

LETTER

Risk of severe illness from COVID-19 in patients with metabolic dysfunction-associated fatty liver disease and increased fibrosis scores

A recent study reported that patients with severe COVID-19 were more likely to have non-alcoholic fatty liver disease (NAFLD) compared with those with non-severe COVID-19 illness.¹ However, the prognosis of NAFLD (recently renamed metabolic dysfunction-associated fatty liver disease (MAFLD)²) is determined by the severity of liver fibrosis.^{3,4} We therefore postulated that patients with MAFLD with increased non-invasive liver fibrosis scores are at higher risk of severe illness from COVID-19.

We studied 310 patients with laboratory-confirmed COVID-19 who were consecutively hospitalised at four sites in Zhejiang Province, China, between January and February 2020. Some of these patients (n=150) have been included in a prior study examining the association between obesity and COVID-19 severity.⁵ Patients with viral hepatitis, excessive alcohol consumption, chronic pulmonary diseases or active cancers were excluded. Clinical and laboratory data were collected at hospital admission. All patients were screened for hepatic steatosis by computed tomography and subsequently diagnosed as MAFLD.⁶ The originally validated cut-points for fibrosis-4 (FIB-4) index and NAFLD fibrosis score (NFS) were used to categorise liver fibrosis probability as low, intermediate or high.⁷ COVID-19 severity was classified as severe and non-severe.⁸ The study protocol was approved by the local ethics committees of the four hospitals.

In our cohort of 310 confirmed cases of COVID-19, 94 (30.3%) patients had MAFLD. As shown in table 1, patients with MAFLD with intermediate or high FIB-4 scores were more likely to be older, obese, have diabetes and have higher NFS, higher liver enzymes, higher C reactive protein, as well as lower levels of lymphocyte count, platelet count, triglycerides and high-density lipoprotein cholesterol compared with their counterparts with low FIB-4 score or those without MAFLD. Notably, the severity of COVID-19 illness markedly increased among patients with MAFLD with intermediate or high FIB-4 scores.

The severity of COVID-19 illness was associated with intermediate (unadjusted OR 4.32, 95%CI 1.94 to 9.59) or high (unadjusted-OR 5.73, 95%CI 1.84 to 17.9)

Table 1 Main clinical and biochemical characteristics of patients with laboratory-confirmed COVID-19, stratified by the presence or absence of imaging-defined MAFLD with increasing levels of FIB-4 score

	No MAFLD	MAFLD with low FIB-4 (≤ 1.3)	MAFLD with intermediate FIB-4 (1.3–2.67)	MAFLD with high FIB-4 (>2.67)	P value
N	216	44	36	14	
Age (years)	45.9±15.4	41.2±14.2	54.2±10.8	59.9±9.1	<0.001
Male sex (%)	43.5	54.6	63.9	57.1	0.086
BMI (kg/m ²)	23.0±3.2	26.5±4.7	26.6±3.2	26.1±2.8	<0.001
Obesity (%)	25.9	61.4	75.0	64.3	<0.001
Current smokers (%)	8.8	2.3	11.1	7.1	0.768
Systolic blood pressure (mm Hg)	130±17	136±15	136±16	136±14	0.047
Diastolic blood pressure (mm Hg)	80±11	84±11	82±11	79±8	0.117
Prior diabetes (%)	7.4	20.4	13.9	28.6	0.010
White blood count ($\times 10^9/L$)	4.90 (3.9–6.2)	5.35 (4.5–6.7)	4.69 (3.8–6.6)	4.90 (3.2–5.8)	0.254
Neutrophil count ($\times 10^9/L$)	3.05 (2.2–4.1)	3.49 (2.6–4.2)	3.0 (2.4–4.7)	3.20 (1.9–4.5)	0.759
Lymphocyte count ($\times 10^9/L$)	1.20 (0.9–1.6)	1.40 (1.2–2.0)	1.10 (0.7–1.3)	0.92 (0.7–1.14)	<0.001
Haemoglobin (g/L)	132.4±16	136.6±15	135.5±12	135.8±17	0.305
Platelet count ($\times 100\ 000/mm^3$)	211±72	246±75	183±64	125±24	<0.001
Prothrombin time (s), n=211	12.1±1.3	11.9±2.3	12.2±1.2	12.3±1.1	0.754
APTT (s), n=211	31.5±3.8	32.7±5.4	32.8±3.7	35.3±7.4	0.026
D-dimer (mg/L), n=183	0.18 (0.11–0.29)	0.21 (0.1–0.5)	0.21 (0.1–0.4)	0.27 (0.1–0.4)	0.432
C reactive protein (mg/L)	7.9 (1.9–24.9)	8.6 (2.9–23.6)	25.4 (8.7–53.3)	25.5 (8.7–53.4)	<0.001
Procalcitonin (ng/mL), n=190	0.05 (0.03–0.25)	0.08 (0.04–0.25)	0.06 (0.04–0.10)	0.20 (0.08–0.25)	0.077
Albumin (g/L), n=286	41.4 (38.0–44.5)	42.3 (37.3–44.6)	41.0 (36.9–43.1)	38.6 (37.7–42.8)	0.348
Total bilirubin ($\mu\text{mol/L}$)	10.5 (7.0–15.5)	10.1 (7.5–15.6)	12.0 (9.5–16.6)	15.3 (11.8–17.3)	0.025
AST (IU/L)	22 (17–27)	23 (19–32)	32 (26–50)	44 (29–83)	<0.001
ALT (IU/L)	18 (13–27)	30 (22–52)	28 (22–48)	27 (16–82)	<0.001
GGT (IU/L)	21 (14–33)	31 (22–54)	51 (24–81)	49 (32–102)	<0.001
Elevated AST >40 IU/L (%)	7.9	9.1	27.8	57.1	<0.001
Elevated ALT >40 IU/L (%)	13.0	29.6	30.6	42.9	<0.001
Total cholesterol (mmol/L)	3.98±0.8	4.11±0.9	3.73±0.7	3.82±1.0	0.217
Triglycerides (mmol/L)	1.15 (0.9–1.7)	1.61 (1.0–2.1)	1.48 (1.1–1.8)	1.11 (0.9–1.6)	<0.005
HDL cholesterol (mmol/L)	1.18±0.4	0.96±0.2	1.06±0.2	1.09±0.4	<0.005
LDL cholesterol (mmol/L)	2.23±0.7	2.54±0.8	2.10±0.7	2.20±0.8	0.116
Hospital stay (days)	18 (13–24)	18 (13–22)	22 (16–29)	17 (6–23)	0.122
NAFLD fibrosis score, n=286	−1.82 (−2.8 to −1.0)	−2.61 (−3.1 to −1.9)	−0.68 (−1.4 to −0.2)	+0.26 (−0.03 to 1.6)	<0.005
COVID-19 severity					<0.001
Non-severe (%)	88.4	86.4	63.9	57.1	
Severe (%)	11.6	13.6	36.1	42.9	

Sample size, n=310, except where indicated. Diabetes was diagnosed as self-reported history of disease and/or specific drug treatment. Obesity was diagnosed as BMI >25 kg/m². Data are expressed as means±SD, medians and IQRs (in parentheses) or frequencies. Differences among the four groups of patients were tested by Fisher's exact test for categorical variables, one-way analysis of variance for normally distributed continuous variables or the Kruskal-Wallis test for not normally distributed continuous variables. For the sake of clarity, significant p values are highlighted in bold.

ALT, alanine aminotransferase; APTT, activated partial thromboplastin time; AST, aspartate aminotransferase; BMI, body mass index; FIB-4, fibrosis-4; GGT, gamma-glutamyltransferase; HDL, high-density lipoprotein cholesterol; LDL, low-density lipoprotein cholesterol; MAFLD, metabolic dysfunction-associated fatty liver disease; NAFLD, non-alcoholic fatty liver disease.

FIB4 scores among patients with MAFLD (table 2, model 1A). This association remained significant after adjusting for sex, obesity and diabetes (model 1B). We did not additionally adjust for age, because this variable was incorporated in the FIB-4 score. Similar to the main analysis, when the intermediate and high FIB-4 categories were combined, the risk of severe COVID-19 illness was increased with intermediate/high FIB-4 score in unadjusted (model 2A) and multivariate-adjusted analyses (model 2B).

Similarly, the intermediate/high NFS (unadjusted-OR 5.21, 95%CI 2.39 to 11.3) was associated with a higher risk of severe COVID-19 illness. This significant association persisted in multivariate-adjusted analyses after controlling for sex, obesity and

diabetes (adjusted OR 2.91, 95%CI 1.20 to 7.06).

When we included FIB-4 or NFS as continuous measures in multivariable regression models, increasing FIB-4 (adjusted OR 1.90, 95%CI 1.33 to 2.72) or NFS (adjusted OR 2.57, 95%CI 1.73 to 3.82) were significantly associated with greater COVID-19 severity, even after adjusting for sex, obesity, diabetes and presence/absence of MAFLD.

Our study has some limitations, including the relatively modest sample size, the Asian ancestry of the cohort and the use of non-invasive fibrosis scores without a histological diagnosis of liver fibrosis. Despite these limitations, our study is the first to examine the impact of FIB-4 or NFS on COVID-19 severity in patients with imaging-defined

Table 2 Association between imaging-defined MAFLD with increasing levels of FIB-4 score and risk of having severe illness associated with COVID-19

Logistic regression analyses	ORs	95% CI	P value
Severity of COVID-19 illness (mild/moderate vs severe/critical)			
Unadjusted model 1A			
MAFLD/FIB-4 status			
No MAFLD (n=216)	Ref.	Ref.	
MAFLD with low FIB-4 (≤ 1.3 ; n=44)	1.21	0.46 to 3.14	0.701
MAFLD with intermediate FIB-4 (1.3–2.67, n=36)	4.32	1.94 to 9.59	<0.001
MAFLD with high FIB-4 (>2.67, n=14)	5.73	1.84 to 17.9	<0.005
Adjusted model 1B			
MAFLD/FIB-4 status			
No MAFLD (n=216)	Ref.	Ref.	
MAFLD with low FIB-4 (≤ 1.3 , n=44)	0.82	0.30 to 2.24	0.696
MAFLD with intermediate FIB-4 (1.3–2.67, n=36)	2.59	1.09 to 6.13	0.030
MAFLD with high FIB-4 (>2.67, n=14)	4.04	1.22 to 13.3	0.021
Sex (men vs women)	1.78	0.93 to 3.44	0.079
Obesity (yes vs no)	2.62	1.31 to 5.24	<0.005
Prior diabetes (yes vs no)	1.04	0.40 to 2.80	0.928
Unadjusted model 2A			
MAFLD/FIB-4 status			
No MAFLD (n=216)	Ref.	Ref.	
MAFLD with low FIB-4 (≤ 1.3 , n=44)	1.21	0.46 to 3.14	0.701
MAFLD with intermediate/high FIB-4 (>1.3, n=50)	4.68	2.31 to 9.49	<0.001
Adjusted model 2B			
MAFLD/FIB-4 status			
No MAFLD (n=216)	Ref.	Ref.	
MAFLD with low FIB-4 (≤ 1.3 , n=44)	0.82	0.30 to 2.24	0.696
MAFLD with intermediate/high FIB-4 (>1.3, n=50)	2.95	1.37 to 6.34	<0.005
Sex (men vs women)	1.79	0.94 to 3.45	0.084
Obesity (yes vs no)	2.60	1.30 to 5.16	<0.005
Prior diabetes (yes vs no)	1.09	0.41 to 2.89	0.862

Sample size, n=310. Data are expressed as ORs and 95% CI as tested by univariable (unadjusted) and multivariable (adjusted) logistic regression analysis. Diabetes was diagnosed as self-reported history of disease and/or specific drug treatment. Obesity was diagnosed as BMI>25 kg/m². In the adjusted logistic regression models, we did not additionally adjust also for age, because this variable is already incorporated in the FIB-4 score. FIB-4, fibrosis-4; MAFLD, metabolic associated fatty liver disease; Ref, reference category.

MAFLD. These non-invasive fibrosis scores have been shown to predict histological fibrosis stage with reasonable accuracy in cohorts of patients with MAFLD,⁷ and are also associated with increased overall and disease-specific mortality in population-based studies.^{9,10} Our data demonstrate that patients with MAFLD with increased FIB-4 or NFS are at higher likelihood of having severe COVID-19 illness, irrespective of metabolic comorbidities. In the context of COVID-19, the presence of MAFLD with significant/advanced fibrosis might exacerbate the virus-induced cytokine ‘storm’, possibly through the hepatic release of multiple proinflammatory cytokines, thereby contributing mechanistically to severe COVID-19. Further research is needed to better understand the mechanistic link of advanced MAFLD to the viral disease process.

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