

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:

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1.1 Antibodies

Name	Citation	Supplier	Cat no.	Clone no.

1.2 Cell lines

Name	Citation	Supplier	Cat no.	Passage no.	Authentication test method

1.3 Organisms

Name	Citation	Supplier	Strain	Sex	Age	Overall n number

1.4 Sequence based reagents

Name	Sequence	Supplier
AB 3730 XL DNA Analyzer using Big Dye 3.1 terminator chemistry	Sanger	Thermo Fisher Scientific, Burlington, ON, Canada

1.5 Biological samples

Description	Source	Identifier
Serum or plasma	human	

1.6 Deposited data

Name of repository	Identifier	Link
GenBank	Accession numbers MZ712366-MZ712572	Once processed by NCBI, the data will be available

		at Home - Nucleotide - NCBI (nih.gov)
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1.7 Software

Software name	Manufacturer	Version
MAFFT	https://mafft.cbrc.jp/alignment/server/index.html	7
DIVEIN	https://indra.mullins.microbiol.washington.edu/DIVEIN/	
GraphPad Prism	GraphPad Software	9.0.0

1.8 Other (e.g. drugs, proteins, vectors etc.)

1.9 Please provide the details of the corresponding methods author for the manuscript:

HDV antibody and RNA detection and sequencing: Carla Osiowy
(Carla.osiowy@phac-aspc.gc.ca)
Patient data collection and statistical analysis: Sarah Haylock-Jacobs
(s.haylockjacobs@gmail.com)

2.0 Please confirm for randomised controlled trials all versions of the clinical protocol are included in the submission. These will be published online as supplementary information.

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