1

# Relationship between Autoimmunity and Cancer or Metabolic Abnormality in Liver Diseases

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**Abstract:** Recent advances in the serological analysis of recombinant cDNA expression libraries and proteomic analysis have enabled the identification of numerous kinds of novel autoantigens, especially tumor-associated antigens (TAAs). The emergence of antibodies to TAAs might imply that an autoimmune response was involved in the malignant transformation. These autoantibodies to TAAs often have predictive value as well as diagnostic relevance. Similarly, the relationship between autoimmunity and metabolic abnormalities, including insulin resistance and hepatic steatosis, has been also documented. Metabolic abnormalities are frequently associated with autoimmune diseases. In contrast, autoimmune reactions were occasionally involved in the process of metabolic dysregulation. This review primarily focuses on the current trends for the relationship between autoimmunity and cancer or metabolic abnormalities in liver disease and the current interpretations of autoantibodies in those diseases.

**Keywords:** Autoantibodies, cancer, liver diseases, metabolic abnormalities, oxidatively modified autoantigens, stress proteins, tumor-associated antigens.

#### **INTRODUCTION**

Autoantibodies are generally produced by humoral immune responses against self-cellular proteins and nucleic acids [1], and have been well established as serological hallmarks of autoimmune disease. Recent advances in the molecular technology using a recombinant cDNA expression library (serological identification of antigens by recombinant expression cloning: SEREX) have enabled the isolation of novel autoantigens from the sera of patients with autoimmune or malignant diseases. Some of these autoantibodies often have clinical value for the diagnosis, and classification of the autoimmune disease and its disease activity. Other autoantibodies can even predict the prognosis of the autoimmune diseases and malignant diseases. Such autoantigens are ordinarily engaged in essential cellular functions including DNA replication, DNA transcription, and RNA processing [2].

The detection of autoantibodies is essential in the process of diagnosing autoimmune liver diseases, including autoimmune hepatitis (AIH), primary biliary cirrhosis (PBC) or primary sclerosing cholangitis (PSC). However, some of these autoantibodies are occasionally present even in the sera of patients with liver disease other than autoimmune liver diseases [3-7]. Such autoantibodies can be detected in the sera of patients with viral hepatitis, alcoholic liver disease, non-alcoholic fatty liver disease (NAFLD), drug-induced hepatitis, and hepatocellular carcinoma (HCC).

On the other hand, the relationship between autoimmunity and insulin resistance has been widely studied [8]. Patients with systemic lupus erythematosus (SLE) are often associated with metabolic syndrome [9, 10]. In addition, a recent article described the relationship between thyroid autoantibodies and obesity [11]. These findings may imply that the metabolic abnormalities trigger the production of a variety of autoantibodies.

This review mainly discusses the current trends of the relationship between autoimmunity and cancer or metabolic abnormalities in the liver diseases, and describes the interpretation of autoantibodies in those liver diseases.

# AUTOANTIBODIES IN THE SERA OF PATIENTS WITH HEPATOCELLULAR CARCINOMA

### 1) Methods for the Detection of Antibodies to Tumor-Associated Antigens

It has been well established that patients with autoimmune disease including Sjögren's syndrome are frequently complicated with the development of lymphoproliferative disorders, especially non-Hodgikin's lymphoma [12]. Inversely, autoimmune response is occasionally observed in cancer patients during the process of malignant transformation [13]. The autoantibodies detected in cancer patients are directed against nuclear antigens, against cytoplasmic antigens, and even against extracellular antigens.

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#### 2 The Open Autoimmunity Journal, 2014, Volume 6

HCC has been well recognized as one of the most common malignant tumors in the world with a high mortality. HCC has a clinical characteristic that develops from precursor conditions such as chronic hepatitis or liver cirrhosis [14]. Imai and colleagues documented that antinuclear antibodies (ANA) emerged *de novo* or the titers of ANA had been increased during the development of HCC from the precursor conditions [15]. These findings may support the hypothesis that an autoimmune response is involved in the process of malignant transformation.

Numerous types of autoantibodies to tumor-associated antigens (TAAs) have been identified in the sera of cancer patients. TAAs primarily consist of membrane receptors such as HER-2/neu oncoprotein [16], tumor suppressor gene proteins like p53 [17], proliferation-associated antigens including cyclin B1 [18] and centromere protein F (CENP-F) [19], and onconeural proteins such as Hu [20]. Table **1** summarizes the autoantibodies to TAAs identified from patients with HCC [18,19, 21-41]. Isolation of several kinds of TAAs has been accomplished using the recombinant cDNA expression library called serological identification of antigens by recombinant expression cloning (SEREX) [42]. Later, a method of serological proteome analysis (SERPA) (also called PROTEOMEX) has been proposed to trap autoantibodies to TAAs [43]. This approach is mainly based on a classical proteomics workflow combining effective separation on 2dimensional gel electrophoresis and identification by mass spectrometry. More recently, proteosome-covered array chips have been introduced as a novel technique for the detection of TAAs [44].

Table 1	Clinical significance of autoantibodies to TAAs in patients with HCC.

Intracellular Component	Autoantigens	<b>Biological functions</b>	Frequency of autoantibodies	References
Nucleus	c-myc	oncogenic protein	2-20%	[21,22]
	p53	tumor supressor gene protein	5-52%	[22-28]
	cyclin B1	cell-cycle related protein	15%	[29]
	p16	cyclin-dependent kinase inhibitor	21%	[30]
	survivin	apoptotic inhibitor	24%	[31]
	Ku86	DNA double-strand break repair	61%	[32]
	glucose-regulated protein 78	regulation of ER stress	36%	[33]
	DNA topoisomerase II	DNA replication/ transcription	N/A	[34]
	CENP-F	mitotic function	N/A	[19]
	HCC-1	mRNA splicing	N/A	[35]
	Fibrillarin	rRNA processing	N/A	[36]
	NOR-90/hUBF	RNA pol I transcription	N/A	[37]
	B23	Ribosome maturation	N/A	[36]
Cytoplasm	Golgi	processing, transporting and sorting	6%	[38]
		of intracellular proteins		
	IMP1	regulation of IGF-II mRNA	4-17%	[22,39]
	p62/IMP2	regulation of IGF-II mRNA	1-14%	[22,39]
	Koc/IMP3	regulation of IGF-II mRNA	5-10%	[22,39]
	fatty acid synthase	oncogenic protein	97%	[40]
	heat shock protein 70	stress protein	47%	[41]
	peroxiredoxin	cellular redox regulation	33%	[41]
	Mn-SOD	scavenger of oxidative stress	33%	[41]
			N/A,not available	

# 2) Putative Mechanism by which Autoantibodies to TAAs are Produced

Among these anti-TAAs, antibodies to p53 have been the most widely studied. The putative mechanism by which the autoantibodies are produced is as follows: the gene mutation of p53 causes an increase in the half-life of p53, leading to the accumulation of non-functional p53 protein in the nucleus of tumor cells. The accumulated p53 proteins are likely to become immunogenic, and subsequently evoke a humoral response to the p53 protein [17]. Similarly, antibodies to K-ras are present in approximately 20% of patients with esophageal carcinoma [45]. The production of antibodies to K-ras may also derive from the mutation of the K-ras gene [46]. Unfortunately, there is no study regarding antibodies to K-ras in patients with HCC so far.

We previously investigated the clinical significance of autoantibodies to <u>I</u>GF-II <u>m</u>RNA-binding proteins (anti-IMPs) in patients with HCC. The HCC patients with circulating anti-IMPs have overexpression of IMPs in the tumor cells, supporting the hypothesis that these autoantibodies are produced by an antigen-driven immune system [38].

It has been recently documented that gene promoter methylation plays important roles in the regulation of gene expression. Aberrant promoter methylation of the genes can directly lead to transcriptional inactivation and loss of gene function [47]. The methylation of tumor-associated genes may be involved in carcinogenesis [48-50] and may trigger the production of autoantibodies to TAAs (anti-TAAs).

#### 3) Diagnostic Significance of Anti-TAAs

In general, the sensitivities of anti-TAAs in patients with HCC are comparatively low, although their specificities seem to be extremely high. For example, the sensitivities of antibodies to p53 (anti-p53) in patients with HCC ranged from 5-52%, while their specificities ranged 96-100% in those patients [22-28]. It is notable that the status of anti-p53 appears to be independent of the elevation of serum  $\alpha$ -fetoprotein (AFP) levels, the tumor marker for HCC [23].

Recently, an enzyme-linked immunosorbent assay (ELISA) technique for the detection of antibodies to CENP-F was developed by Welner and colleagues [51]. They used overlapping synthetic peptides covering the predicted structural maintenance of chromosomes (SMC) domain. Half of the patients seropositive for anti-CENP-F were diagnosed with cancer using the ELISA method.

We previously examined the frequency of anti-TAAs, including anti-p53, c-myc, survivin and IMPs, in patients with HCC. Eight of 86 (9%) patients with HCC had one or more of these autoantibodies [21]. However, the levels of serum AFP in 7 of those 8 patients with anti-TAAs remained within normal limits, indicating that anti-TAAs are complementary tools for the diagnosis of HCC in those patients without elevation of AFP levels.

Interestingly, Akada and colleagues isolated antibodies to heat shock protein (HSP) 70 from the sera of patients with HCV-related HCC alone, implying that persistent HCV infection may induce a tumor-associated antigen during malignant transformation [44]. Recently, glucose-regulated protein 78 (GRP78), which is a chaperone protein belonging to the HSP70 family, was captured as a novel TAA in patients with HCC [32]. Fatty acid synthase was also identified as a novel TAA from those patients [39]. It is of interest that these TAAs are the proteins which originally contribute to glucose and lipid metabolism, respectively, implying that the metabolic dysregulation may evoke an autoimmune response during malignant transformation. Unfortunately, the clinical significance of these antibodies to HSP70, GRP78 and fatty acid synthase remains uncertain. Further examinations will be required to clarify it.

#### 4) Predictive Values of Anti-TAAs

A previous study documented that HCC patients with anti-p53 seemed to predict an unfavorable prognosis [26]. In, addition, an association between the emergence of anticentromere antibodies (ACA) and malignancy has been shown. ACA may be a risk factor for cancer in patients with systemic sclerosis [52]. We previously elucidated that ACA were rarely found in the sera of patients with HCV-related chronic liver disease (CLD) [53] as well as those with AIH and PBC [54], and that patients with HCV-related CLD seropositive for ACA often progressed to HCC [53].

On the other hand, we experienced a HCC patient who had circulating anti-p53, anti-IMP1 and anti-IMP3 simultaneously prior to the clinical diagnosis of HCC, suggesting that the emergence of these autoantibodies may predict the development of HCC [21].

#### 5) Anti-p53 in Autoimmune Liver Disease

Anti-p53 are sometimes present in the sera of patients with autoimmune diseases such as SLE [55], rheumatoid arthritis [56] and dermatomyositis/polymyositis [57]. It is of interest that the emergence of anti-p53 was not found to be associated with the mutation of p53 gene in patients with SLE [55].

Anti-p53 is considered to be useful to discriminate patients with AIH from those with PBC [58, 59]. The prevalence and titers of circulating anti-p53 were significantly higher in patients with AIH than in those with PBC. Cellmediated cytotoxicity is involved in the pathogenesis of AIH, while liver damage in patients with PBC is primarily caused by cholestasis. The existence of anti-p53 is likely to correspond to a secondary hallmark for autoimmune inflammation and stress. Taking these findings into consideration, the lower frequency of anti-p53 in the sera of patients with PBC can be explained.

It is remarkable that circulating anti-p53 detected in patients with AIH were predominantly directed against the Cterminal domain of p53 protein [58]. Moreover, the titers of anti-p53 in patients with AIH were significantly associated with the titers of antibodies to ds-DNA [59], suggesting that DNA damage may trigger the production of anti-p53.

## AUTOANTIBODIES IN SERA OF PATIENTS WITH LIVER DISEASE ASSOCIATED WITH METABOLIC ABNORMALITIES

#### 1) Relationship between Autoimmunity and Metabolic Abnormalities in Liver Diseases

In the field of autoimmune liver disease, insulin resistance is commonly observed in patients with AIH [60] and PBC [61]. The phenomena may explain that insulin resistance contributes to the development of autoimmune liver disease. Meanwhile, approximately 20-30% of patients with NAFLD, which is characterized by obesity, insulin resistance and other excessive nutritional intake, had ANA in their sera, suggesting that the autoimmune response might be involved in the process of insulin resistance [62-65]. The elevation of serum B-lymphocyte activating factor (BAFF) levels in patients with nonalcoholic steatohepatitis (NASH) may be attributed to insulin resistance [66].

Oxidative stress also plays crucial roles in the metabolic abnormalities including insulin resistance, hepatic steatosis, and iron overload [67, 68]. Oxidation of the host proteins appears to be responsible for the humoral response to the host proteins. Indeed, oxidatively modified autoantigens, including oxidized low-density lipoprotein (oxLDL), 4hydroxy-2-noneal (HNE)-modified 60 kD Ro and 8oxodeoxyguanine, were identified from patients with SLE as novel autoantigens [69].

Recent studies have revealed that oxidative modification of self-antigens may initiate a process of oxidative posttranslational modification intolerance, resulting in a primary B cell response against the posttranslationally modified selfantigen. Moreover, a secondary response may involve the development of autoreactive B cells by way of "epitope spreading", which is defined as the progression of an autoimmune response from initial activation to a chronic state involving increased targeting of autoantigens by T cells and antibodies [70].

#### 2) Clinical Significance of Autoimmunity in Liver Diseases Associated with Metabolic Abnormalities

A recent study focused on the relationship between ANA status and insulin resistance in patients with NAFLD. Loria and colleagues documented that the high titers of ANA may be an indicator for a severe insulin resistance in those patients [71].

Here, autoantibodies to stress proteins in the field of liver diseases and their significance are summarized in Table 2. Antibodies to oxidatively modified autoantigens were also trapped in the sera of patients with NAFLD and chronic hepatitis C (CH-C). Titers of malondialdehyde (MDA)-adducted

 Table 2.
 Clinical significance of autoantibodies to stress proteins in liver diseases

Autoantibodies	Molecular Target	Associated Liver Disease	Clinical Significance	References
anti-MDA	MDA	HCV	correlation with hepatic steatosis	[72]
		NAFLD	correlation with hepatic fibrosis	[73]
anti-oxLDL	oxLDL	HCV	correlation with hepatic steatosis	[75]
anti-SOD	SOD	AIH	false-positivity for anti-HCV	[78]
anti-cardiolipin	cardiolipin	HCV	correlation with oxidative stress	[79]
		HCV	higher prevalence of portal hypertension	[86]
		HCV	lack in the symptoms for anti-phospholipid syndrome	[79,81-84]
		HCV	concurrent oral lichen planus	[87]
		HCV	concurrent mixed cryoglobulin	[88]
		NASH	N/A	[79]
		AIH	Indicator for more advanced liver fibrosis	[89]
		PBC, PSC	indicator for more advanced liver fibrosis	[90]
anti-CYP2E1	CYP2E1	halothene-induced hepatitis	specific to IgG4 type of anti-CYP2E1	[101]
		alcoholic liver cirrhosis	N/A	[95,96]
		AIH	N/A	[97-99]
		HCV	indicator for more severe necroinflammation	[97,102]
anti-HSP	HSP65	AIH	molecular mimicry between HSP and SOD	[105]
	HSP70	PBC	indicator for disease activity	[106]
	HSP70	HCV	inverse correlation with diseae-activity	[108]

human serum albumin (HSA) were elevated in CH-C patients with severe hepatic steatosis [72]. To the contrary, an increase in titers of antibodies to MDA-HSA was independently associated with the progression of liver fibrosis in patients with NAFLD [73]. We previously elucidated that serum oxLDL concentration was significantly higher in patients with HCV-related CLD than in normal healthy controls [74] and that the titers of antibodies to oxLDL were significantly associated with the extent of hepatic steatosis in those patients [75]. However, the titers of antibodies to oxLDL were independent of insulin resistance in those patients. In contrast, antibodies to oxLDL were closely related to atherosclerosis in patients with rheumatoid arthritis [76]. Atherosclerosis has been considered one of the metabolic abnormalities caused by HCV infection [77]. Therefore, the emergence of antibodies to oxLDL may predict atherosclerosis in patients with CH-C. Further examinations are required to clarify the association.

An interesting study on the relationship antibody to superoxide dismutase (SOD) and anti-HCV in patients with AIH was previously reported by Ikeda and colleagues [78]. They revealed that the titers of anti-HCV were closely linked to the titers of antibodies to SOD in those patients, suggesting that antibodies to SOD may account for the false-positive result of anti-HCV status in those patients.

On the other hand, our previous study elucidated that the titers of antibodies to cardiolipin (anti-cardiolipin), the serological hallmark for anti-phospholipid syndrome, were associated with the extent of hepatic 8-hydroxy-2'-deoxyguanosine (8-OHdG) expression in patients with CH-C [79]. 80HdG is well recognized as oxidatively generated DNA damage [80], implying that the emergence of anti-cardiolipin was associated with oxidative stress in those patients. However, none of the CH-C patients seropositive for anticardiolipin fulfilled the category for anti-phospholipid syndrome [79, 81-84]. It is of interest that anti-cardiolipin in patients with CH-C were in generally low titers, and were not associated with antibodies to \u03b32-glycoprotein-I. Crossreactivity between HCV antigen and phospholipid is speculated to trigger the production of anti-cardiolipin. Persistent HCV infection may disrupt cell membrane and favor the exposure of hidden phospholipids [85]. Patients with HCVrelated CLD seropositive for anti-cardiolipin had a trend toward portal hypertension [86] or advanced hepatic fibrosis [79]. Anti-cardiolipin may be also indicative for concurrent oral lichen planus [87] and mixed cryoglobulinemia [88] in CH-C patients, although the findings remain controversial [79].

Anti-cardiolipin were also found in the sera of patients with autoimmune liver disease including AIH, PBC and PSC [89, 90]. Similarly, those patients with anti-cardiolipin had a higher prevalence of liver cirrhosis than those patients without anti-cardiolipin.

In addition, cytochrome p450 2E1 (CYP2E1) is an endoplasmic monooxygenase and major source of oxidative stress in microsomes [91]. Overexpression of CYP2E1 in the liver tissues is frequently observed in patients with NASH [92] and CH-C [93]. Interestingly, the hepatic CYP2E1 expression was related to the severity of hepatic steatosis in patients with CH-C [93]. In contrast, circulating antibodies to CYP2E1 were originally detected in the sera of patients with halothane-induced hepatitis [94] and alcoholic liver cirrhosis [95, 96]. Recently, CYP2E1 was also identified as a target antigen from patients with AIH [97-99], while CYP2D6 has been well recognized as the target antigen of antibodies to liver-kidney microsome type 1 (anti-LKM1) [100]. More recently, Njoku and colleagues revealed that the IgG4 type of antibodies to CYP2E1 was specific to the patients with anesthetic-induced hepatitis [101]. On the other hand, approximately 5-40% of patients with CH-C possessed circulating antibodies to CYP2E1 in their sera [97, 102]. The molecular mimicry between HCV NS5 and CYP2E1 may initiate the humoral response to CYP2E1 in those patients [103]. CH-C patients with circulating antibodies to CYP2E1 had more severe necroinflammation in the liver than CH-C patients without antibodies to CYP2E1 [102], as usually did CH-C patients seropositive for ANA [104].

The HSPs, a family of ubiquitous and highly conserved proteins which play essential roles as molecular chaperones, appear to be antigenic. Antibodies to HSP65 detected in patients with AIH had cross-reactivity against SOD [105]. In contrast, antibodies to HSP 70 (anti-HSP 70) were found the sera of patients with PBC [106] and CH-C [107, 108]. Anti-HSP 70 seemed to reflect the disease activity in patients with PBC, while the titers of anti-HSP 70 were inversely associated with serum alanine aminotransferase (ALT) levels and the loads of HCV-RNA in patients with CH-C [108].

### CONCLUSION

In summary, autoimmune response is involved in the process of malignant transformation and metabolic dysregulation. Some types of autoantibodies to TAAs seem to be useful as a complementally diagnostic marker or predictive marker of HCC development. Other types of autoantibodies to stress proteins frequently reflect the severity of metabolic dysregulation and oxidative stress in the liver diseases.

#### **CONFLICT OF INTEREST**

The authors have no conflict of interest to declare.

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#### 8 The Open Autoimmunity Journal, 2014, Volume 6

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