

Mac-2 binding protein glycan isomer (M2BPGi) is a new serum biomarker for assessing liver fibrosis: more than a biomarker of liver fibrosis

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Abstract Assessing liver fibrosis is important for predicting the efficacy of antiviral therapy and patient prognosis. Liver biopsy is the gold standard for diagnosing liver fibrosis, despite its invasiveness and problematic diagnostic accuracy. Although noninvasive techniques to assess liver fibrosis are becoming important, reliable serum surrogate markers are not available. A glycoproteomics study aimed at identifying such markers discovered Mac 2-Binding Protein Glycan Isomer (M2BPGi), which is a reliable marker for assessing liver fibrosis in patients with viral hepatitis and other fibrotic liver diseases such as primary biliary cholangitis, biliary atresia, autoimmune hepatitis, and nonalcoholic fatty liver disease. M2BPGi predicts the development of hepatocellular carcinoma (HCC) in patients infected with hepatitis B and C as well as the prognosis of liver cirrhosis in those with HCC after therapy. The unique features of M2BPGi are as follows: (1) cut-off values differ for the same stages of fibrosis according to the cause of fibrosis; and (2) M2BPGi levels rapidly decrease after patients achieve a sustained antiviral

response to hepatitis C virus. These observations cannot be explained if M2BPGi levels reflect the amount of fibrotic tissue. Hepatic stellate cells (HSCs) secrete M2BPGi, which may serve as a messenger between HSCs and Kupffer cells via Mac-2 (galectin 3) that is expressed in Kupffer cells during fibrosis progression. Here we show that M2BPGi is a surrogate marker for assessing HSC activation. These findings may reveal the roles of HSCs in extrahepatic fibrotic disease progression.

Keywords Liver fibrosis · M2BPGi · Hepatocarcinogenesis

Introduction

Chronic liver disease (CLD) is a serious health concern worldwide, and infections associated with hepatitis B virus (HBV) and hepatitis C virus (HCV), alcohol abuse, primary biliary cholangitis, and nonalcoholic fatty liver disease (NAFLD) are the common predisposing conditions for developing liver fibrosis and cirrhosis [1]. Without proper management, CLD progresses to liver fibrosis and consequently leads to liver cirrhosis, which increases morbidity and mortality caused by portal hypertension, hepatic insufficiency, and the development of hepatocellular carcinoma (HCC) [2, 3]. The prognosis and treatment of patients with CLD varies depending on the stage of fibrosis, and therefore the diagnosis of the stage of fibrosis is clinically important [4–6]. However, the diagnosis of liver cirrhosis is challenging [1].

Although liver biopsy is considered the gold standard for stratifying hepatic fibrosis [7, 8], it is invasive, its diagnostic value is limited by sampling error, and histologic interpretations can vary [1, 9–12]. Furthermore,

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repeated liver biopsies to monitor liver fibrosis are not feasible in a routine clinical setting [1, 13]. Therefore, noninvasive tests to assess liver fibrosis have emerged, which include elastographic techniques [14–17], serum biomarkers such as hyaluronic acids, type IV collagen, and type III procollagen-N-peptide [18] as well as surrogate markers (e.g., the AST-to-platelet ratio index (APRI) [19], and the FIB-4 index [20]). Among elastographic techniques, transient elastography (TE) and magnetic resonance elastography (MRE) are widely accepted. Several major manufacturers of ultrasound instrumentation recently have developed ultrasound systems that perform elastographic imaging. Furthermore, a novel marker for assessing liver fibrosis, M2BPGi (called WFA⁺-M2BP), was recently introduced. In this review, we summarize the biological profiles and clinical applications of M2BPGi.

Mac-2 binding protein

The 90K/Mac-2 binding protein (M2BP) is a secreted glycoprotein that is present in the extracellular matrix of several tissues [21], and elevated levels are present in certain tumors and in people with viral infections [21]. In vitro, human M2BP induces the production of interleukin (IL)-1, IL-6, and other cytokines by blood monocytes [22]. A prominent feature of human M2BP is its oligomerization to form large ring structures, which resemble a “powdered sugar doughnut” covered with 70–112 *N*-glycans [23]. M2BP is extensively glycosylated and interacts with galectin-3 (former name, Mac-2), and it interacts with other extracellular proteins such as collagens IV, V, and VI, fibronectin, and nidogen [23]. M2BP binds galectin-3 on the cell surface and induces homotypic cell aggregation [23].

Sources and the role of M2BPGi in liver fibrosis

Bekki et al. [24] found that hepatic stellate cells (HSCs) are the source of M2BPGi in subpopulations of liver-derived cells such as HSCs, Kupffer cells, endothelial cells, biliary epithelial cells, and hepatocytes. An in vitro study found that the addition of exogenous M2BPGi enhances Mac-2 (galectin 3) expression by Kupffer cells. Furthermore, in cocultures of HSCs and Kupffer cells, alpha-SMA expression by HSCs is increased, which is reduced by Mac-2 depletion from Kupffer cells. These findings suggest that M2BPGi is a juxtacrine-acting messenger sent by HSCs to Kupffer cells during liver fibrosis and that it plays an important role in the progression of fibrosis (Fig. 1). Thus, M2BPGi levels reflect the activation of HSCs during the progression of liver fibrosis, in contrast to the amount of

collagen. This may explain the rapid decrease of M2BPGi levels after patients with hepatitis C achieve a sustained virus response (SVR).

An immunohistochemical analysis of cirrhotic human liver found that Mac-2 (galectin 3) and M2BPGi are expressed by CD68-positive cells, which are likely Kupffer cells. Therefore, M2BPGi may interact with Mac-2-positive cells to induce biological activity. Mac-2 is involved in diverse functions such as cell adhesion, growth regulation, cytokine production, T cell apoptosis, and immune responses. For example, Kianoush et al. [25] suggest that Mac-2 induces M2 polarization of macrophages, and other studies indicate that Mac-2 may stimulate cancer progression [26]. Thus, the biological activities of M2BPGi mediated by Mac-2 may explain the high incidence of the development of HCC development in patients with high M2BPGi levels.

Discovery of a new marker for diagnosing liver fibrosis

Kuno et al. [27, 28] and Narimatsu [29] used glycoproteomics techniques to identify biomarkers of liver fibrosis. Glycoproteomics uses comprehensive high-throughput methods to analyze glycosylation sites, glycan structures, and peptide sequences. The investigators cited above focused on the glyco-alteration of α 1-acid glycoprotein (AGP), an acute-phase protein that is secreted mainly by the liver. Glyco-alteration of AGP occurs during cirrhosis and acute and chronic inflammation. Thus, the signal patterns of 12 selected lectins reflect fibrosis-associated glyco-alterations of AGP [27]. Furthermore, a fully automated glycan-based immunoassay, called FastLec-Hepa, was developed to search for serum biomarkers for liver fibrosis using a single glycoprotein rather than sets of glycoproteins [28]. FastLec-Hepa automatically detects unique fibrosis-related glyco-alterations of hyperglycosylated Mac-2 binding protein present in serum. Six candidate lectins were selected that bind M2BP [28]. After analyzing sera from patients with different stages of fibrosis, Wisteria floribunda agglutinin (WFA) was found to be superior to the other five lectins for diagnosing liver fibrosis. This glycoprotein (WFA⁺-Mac2 binding protein) is named Mac-2 binding protein glycosylation isomer (M2BPGi).

Measurement of M2BPGi

M2BPGi is measured using a sandwich immunoassay with anti-WFA and anti-M2BP antibodies. M2BP enriched from serum is added to WFA-coated agarose in a microtube, immunoprecipitated, and analyzed using 5–20% gradient SDS-polyacrylamide electrophoresis under reducing

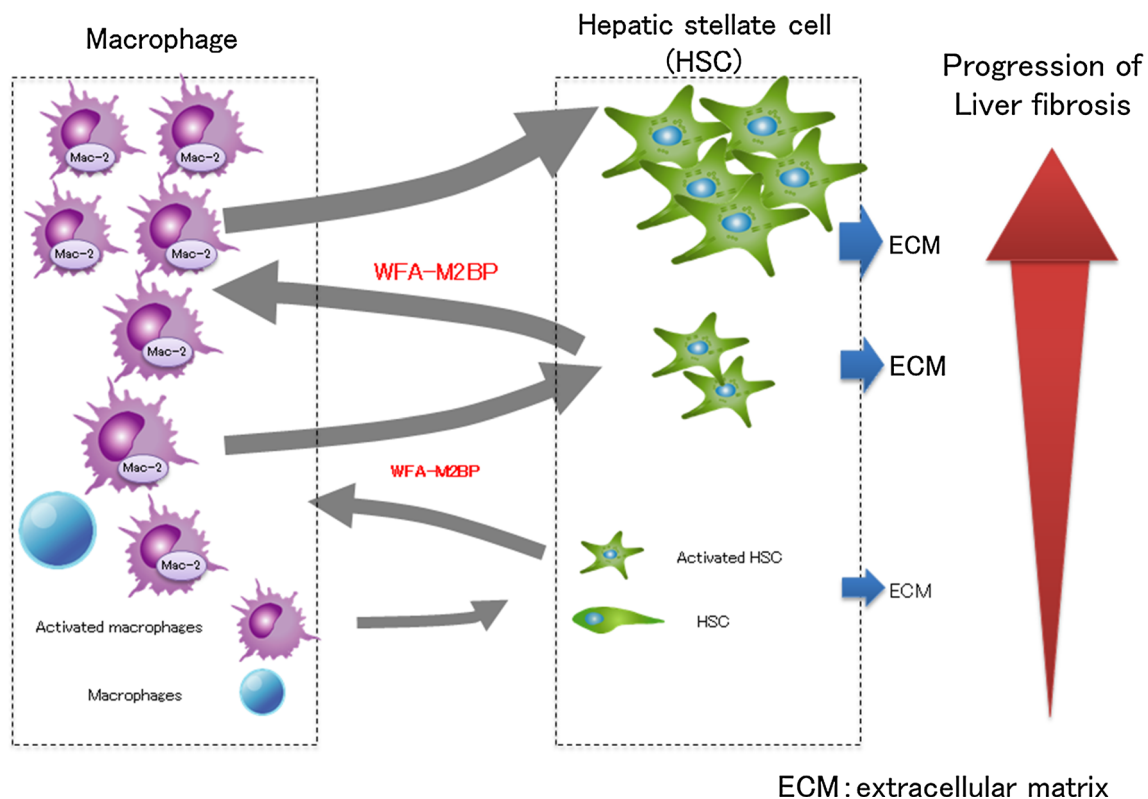


Fig. 1 The role of M2BPGi in the progression of liver cirrhosis

conditions [28]. Alternatively, M2BPGi is measured using a fully automated HSCL-2000i Immunoanalyzer (Sysmecs Co., Hyogo, Japan) [29]. The values of M2BPGi conjugated to WFA are indexed to the values calculated as follows:

$$\text{Cut-off index (COI)} = \frac{([\text{M2BPGi}]_{\text{sample}} - [\text{M2BPGi}]_{\text{NC}})}{([\text{M2BPGi}]_{\text{PC}} - [\text{M2BPGi}]_{\text{NC}})},$$

where $[\text{M2BPGi}]_{\text{sample}}$ represents the M2BPGi count of the serum sample (PC, positive control; NC, negative control). The positive control is supplied as a calibration solution that is preliminarily standardized to yield $\text{COI} = 1.0$.

Clinical significance of M2BPGi for the assessment of liver fibrosis

Kuno et al. [28] found that M2BPGi is the most precise predictor of severe fibrosis or liver cirrhosis compared with markers such as the FIB-4 index and hyaluronic acid. Moreover, Toshima et al. [30] found the following: 1. Only histologically diagnosed fibrosis stage correlates with M2BPGi levels. 2. The area under the curve (AUC) of fibrosis ($F \geq 3$) using M2BPGi is similar to that determined using ultrasound-based virtual touch tissue quantification. 3. The AUC values of M2BPGi are superior to those of serum

surrogate markers such as APRI, hyaluronic acids, and type 4 collagen. These findings suggest that M2BPGi may serve as the most reliable serum biomarker. Moreover, the usefulness of M2BPGi as a marker for assessing liver fibrosis is shown in a study that enrolled patients infected with chronic hepatitis C [28, 31, 32] or B virus [33–37]. Recent studies show that M2BPGi is a useful marker for monitoring the improvement of patients with liver fibrosis after achieving an SVR following antiviral therapy [38, 39].

M2BPGi is useful for assessing liver fibrosis in patients with primary biliary cirrhosis [40, 41], biliary atresia [42], autoimmune hepatitis [43], nonalcoholic fatty liver disease [44–48], and hepatitis B or C infection. A meta-analysis confirms that M2BPGi is a reliable predictor for staging liver fibrosis [49]. Although M2BPGi is a novel marker for assessing liver fibrosis, M2BPGi COI values may differ among patients stratified according to the cause of liver fibrosis, even for those with the same stage of fibrosis (Table 1). Nishikawa et al. [50] found that M2BPGi levels in patients with chronic hepatitis C are higher compared with those of patients with chronic hepatitis B infection, even for those with the same degree of liver fibrosis. Patients with nonalcoholic steatohepatitis (NASH) have the lowest M2BPGi levels compared with those of patients with chronic hepatitis C or B infection. The histological criteria used to diagnose liver fibrosis in patients with NASH differ

Table 1 Differences in M2BPGi COI values between stages of fibrosis among patients with liver disease

Fibrosis stages	Liver diseases				
	HCV* [31]	HBV** [35]	NAFLD* [46]	AIH** [43]	PBC** [40]
F0	1.3 ± 0.1 (n = 274)	0.9 (n = 108)	0.46 (n = 71)		0.4 (n = 10)
F1			0.57 (n = 90)	1.5 (n = 18)	0.6 (n = 71)
F2	2.2 ± 0.1 (n = 37)	1.4 (n = 37)	0.64 (n = 16)	2.1 (n = 24)	1.4 (n = 27)
F3	3.3 ± 0.2 (n = 120)	1.6 (n = 28)	0.61 (n = 36)	3.3 (n = 24)	2.0 (n = 19)
F4	5.2 ± 0.3 (n = 120)	3.1 (n = 16)	0.91 (n = 7)	9.8 (n = 18)	3.0 (n = 11)

Stages of fibrosis were defined using the criteria of the French METAVIR Cooperative Study Group [76] and the scoring system for nonalcoholic steatohepatitis of the Clinical Research Network [51]

HCV hepatitis C virus, HBV hepatitis B virus, NAFLD nonalcoholic fatty liver diseases, AIH autoimmune hepatitis, PBC Primary biliary cholangitis

*The data are expressed as the mean (± standard error)

**The data are expressed median values

from those with chronic hepatitis B or C infection. However, in patients with advanced cirrhosis with NASH [51], the M2BPGi levels are much lower compared with those of patients with chronic hepatitis B or C infection.

Another unique aspect of M2BPGi is its rapid decrease after treatment of hepatitis C. For example, Nagata et al. [52] found that the M2BPGi values of 64 patients who achieved SVR significantly decreased after completing treatment. However, in 12 patients without SVR, there were no significant differences between serum M2BPGi values before and after treatment. Because rapid improvement of liver fibrosis beyond 24-week post-treatment is unlikely, even in SVR patients, these studies suggest that M2BPGi values may not reflect the amount of fibrous tissue.

M2BPGi reflects liver function, the effects of antiviral therapy, and liver failure

Yamasaki et al. [31] studied patients with chronic hepatitis C and found that M2BPGi correlates with the stage of fibrosis, alpha-fetoprotein (AFP), albumin, AST, platelets, sex, HCV core antigen, total bilirubin, and age. Other studies show that M2BPGi levels significantly correlate with the Child–Pugh

class and the MELD score, and serve as a marker of fibrosis [53, 54]. Moreover, M2BPGi is a useful marker for predicting the efficacy of direct-acting agent-based therapy, including regimens with or without interferon for treating chronic hepatitis C infection [55, 56]. In patients with hepatitis B infection, M2BPGi levels that are > 1.55 COI are poor significant predictors that are linked to the loss of HBeAg or seroconversion of patients positive for hepatitis B e-antigen [57]. Furthermore, Hanai et al. [53] and Hasegawa et al. [58] found that M2BPGi is a useful predictor of the prognosis of patients with liver cirrhosis (Table 2).

Okuda et al. [59] found that preoperative M2BPGi predicts posthepatectomy liver failure, and Morio et al. [60] found that M2BPGi increases in patients with acute liver injury and decreased after recovery. They suggested that M2BPGi might reflect liver fibrosis and other factors such as liver inflammation, liver damage, and hepatocyte regeneration.

Predicting the development of HCC

Several studies show that elevated serum M2BPGi levels predict the development of HCC (Table 3). For example, Yamasaki et al. [31] found that in patients with chronic

Table 2 The prognostic value of M2BPGi for patients with liver cirrhosis and hepatocellular carcinoma

Authors	Diseases	Therapy	Cut-off values (COI)
Hanai et al. [53]	LC	No	5.0
Hasegawa et al. [58]	Compensated LC (HCV)	No	6.15
Toyoda et al. [71]	Early HCC	Hepatectomy	3.00
Fujiyoshi et al. [70]	HCC	Hepatectomy	1.435 (non-HCV), 4.615 (HCV)

LC liver cirrhosis, HCV hepatitis C virus, HCC hepatocellular carcinoma

Table 3 M2BPGi as a risk factor for the development of HCC

Authors	Diseases	Cut off values (COI)	HR (95% CI)
Yamasaki et al. [31]	HCV	< 1	1
		1–4	5.155 (1.18–22.5)
		≥ 4	8.318 (1.78–38.79)
Tamaki et al. [61]	HCV	< 4.2	1
		≥ 4.2	4.2 (2.6–26)
		ΔM2BPGi/year < 0.3	1
		ΔM2BPGi/year < 0.3	3.1 (1.1–9.3)
Nagata et al. [52]	HCV	< 2.2	Not available
		After viral Tx	≥ 2.2
Sato et al. [62]	HCV postSVR	< 2.8	1
		≥ 2.8	15.21 (1.77–130.9)
Sasaki et al. [63]	HCV postSVR	≤ 2.0	1
		> 2.0	7.30 (2.20–24.17)
Heo et al. [64]	HBV	< 1.8	1
Kim SU, et al. [65].	HBV	> 1.8	11.5 (1.4–97.2)
		< 1.8	1
Ichikawa Y, et al. [66].	HBV	≥ 1.8	1.539(1.117-2.120) < 0.71
		< 0.71	1
Cheung KS, et al. [67]	HBV	≥ 0.71	8.32 (1.03-67.0)
		< 0.69	1
		≥ 0.69	4.80 (1.83-12.58)

ΔM2BPGi/year is defined as follows: (M2BPGi at follow up—M2BPGi at liver biopsy)/interval between two measurements (years)

HCV hepatitis C virus, HBV hepatitis B virus, HR hazard ratio, CI confidence interval, Tx treatment, SVR sustained viral response

hepatitis C infection with M2BPGi ≥ 4, the cumulative 5-year incidence of HCC is 77%, although for those with COI = 1–4 or those with COI < 1, the cumulative 5-year incidence is 31.6% and 3.1%, respectively.

There is a high incidence of HCC in patients with chronic hepatitis C infection and elevated serum M2BPGi levels [61]. In patients with chronic hepatitis C infection who reach an SVR and who receive antiviral therapy, a high M2BP value is a significant risk factor for the development of HCC [28, 31, 62, 63]. Furthermore, there is a high incidence of HCC in patients with chronic hepatitis B infection and elevated serum M2BPGi levels [64–67]. A

meta-analysis supports the hypothesis that an elevated M2BPGi level is a risk factor for developing HCC [49].

Elevated AFP levels correlate with a high incidence of the development of HCC [68]. However, Yamasaki et al. [31] found that the M2BPGi assay is superior to other assays that assess AFP levels for predicting the development of HCC. Furthermore, the progression of liver fibrosis increases the risk of developing HCC [69]. Therefore, predicting the development of HCC using M2BPGi may reflect liver fibrosis. Yamasaki et al. [31] found that elevated M2BPGi levels are a significant risk factor, even after stratification of patients according to histologically

confirmed liver fibrosis. Thus, in each fibrosis stage (F0/1, F2/3, and F4), elevated M2BPGi levels indicate a significant risk for developing HCC. Therefore, elevated M2BPGi levels may serve as a risk factor for developing HCC, independent of the stage of fibrosis.

M2BPGi as a prognostic factor after therapy for HCC

M2BPGi levels are a prognostic factor for patients with HCC (Table 1). For example, Fujiyoshi et al. [70] found that an elevated M2BPGi level is a significant risk factor for tumor recurrence and shorter overall survival in patients with HCC who undergo hepatectomy. Moreover, M2BPGi levels are not affected by tumor-related factors such as AFP, AFP L3, PIVKA II, tumor number, tumor size, lymph node metastasis, vascular invasion, and tumor differentiation. Toyoda et al. [71] found that an elevated M2BPGi level is a risk factor for tumor recurrence and shorter overall survival of patients with early-stage HCC who underwent curative resection. These investigators were unable to demonstrate a correlation between tumor-related factors and M2BPGi levels, and they suggested that M2BPGi represents the liver's background degree of oncogenic potential.

M2BPGi in other diseases

M2BPGi levels are elevated in organ fibrosis, such as idiopathic pulmonary fibrosis [72] and chronic pancreatitis [73], and correlate with the severity of disease. Moreover, M2BPGi levels are increased in pancreatic ductal adenocarcinoma [74], and elevated M2BPGi levels are associated with metastatic lesions. Pancreatic ductal adenocarcinoma is a cancer typical of those with dense fibrosis, and activation of stellate cells plays an important role in its progression [75]. Therefore, elevated M2BPGi levels in pancreatic ductal adenocarcinoma may reflect activation of cancer-associated stellate cells.

In conclusion, M2BPGi is a new, reliable surrogate marker of liver fibrosis and liver function. M2BPGi predicts the development of HCC, the prognosis of patients with liver cirrhosis, and patient prognosis after curative treatment for HCC. Moreover, elevated M2BPGi levels may reflect the activation of HSCs. Further clinical studies of HSC activation will likely contribute to a better understanding of disease.

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