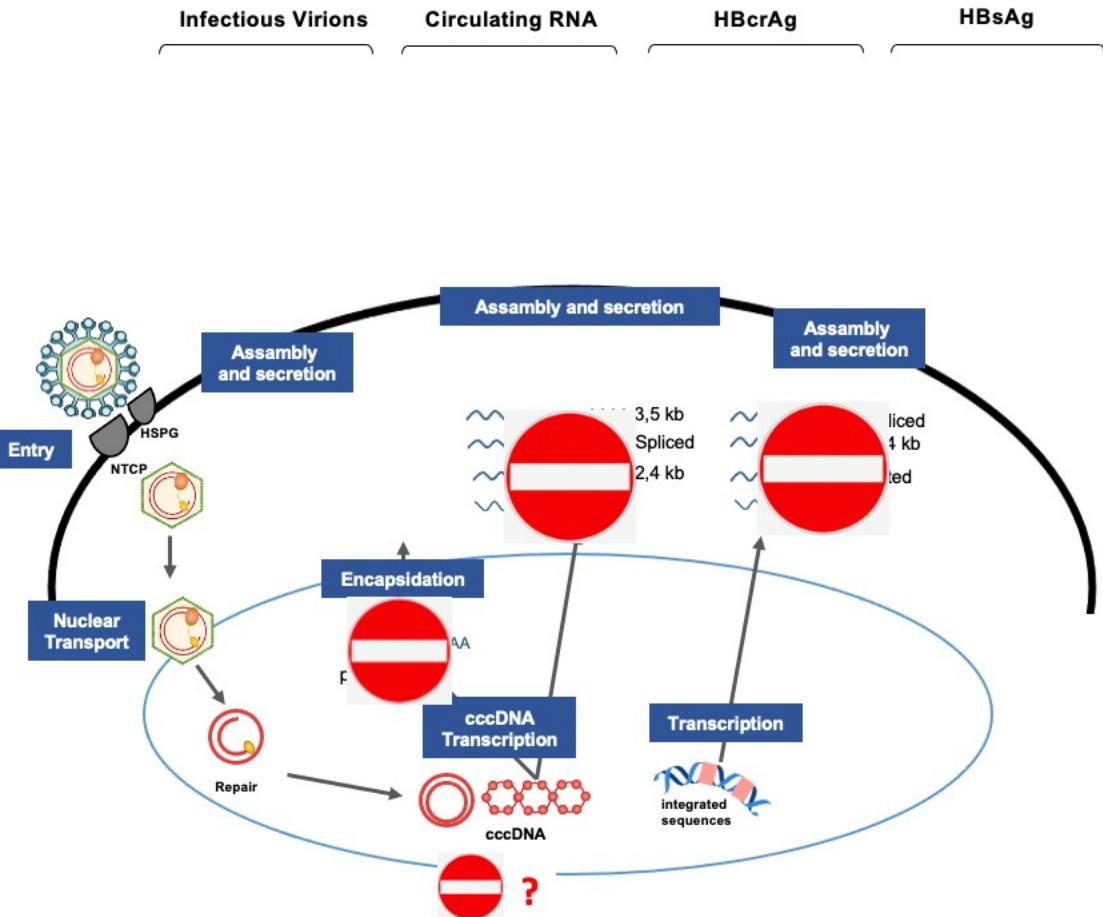


- mainly target engagement
- effect on cccDNA pool size not established
- effect on cccDNA activity not established

The molecular biology of HBV RNAs: insights on new biomarkers

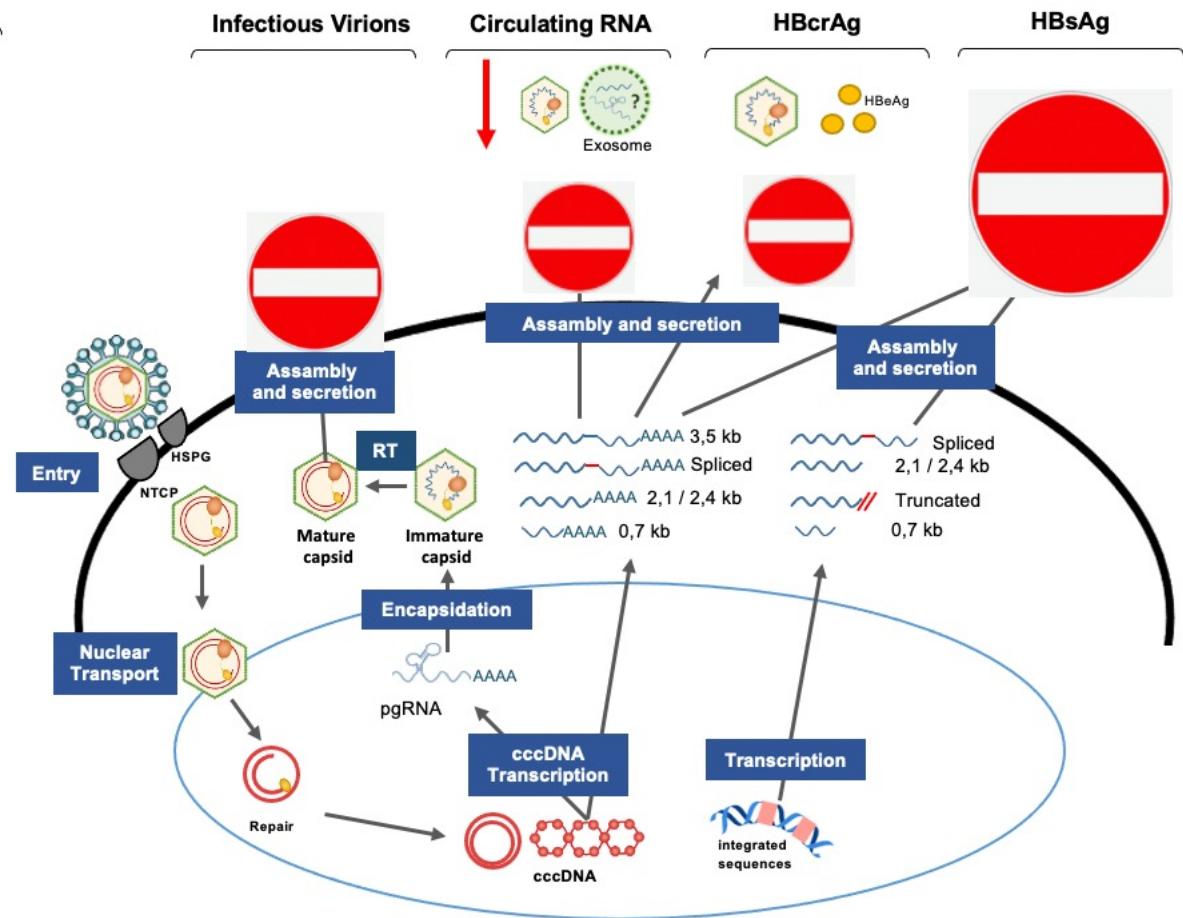
Serum HBV RNAs in CHB treated patients: target engagement vs endpoints prediction

siRNAs (ASO, RNA destabilizers)



- target engagement
- effect on cccDNA pool size not established
- effect on cccDNA activity not established

Inhibitors of HBsAg release



- mainly target engagement
- effect on cccDNA pool size not established
- effect on cccDNA activity not established

RHU « CirB-RNA »
ANR-17-RHUS-0003

International Workshop on Viral Biomarkers

September 7, 2023
Institut Lumière, Lyon

Thank you for your attention