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:

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
TaqMan SNP assay for rs738409	C_____7241_10	Thermo Fisher Scientific
TaqMan SNP assay for rs58542926	C__89463510_10	Thermo Fisher Scientific
TaqMan SNP assay for rs641738	C_____8716820_10	Thermo Fisher Scientific
TaqMan SNP assay for rs1260326	C___2862880_1_	Thermo Fisher Scientific
Custom TaqMan SNP assay for rs72613567	See doi.org/10.1002/hep.30799	Thermo Fisher Scientific

1.5   Biological samples

Description	Source	Identifier
DNA from PBMC of 2,564	Multicentric study, leading	NAFLD/MAFLD cohort

patients with NAFLD/MAFLD and healthy controls (see Methods – patients section, and Supplementary Methods – study cohort characterization)	center: Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico Milano (see Methods – patients section, and Supplementary Methods – study cohort characterization)	
DNA from PBMC of 427 patients with NAFLD/MAFLD (see Methods – patients section, and Supplementary Methods – study cohort characterization)	Patients referred to the University Clinic of Leipzig and Universital Dresden (see Methods – patients section, and Supplementary Methods – study cohort characterization)	German NAFLD/MAFLD cohort

1.6   Deposited data

Name of repository	Identifier	Link
UK BioBank	UKBB	https://www.ukbiobank.ac.uk/
1000 Genome Project	1000 Genome Project	https://www.internationalgenome.org/

1.7   Software

Software name	Manufacturer	Version
JMP Pro Statystical Analysis Software	SAS Institute, Cary, NC	14.0
R	CRAN	3.5.2

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

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

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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.