



# Clinical efficacy of Mac-2-binding protein glycosylation isomer as a biomarker for albumin-bilirubin grade and the Controlling Nutritional Status score in chronic liver disease: investigation of cut-off values by the type of chronic liver disease

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**Background:** Mac-2-binding protein glycosylation isomer (M2BPGi), a novel noninvasive biomarker for fibrosis, is a prognostic factor for liver disease; however, its relationship with hepatic function reserve and nutritional status remains unclear. Furthermore, the cut-off value of this marker varies with the underlying liver disease. This study aims to clarify that M2BPGi can be clinically used as a hepatic function reserve marker and nutritional index without pushing the search for alternative markers to the forefront in clinical practice as an important biomarker.

**Methods:** Seven hundred and forty-three outpatients with chronic liver disease (CLD) were enrolled. We evaluated the relationship among M2BPGi, albumin-bilirubin (ALBI) grade, and Controlling Nutritional Status (CONUT) score as nutritional status markers. Diagnostic performance of M2BPGi values in distinguishing different modified ALBI (mALBI) grade and CONUT score were compared using receiver operating characteristic (ROC) curve analysis.

**Results:** The M2BPGi level increased with ALBI and mALBI grades. The correlation coefficient ( $r^2$ ) between M2BPGi and ALBI grade was 0.40 ( $r=0.63$ ), indicating a positive correlation between M2BPGi and ALBI grade. The cut-off for M2BPGi to predict mALBI G1 *vs.* G2–G3 was 1.07, G1–2a *vs.* G2b–3 was 1.73, and mALBI G1–2 *vs.* G3 was 5.83 under the ROC curves. The cut-off for M2BPGi to predict CONUT score normal *vs.* light-severe was 1.60, normal-light *vs.* moderate-severe was 1.74, and normal-moderate *vs.* severe was 5.83 under the ROC curves. M2BPGi correlates with ALBI grade and is useful for diagnosing ROC analysis results, especially G2 and above. M2BPGi also correlates with the CONUT score and is useful for diagnosing ROC analysis results, especially moderate or higher. These results showed similar diagnostic performance regardless of the etiology of the background liver disease.

**Conclusions:** Although the predictive cut-off value varied with the type of liver disease, M2BPGi was found to be a single predict biomarker of ALBI and CONUT, and thus, is an effective indicator of CLD status. Further investigation is warranted to determine the clinical utility of this marker.

**Keywords:** Mac-2-binding protein glycosylation isomer (M2BPGi); chronic liver disease (CLD); Controlling Nutritional Status (CONUT); albumin-bilirubin (ALBI) grade; cut-off values

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## Introduction

Hepatic function reserve and nutritional status determine the prognosis for patients with chronic liver disease (CLD) (1-3). The Child-Turcotte classification was first published in 1964 as a prognostic criterion for cirrhosis and was used to classify patients receiving treatments such as transabdominal esophageal varicectomy or sclerotherapy. This classification was modified in 1973 by Pugh *et al.* as the Child-Pugh score (4). The major limitation of the Child-Pugh scoring system is that it includes several subjective parameters (hepatic encephalopathy and ascites) and interrelated parameters (ascites and serum albumin). Ascites can be easily influenced by diuretic use or dehydration state. Diagnosing minimal or covert hepatic encephalopathy involves difficulties. Hence, the albumin-bilirubin (ALBI) score uses only objective parameters, albumin (Alb) and total bilirubin (T. Bil), enabling a better evaluation. However, the limit of ALBI score is log calculation, so it is a little complicated in clinical use (5). In addition, the modified ALBI (mALBI) grade was proposed based on the relationship with indocyanine green values, and it has been utilized in clinical practice (6,7).

Nutritional status is also important for the assessment of CLD. The effect of immunological and nutritional status on the long-term prognosis of patients with CLD, including those with hepatocellular carcinoma, has been described. The effectiveness of the Controlling Nutritional Status (CONUT) score, which focuses on Alb, total lymphocyte count (TLC), and total cholesterol (TC) level, for systemic nutritional assessment, has been discussed by the nutrition support team (NST) and others (8). However, this tool is also cumbersome to use, requiring several items and the limit of CONUT score is leukocyte fraction, so it is affected by infectious diseases; thus, a single marker with a cut-off value for diagnosis is needed to examine the relationship with ALBI grade and CONUT score, and examine clinical application in hepatic reserve and nutritional status evaluation of patients with CLD.

In 2015, Mac-2-binding protein glycosylation isomer (M2BPGi), a carbohydrate antigen marker associated with fibrosis, was identified as a novel marker for liver fibrosis. M2BPGi is useful for the diagnosis of fibrosis in various liver

diseases and changes in liver fibrosis. It has also been reported as a predictor of carcinogenesis in chronic hepatitis C and sustained virologic response of chronic hepatitis C (9).

Although M2BPGi is a biomarker of liver fibrosis, it is considered to have a function other than fibrosis because it is increased even in patients at high acute liver injury and inflammatory disease (10,11).

However, the relationship between M2BPGi and nutritional indices and assessments of hepatic function reserve in patients with CLD has not been investigated in detail. Thus, in this study, we aimed to evaluate the use of M2BPGi as a biomarker in patients with CLD to examine clinical application. We present the following article in accordance with the STROBE reporting checklist (available at <https://apm.amegroups.com/article/view/10.21037/apm-22-270/rc>).

## Methods

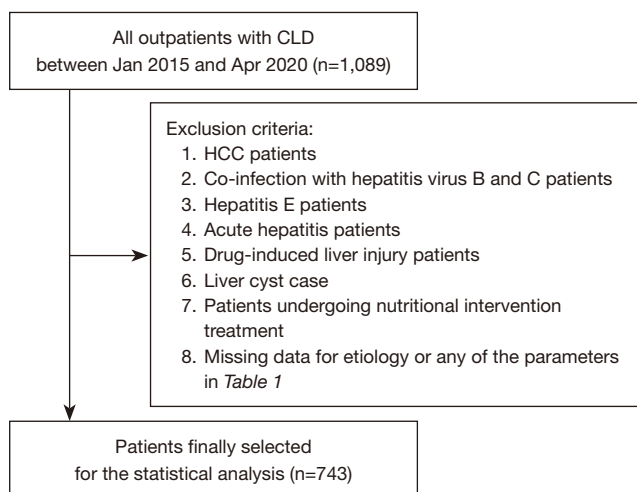
The relationship between the M2BPGi-CONUT score and M2BPGi-ALBI grade was investigated among 1,089 outpatients whose M2BPGi value was measured at their first visit to Saiseikai Niigata Hospital for CLD between January 2015 and April 2020. Seven hundred and forty-three patients whose CONUT score, ALBI grade, and M2BPGi were calculated from the same blood sample, and who did not meet any of the following exclusion criteria were included in the study (*Figure 1*). Patients with complications from hepatocellular carcinoma, those with missing data, and those in whom nutritional interventions such as branched-chain amino acids were already being used were excluded from the study.

ALBI grade was calculated using Alb and T. Bil as follows:

$$\begin{aligned} \text{ALBI score} = & 0.66 \times \log(\text{T. Bil, mmol/L}) \\ & - 0.085 \times (\text{Alb, g/L}) \end{aligned} \quad [1]$$

Patients with ALBI grade 1, 2 and 3 were allocated a score of 1, 2 and 3 points, respectively. Patients with mALBI grade 1, 2a, 2b and 3 were allocated a score of 1, 2 and 3 points, respectively.

The CONUT score was calculated using Alb, TLC and



**Figure 1** Flowchart of the patient process. CLD, chronic liver disease; HCC, hepatocellular carcinoma.

TC using the following formula:

$$\text{CONUT score} = \text{serum albumin score} + \text{TC score} + \text{TLC score} \quad [2]$$

where, serum albumin scores were 0:  $\geq 3.5$ ; 2: 3.0–3.49; 4: 2.50–2.99; 6:  $< 2.50$  g/dL; TC scores were 0:  $\geq 180$ ; 1: 140–179; 2: 100–139; 3:  $< 100$  mg/dL; and TLC scores were 0:  $\geq 1.6109$ ; 1: 1.20–1.59; 2: 0.80–1.19; 3:  $< 0.8$  g/L (8).

CONUT grade normal, light, moderate and severe were allocated of CONUT score.

FIB-4 was estimated using the following formula:

$$\text{FIB-4} = \frac{[\text{age (years)} \times \text{AST (U/L)}]}{[\text{PLT} (\times 10^9 / \text{L}) \times \text{ALT (U/L)}]^{0.5}} \quad [3]$$

M2BPGi value was evaluated using HISCL M2BPGi Assay Kit with an automated immunoassay system HISCL-5000 (Sysmex, Hyogo, Japan), which takes 17 min and requires 10  $\mu\text{L}$  sample of serum, allowing the results to be obtained on the date of blood sampling. M2BPGi levels were indexed using the following equation: cut-off index (COI) = (S-N)/(C-N), where S represents the light intensity in sample, N represents the light intensity in HISCL negative control and C represents the cut-off value.

The baseline characteristics of the 743 outpatients included in this study are shown in *Table 1*. The median age of patients was 67 (range, 59–75) years; 330 patients (44.41%) were males and 413 (55.59%) were females. CLD was related to hepatitis C virus (HCV) in 305 (41.0%)

patients, hepatitis B virus (HBV) in 120 (16.2%) patients, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis (NAFLD/NASH) in 167 (22.5%) patients, alcohol in 84 (11.3%) patients, and autoimmune hepatitis (AIH) in 67 (9.0%) patients. As the study was limited to patients with CLD under outpatient care, only six patients (0.81%) with ALBI grade 3 were included.

In this study, we determined the relationship between M2BPGi and nutritional status and hepatic function reserve in patients with CLD.

### Ethics statement

The study was approved by the Institutional Review Board of Saiseikai Niigata Hospital (No. E18-18) and was performed in accordance with the principles of the Declaration of Helsinki (as revised in 2013). All patients provided written informed consent.

### Statistical analysis

Data are expressed as median and interquartile range unless otherwise stated. Spearman's rank correlations were performed to evaluate the serum M2BPGi with ALBI grade and other clinical features. Results with  $0 \leq r < 0.3$  were considered almost irrelevant. Results with  $0.3 \leq r < 0.5$  were considered almost a very weak correlation. Results with  $0.5 \leq r < 0.7$  were considered almost a correlation. Results with  $0.7 \leq r < 0.9$  were considered almost a strong correlation. Pairwise comparisons were made using Wilcoxon's rank-sum test for between-group comparisons. To evaluate the diagnostic performance of M2BPGi in assessing hepatic function reserve and nutritional status, receiver operating characteristic (ROC) curve analysis was performed. Diagnostic accuracy is expressed as specificity, sensitivity, positive predictive value (PPV), negative ROC curve predictive value, and area under the ROC curve. The optimal cut-off values were obtained by maximizing Youden's index (sensitivity + specificity-1). Results with  $P < 0.05$  were considered statistically significant. All statistical analyses were performed using JMP15.2.0 (SAS Institute Inc., Cary, NC, USA).

### Results

The distribution of M2BPGi, CONUT, and ALBI by the underlying liver disease type is shown in *Figures 2-4*, respectively. The pairwise Wilcoxon test revealed significant

**Table 1** Clinical characteristics of 743 patients with CLD

Parameter	Total (n=743)	HCV (n=305)	HBV (n=120)	NAFLD/NASH (n=167)	Alcoholic (n=84)	Autoimmune (n=67)
Age (years)	67 [59–75]	72 [65–80]	61.5 [48–68]	65 [56–74]	65 [60–70]	69 [63–76]
Sex, n (%)						
Male	330 (44.41)	111 (36.39)	69 (57.50)	71 (42.51)	65 (77.38)	14 (20.90)
Female	413 (55.59)	194 (63.61)	51 (42.50)	96 (57.49)	19 (22.62)	53 (79.10)
AST (U/L)	25 [20–32]	23 [19–27]	22 [19.25–27]	30 [22–44]	29 [22–44.5]	26 [22–30]
ALT (U/L)	17 [12–26]	14 [11–20]	16.5 [13–23]	29 [17–49]	22 [13.25–33]	17 [13–22]
Serum albumin (g/dL)	4.2 (3.9–4.4)	4.2 (3.9–4.4)	4.2 (4–4.38)	4.2 (4–4.4)	4.1 (3.6–4.4)	4.1 (3.8–4.3)
Total bilirubin (mg/dL)	0.59 (0.45–0.81)	0.57 (0.43–0.76)	0.74 (0.56–0.99)	0.59 (0.46–0.79)	0.57 (0.46–0.86)	0.56 (0.42–0.67)
Prothrombin time (%)	106.8 (96.8–115.7)	107.7 (97.3–117.5)	103.65 (92.85–114.88)	109.9 (100.6–116.7)	103.2 (89.65–110.18)	105.8 [96–115]
Platelet count ( $\times 10^4$ /mL)	19.3 (15.5–23.6)	18.2 (15–22.25)	19.85 (14.95–22.78)	20.5 (17–24.4)	18.65 (14.03–22.83)	23.2 (19–30.5)
CONUT score	1 [0–2]	1 [0–2]	2 [0–2]	1 [0–2]	1 [0–2]	2 [1–3]
TC (mg/dL)	185 [161–207]	188 [165.5–212.5]	185 [164–199]	177 [157–197]	182 [161–202]	189 [168–211]
Lymphocyte count (%)	15.8 (11.9–21.3)	15.8 (12–21.7)	13.65 (10.32–18.1)	17.3 (14.7–23.3)	15.95 (11.93–20.90)	13.6 (9.9–19.1)
M2BPGi (COI)	0.82 (0.57–1.32)	0.95 (0.67–1.55)	0.66 (0.47–1.03)	0.69 (0.48–1.1)	0.91 (0.68–1.61)	0.93 (0.57–1.35)
Fib-4 index	2.17 (1.43–3.09)	2.4 (1.75–3.27)	1.84 (1.03–2.84)	1.88 (1.19–2.71)	2.37 (1.72–3.77)	1.8 (1.21–2.8)
ALBI score	–2.9 (–3.1, –2.7)	–2.9 (–3.1, –2.7)	–2.8 (–3.01, –2.6)	–2.9 (–3.1, –2.7)	–2.8 (–3.08, –2.45)	–2.9 (–3.1, –2.6)
mALBI grade, n (%)						
1	595 (80.08)	251 (82.30)	97 (80.83)	142 (85.03)	55 (65.48)	50 (74.63)
2a	90 (12.11)	35 (11.48)	14 (11.67)	14 (8.38)	16 (19.05)	11 (16.42)
2b	52 (7.00)	19 (6.23)	6 (5.00)	10 (5.99)	12 (14.29)	5 (7.46)
3	6 (0.81)	0	3 (2.50)	1 (0.60)	1 (1.19)	1 (1.49)

Data are presented as n (%), or median (interquartile range). CLD, chronic liver disease; HCV, hepatitis C virus; HBV, hepatitis B virus; NAFLD/NASH, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis; AST, aspartate aminotransferase; ALT, alanine aminotransferase; CONUT, Controlling Nutritional Status; TC, total cholesterol; M2BPGi, mac-2-binding protein glycosylation isomer; COI, cut-off index; ALBI, albumin-bilirubin; mALBI, modified albumin-bilirubin.

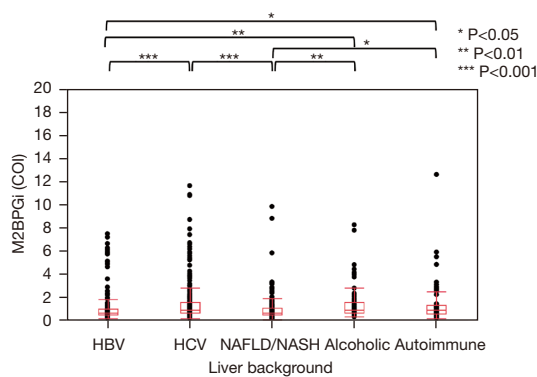
differences among some groups.

The correlation coefficient ( $r^2$ ) between M2BPGi and ALBI grade was 0.40 ( $r=0.63$ ), indicating a positive correlation between M2BPGi and ALBI grade, where M2BPGi increased with ALBI and mALBI grade (Figure 5). The  $r^2$  between FIB-4 index and CONUT score was 0.17 ( $r=0.41$ ), indicating a very weak positive correlation between the FIB-4 index and CONUT score.

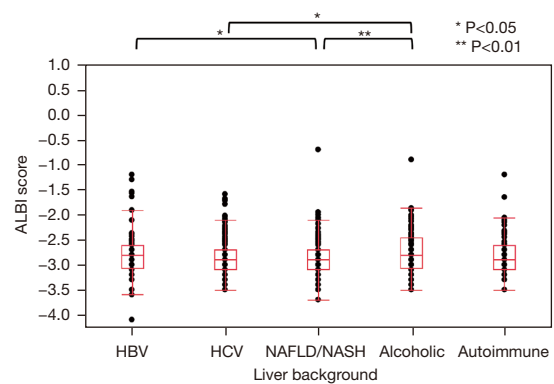
The  $r^2$  between the FIB-4 index and ALBI grade was 0.22 ( $r=0.47$ ), indicating a very weak positive correlation between FIB-4 index and ALBI score.

#### All cases (n=743) (Table 2)

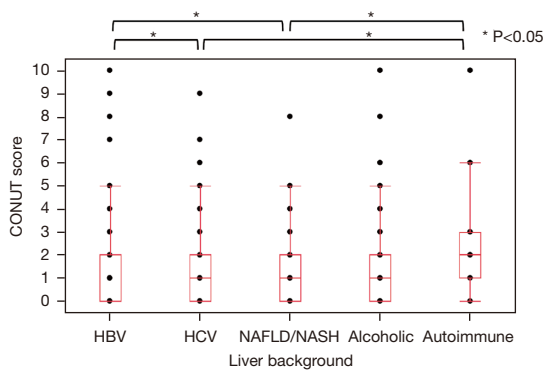
M2BPGi predicted mALBI G1 vs. G2–G3 with a sensitivity of 0.75 and specificity of 0.76 when area under the curve



**Figure 2** Background M2BPGi distribution by the type of liver disease. \*,  $P<0.05$ ; \*\*,  $P<0.01$ ; \*\*\*,  $P<0.001$ . HBV, hepatitis B virus; HCV, hepatitis C virus; NAFLD/NASH, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis; M2BPGi, Mac-2-binding protein glycosylation isomer; COI, cut-off index.



**Figure 4** Background ALBI grade distribution by the type of liver disease. \*,  $P<0.05$ ; \*\*,  $P<0.01$ . HBV, hepatitis B virus; HCV, hepatitis C virus; NAFLD/NASH, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis; ALBI, albumin-bilirubin.

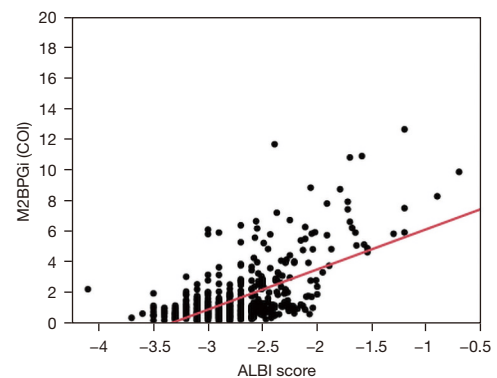


**Figure 3** Background CONUT score distribution by the type of liver disease. \*,  $P<0.05$ . HBV, hepatitis B virus; HCV, hepatitis C virus; NAFLD/NASH, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis; CONUT, Controlling Nutritional Status.

(AUC) =0.82, cut-off 1.07. M2BPGi predicted mALBI G1–2a *vs.* G2b–G3 with a sensitivity of 0.78 and specificity of 0.86 when AUC =0.89, cut-off 1.73. M2BPGi predicted mALBI G1–2 *vs.* G3 with a sensitivity of 1.00 and specificity of 0.97 when AUC =0.99, cut-off 5.83.

#### HCV (n=305)

There were no G3 cases among patients with HCV. M2BPGi predicted mALBI G1–G2a *vs.* G2b with a sensitivity of 0.79 and specificity of 0.82 when AUC =0.86, cut-off 1.78.



**Figure 5** Correlation between the M2BPGi value and ALBI grade. ALBI, albumin-bilirubin; M2BPGi, Mac-2-binding protein glycosylation isomer; COI, cut-off index.

#### HBV (n=120)

M2BPGi predicted mALBI G1–2a *vs.* G2b–G3 with a sensitivity of 1.00 and specificity of 0.96 when AUC =0.97, cut-off 4.63. M2BPGi predicted mALBI G1–G2 *vs.* G3 with a sensitivity of 1.00 and specificity of 0.97 when AUC =0.98, cut-off 5.83.

#### NAFLD/NASH (n=167)

M2BPGi predicted mALBI G1 *vs.* G2–G3 with a sensitivity of 0.92 and specificity of 0.56 when AUC =0.82, cut-off 0.68. M2BPGi predicted mALBI G1–G2a *vs.* G2b–G3 with a sensitivity of 0.82 and specificity of 0.92 when AUC =0.92,

**Table 2** Diagnostic performance of M2BPGi values in distinguishing different mALBI grade

Categories	Cut-off value	AUC	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)	TP	TN	FP	FN
Total (n=743)										
G1 vs. G2–G3	1.07	0.82	75.00	75.80	43.53	92.42	111	451	144	37
G1–G2a vs. G2b–G3	1.73	0.89	77.59	86.42	32.61	97.85	45	592	93	13
G1–G2 vs. G3	5.83	0.99	100.00	97.15	22.22	100.00	6	716	21	0
HCV (n=305)										
G1 vs. G2–G3	1.93	0.82	68.52	90.84	61.67	93.06	37	228	23	17
G1–G2a vs. G2b–G3	1.78	0.86	78.95	82.17	22.73	98.33	15	235	51	4
HBV (n=120)										
G1 vs. G2–G3	1.30	0.83	73.91	93.81	73.91	93.81	17	91	6	6
G1–G2a vs. G2b–G3	4.63	0.97	100.00	96.40	69.23	100.00	9	107	4	0
G1–G2 vs. G3	5.83	0.98	100.00	96.58	42.86	100.00	3	113	4	0
NAFLD/NASH (n=167)										
G1 vs. G2–G3	0.68	0.82	92.00	55.63	26.74	97.53	23	79	63	2
G1–G2a vs. G2b–G3	1.49	0.92	81.82	92.31	42.86	98.63	9	144	12	2
G1–G2 vs. G3	9.86	1.00	100.00	100.00	100.00	100.00	1	166	0	0
Alcoholic (n=84)										
G1 vs. G2–G3	0.88	0.82	89.66	65.45	57.78	92.31	26	36	19	3
G1–G2a vs. G2b–G3	2.80	0.88	69.23	98.59	90.00	94.59	9	70	1	4
G1–G2 vs. G3	8.27	1.00	100.00	100.00	100.00	100.00	1	83	0	0
Autoimmune (n=67)										
G1 vs. G2–G3	1.08	0.87	88.24	78.00	57.69	95.12	15	39	11	2
G1–G2a vs. G2b–G3	1.22	0.83	83.33	75.41	25.00	97.87	5	46	15	1
G1–G2 vs. G3	12.63	1.00	100.00	100.00	100.00	100.00	1	66	0	0

M2BPGi, mac-2-binding protein glycosylation isomer; mALBI, modified albumin-bilirubin; AUC, area under the curve; PPV, positive predictive value; NPV, negative predictive value; TP, true positive; TN, true negative; FP, false positive; FN, false negative; G1, grade 1; G2, grade 2; G3, grade 3; HCV, hepatitis C virus; HBV, hepatitis B virus; NAFLD/NASH, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis.

cut-off 1.49. M2BPGi predicted mALBI G1–G2 vs. G3 with a sensitivity of 1.00 and specificity of 1.00 when AUC =1.00, cut-off 9.86.

#### Alcoholic liver disease (n=84)

M2BPGi predicted mALBI G1–G2 vs. G3, with a sensitivity of 1.00 and specificity 1.00 when AUC =1.00, cut-off 8.27.

#### AIH (n=67)

M2BPGi predicted mALBI G1–G2 vs. G3 with a sensitivity

and specificity of 1.00 when AUC =1.00, cut-off 12.63.

Next, we analyzed the relationship between M2BPGi and CONUT scores (*Table 3*). M2BPGi predicted CONUT score normal vs. light-severe with a sensitivity of 0.38 and specificity of 0.92 when AUC =0.65, cut-off 1.60. M2BPGi predicted CONUT score normal-light vs. moderate-severe with a sensitivity of 0.91 and specificity of 0.85 when AUC =0.93, cut-off 1.74. M2BPGi predicted CONUT score normal-moderate vs. severe with a sensitivity of 1.00 and specificity of 0.97 when AUC =0.99, cut-off 5.83.



**Table 3** Diagnostic performance of M2BPGi values in distinguishing different CONUT score

Categories	Cut-off value	AUC	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)	TP	TN	FP	FN
Total (n=743)										
Grade normal vs. light-severe	1.60	0.65	38.41	92.07	75.51	70.13	111	418	36	178
Grade normal-light vs. moderate-severe	1.74	0.93	90.63	84.81	21.17	99.50	29	603	108	3
Grade normal-moderate vs. severe	5.83	0.99	100.00	97.15	22.22	100.00	6	716	21	0
HCV (n=305)										
Grade normal vs. light-severe	1.60	0.71	49.06	88.44	69.33	76.52	52	176	23	54
Grade normal-light vs. moderate-severe	5.40	0.87	69.23	97.60	56.25	98.62	9	285	7	4
Grade normal-moderate vs. severe	10.89	1.00	100.00	99.67	50.00	100.00	1	303	1	0
HBV (n=120)										
Grade normal vs. light-severe	1.30	0.66	36.07	98.31	95.65	59.79	22	58	1	39
Grade normal-light vs. moderate-severe	4.63	0.96	100.00	95.54	61.54	100.00	8	107	5	0
Grade normal-moderate vs. severe	5.83	0.98	100.00	96.58	42.86	100.00	3	113	4	0
NAFLD/NASH (n=167)										
Grade normal vs. light-severe	1.73	0.60	26.32	98.18	88.24	72.00	15	108	2	42
Grade normal-light vs. moderate-severe	1.74	0.95	100.00	92.07	18.75	100.00	3	151	13	0
Alcoholic (n=84)										
Grade normal vs. light-severe	2.79	0.60	31.25	98.08	90.91	69.86	10	51	1	22
Grade normal-light vs. moderate-severe	1.38	0.92	100.00	75.95	20.83	100.00	5	60	19	0
CONUT grade normal-moderate vs. severe	8.27	1.00	100.00	100.00	100.00	100.00	1	83	0	0
Autoimmune (n=67)										
Grade normal vs. light-severe	1.08	0.69	57.58	79.41	73.08	65.85	19	27	7	14
Grade normal-light vs. moderate-severe	2.78	0.97	100.00	90.63	33.33	100.00	3	58	6	0
Grade normal-moderate vs. severe	12.63	1.00	100.00	100.00	100.00	100.00	1	66	0	0

M2BPGi, mac-2-binding protein glycosylation isomer; CONUT, Controlling Nutritional Status; AUC, area under the curve; PPV, positive predictive value; NPV, negative predictive value; TP, true positive; TN, true negative; FP, false positive; FN, false negative; HCV, hepatitis C virus; HBV, hepatitis B virus; NAFLD/NASH, nonalcoholic fatty liver disease/non-alcoholic steatohepatitis.

### **HCV (n=305) (CONUT score)**

M2BPGi predicted CONUT score normal-moderate *vs.* severe with a sensitivity and specificity of 1.00 when AUC =1.00, cut-off 10.89.

### **HBV (n=120) (CONUT score)**

M2BPGi predicted CONUT score normal-light *vs.* moderate-severe with a sensitivity of 1.00 and specificity of 0.96 when AUC =0.96, cut-off 4.63.

M2BPGi predicted CONUT score normal-moderate *vs.*

severe with a sensitivity of 1.00 and specificity of 0.97 when AUC =0.98, cut-off 5.83.

None of the patients with NADH/NAFLD were evaluated as CONUT severe. M2BPGi predicted CONUT score normal-light *vs.* moderate with a sensitivity of 1.00 and specificity of 0.92 when AUC =0.95, cut-off 1.74.

### **Alcoholic liver disease (n=84) (CONUT score)**

M2BPGi predicted CONUT score normal-light *vs.* moderate-severe with a sensitivity of 1.00 and specificity of

0.76 when AUC =0.92, cut-off 1.38.

### *AIH (n=67) (CONUT score)*

M2BPGi predicted CONUT score normal-light *vs.* moderate-severe with a sensitivity of 1.00 and specificity of 0.91 when AUC =0.97, cut-off 2.78.

## Discussion

Malnutrition is common in CLD, and early intervention is necessary to improve patient prognosis (12,13). Serum albumin is the main factor used to evaluate CLD and is included in both ALBI and Child-Pugh scores. The progression of CLD, including viral and non-viral diseases, involves cirrhosis and hepatocellular carcinoma as a consequence of liver fibrosis. Recently, HISCL M2BPGi, which allows the evaluation of progression of liver fibrosis via glycan markers in serum, has been developed; it is commercially available (14).

Various studies have evaluated the clinical relationship between M2BPGi and CLD pathogenesis (15-18).

Although M2BPGi has the potential for use as a biomarker during treatment and could be used to evaluate prognosis in clinical practice, clinical data are insufficient. In this study, we aimed to clarify the clinical usefulness of measuring the hepatic fibrosis marker M2BPGi at the first outpatient visit in individuals with CLD and to determine whether it would be suitable as a biomarker for nutritional status by ALBI and CONUT, which are indicators of hepatic function reserve.

M2BPGi, *Wisteria floribunda* agglutinin-positive Mac 2-binding protein (WFA<sup>+</sup>-M2BP), is a carbohydrate structure on M2BP that increases in concentration with the progression of liver disease. M2BPGi is a donut-shaped multimeric structure containing 100 glycans, among which, the changes in N-glycan during fibrosis can be identified using lectin WFA. M2BPGi is superior to existing biomarkers in distinguishing cases of advanced fibrosis (F3/F4; 3 COI or more indicates cirrhosis) and increased cirrhosis (up to 20 COI can be measured), and it markedly decreases following the successful treatment of HCV with interferon-based therapies (9,19).

Conversely, the CONUT score is a nutritional index that considers immune status and may be used as an index to evaluate various conditions, including the NST. To date, only a few studies have investigated the relationship between M2BPGi and ALBI grade or CONUT score in

liver disease (20,21).

This is because the cut-off values for M2BPGi differ with the underlying cause of liver disease.

In this study, we examined M2BPGi cut-off values for each liver disease type. We found that M2BPGi was not a suitable biomarker for distinguishing between CONUT normal and light or low in outpatients; however, cut-offs by disease type were effective for other patients. Overall, M2BPGi presented the highest AUC in mALBI G1-G2a and G2b-G3 (AUC =0.88), and CONUT normal-light-moderate and severe (AUC =0.96) groups. Although distinguishing between CONUT normal and light-moderate is challenging, the finding that CONUT moderate and severe can be distinguished by a single marker suggests that active nutritional interventions can be expected at the outpatient group. In addition, the trend did not change significantly when the analysis was performed by primary disease. M2BPGi has also been reported to correlate with indices of the hepatic reserve, and it is speculated that Alb may contribute to this correlation. ALBI grade has also been reported to reflect nutritional status in patients with hepatocellular carcinoma (22).

Although these indices are important for the treatment of patients with hepatocellular carcinoma, shifts in M2BPGi have been reported in patients with hepatocellular carcinoma; thus, we excluded patients with hepatocellular carcinoma from this study. However, it is important to separate M2BPGi grade to decide the timing of treatment in clinical practice. The results of the present study demonstrated that M2BPGi can reflect ALBI grade and metabolic parameters, and thus, nutritional status as CONUT, although the cut-off varied with disease type.

This study had some limitations. First, data were obtained from a single institution. Second, the patients did not have a histological diagnosis of liver disease based on liver biopsy. Third, no indirect calorimetry assay was performed. Finally, all participants were outpatients and a few patients with Child C and ALBI 3 were included.

In the future, it will be necessary to validate these findings using data from a large number of patients obtained at multiple institutions, including hospitalized patients and patients with advanced CLD. In addition, it will be necessary to examine the use of this biomarker to predict carcinogenesis and to determine its relatedness to sarcopenia.

In conclusion, we found that M2BPGi is a single indicator of CLD status, including both ALBI grade and CONUT score.



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