

## Journal of Hepatology

### CTAT methods

Tables for a “Complete, Transparent, Accurate and Timely account” (CTAT) are now mandatory for all revised submissions. The aim is to enhance the reproducibility of methods.

- Only include the parts relevant to your study
- Refer to the CTAT in the main text as ‘Supplementary CTAT Table’
- Do not add subheadings
- Add as many rows as needed to include all information
- Only include one item per row

**If the CTAT form is not relevant to your study, please outline the reasons why:**

All this information is included in the manuscript as supplementary data.
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#### 1.1 Antibodies

Name	Citation	Supplier	Cat no.	Clone no.
CD3		BD	565515	UCHT1
CD4		BD	560768	RPA-T4
CD8		ThermoFisher	56-0086-42	OKT8
CD19		Biologend	363008	SJ25C1
CD56		Biologend	318334	HCD56
Live/Dead		ThermoFisher	L10119	
CD107a		Biologend	328624	H4A3
IFN $\gamma$		Biologend	502525	4S.B3
TNF $\alpha$		Biologend	502946	MAB11

#### 1.2 Sequence based reagents

Name	Sequence	Supplier
HBV overlapping peptides (pangenotypic or genotype D)	On request	Gilead & Peptide Synthesis Core Facility, UPF
ProMix™ CEF Peptide Pool	PX-CEF	Proimmune
Human b-globin assay	Hs00758889_s1	Thermo Fisher
Total HBV DNA assay	Pa03453406_s1	Thermo Fisher
iHBV-RNA forward	GGTCCCCTAGAAGAAGAACTCCCT	Biomers
iHBV-RNA reverse	CATTGAGATTCCCGAGATTGAGAT	Biomers
iHBV-RNA probe	TCTCAATCGCCGCGTCGCAGA	Biomers
cccDNA forward	CCGTGTGCACTTCGCTTCA	Biomers
cccDNA reverse	GCACAGCTTGGAGGCTTGA	Biomers
cccDNA probe	CATGGAGACCACCGTGAACGCC	Biomers
HBV_RT_F	TTCTTAGCGTATTGGAGCTC CTGCTGGTGGCTCCAGTT	Biomers
HBV_RT_R	AAGTATCAGTATTGCGTTGTT GCTAGGAGTTCCGCAGTATGG	Biomers
HBV_RT_F2	TTCTTAGCGTATTGGAGCTC	Biomers
HBV_RT_R2	AAGTATCAGTATTGCGTTGTT	Biomers
Internal RNA control assay	Int-RNA-VIC	Primer Design

## 1.3 Biological samples

Description	Source	Identifier
Serum samples Liver biopsies Peripheral blood mononuclear cells	Liver Unit, Hospital Clínic Barcelona, Spain	

## 1.4 Software

Software name	Manufacturer	Version
FlowJo	BD Bioscience	10
R	The R Foundation for Statistical Computing,	3.4
MEGA X	MEGA X	10
GraphPad Prism	GraphPad Software, Inc	8.1
SPSS	IBM	20

## 1.5 Please provide the details of the corresponding methods author for the manuscript:

Sofía Pérez-del-Pulgar  
Liver Unit, Hospital Clínic, IDIBAPS, Barcelona, Spain.  
Tel.: +34 2275400, ext. 2093.  
Fax: +34 933129405.  
E-mail address: [sofiapp@clinic.cat](mailto:sofiapp@clinic.cat)