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Table 2. Baseline characteristics of the study patients based on HCV RNA detection at 24

weeks

Characteristic		HCV RNA(-) n=56	HCV RNA(+) n=28	P value
Age	years	61.0 (30.0-78.0)	58.5 (36.0-71.0)	0.75
Male:Female	ratio	29:27	14:14	1
Previous IFN therapy	no	11	7	0.88
	yes	40	20	
Body mass index	Kg/m ²	22.2 (18.1-30.8)	22.4 (19.0-31.6)	0.67
Stage of fibrosis	0-2/3,4/ND	25 / 16 / 15	10 / 13 / 5	0.35
IL28B rs8099917	TT/TG/GG/ND	45 / 6 / 1 / 4	14 / 10 / 1 / 3	<u>0.003</u>
HCV-RNA log ₁₀	IU/ml	6.3 (4.7-7.3)	6.3 (4.7-7.3)	1
HCV genotype 1b	No.(%)	56(66.7%)	28(33.3%)	
Platelet count	no./μl	16.7 (9.0-27.2)	15.7 (7.8-36.2)	0.78
Hemoglobin	g/dl	14.3 (11.6-16.3)	13.7 (10.9-17.0)	0.71
ALT*	IU/L	50.5 (12.0-495.0)	64.0 (10.0-246.0)	0.14
AST*	IU/L	51.0 (13.0-182.0)	44.5 (12.0-236.0)	0.094
γ-GTP*	IU/L	35.0 (11.0-282.0)	61.0 (14.0-240.0)	0.057
HbA1c*	%	5.1 (4.6-11.8)	5.1 (3.9-10.2)	0.51
T-Cho*	mg/dl	179.5 (125.0-257.0)	173.0 (116.0-242.0)	0.49

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25(OH)vitamin D ₃	ng/ml	24.0 (12.0-40.0)	20.0 (12.0-39.0)	0.29
Vitamin D +/-	%	33 / 23	9 / 18	<u>0.029</u>
Reduction case	-/PEG/RBV/+	20 / 4 / 19 / 8	11 / 0 / 7 / 8-	
Adherence to PEG IFN	%	69.0%	No case	
Adherence to RBV	%	82.5%	80.8%	0.76
Adherence to both	PEG IFN (%)	83.0%	81.7%	0.66
PEGIFN+RBV	RBV (%)	79.6%	81.7%	0.71

*ALT, alanine aminotransferase; AST, aspartate aminotransferase; γ -GTP, γ -glutamyl

transpeptidase; HbA1c, hemoglobin A1c; T-Cho, total cholesterol

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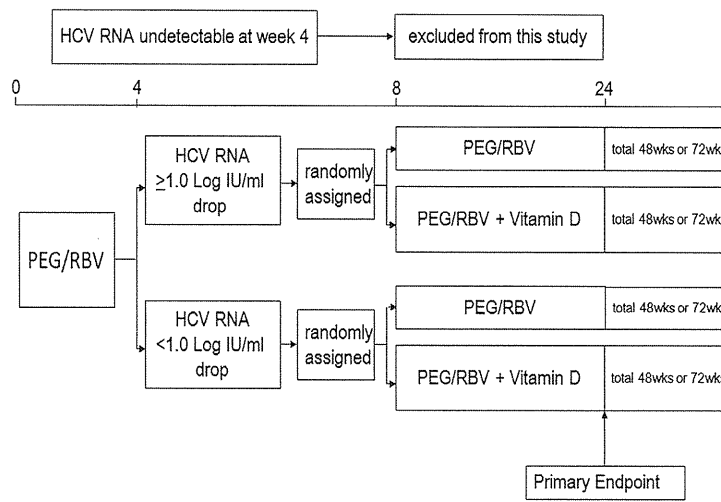
Table 3. Logistic regression analysis of factors contributing to undetectable HCV RNA at week 24.

	Crude OR (95%CI)	P value	Adjusted OR (95%CI)	P value
Age	1.00 (0.96-1.05)	0.81		
Gender	1.29 (0.48-3.46)	0.50		
Platelet count	1 (0.91-1.1)	0.79		
ALT*	0.99 (0.99-1.00)	0.051		
AST*	0.99 (0.98-1.0)	0.054		
γ -GTP*	0.99 (0.98-1)	0.42		
Virus amount	1.03 (0.43-2.46)	0.67		
Previous IFN therapy	2.54 (0.93-6.98)	0.24		
Vitamin D +/-	2.54 (0.93-6.98)	0.062	3.12 (1.03-9.5)	<u>0.04</u>
IL28B	4.96 (1.59-15.41)	0.004	5.85 (1.75-19.57)	<u>0.004</u>

*ALT, alanine aminotransferase; AST, aspartate aminotransferase; γ -GTP, γ -glutamyl transpeptidase

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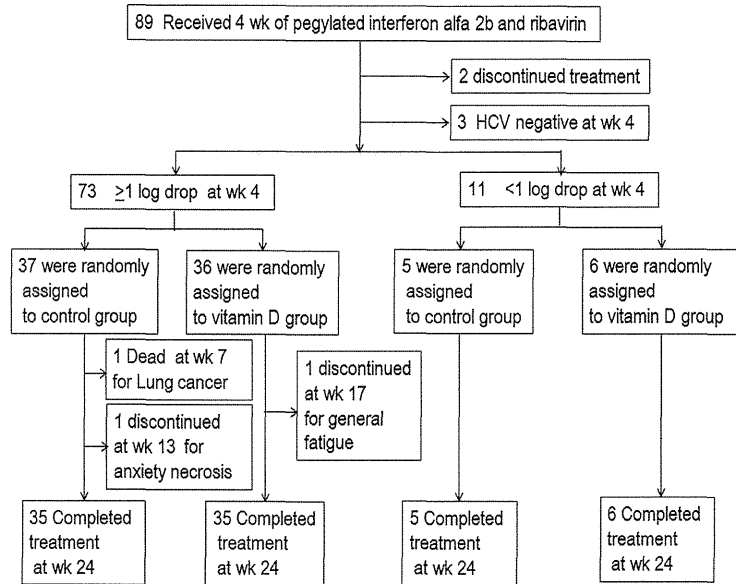
Figure. 1



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Figure. 2

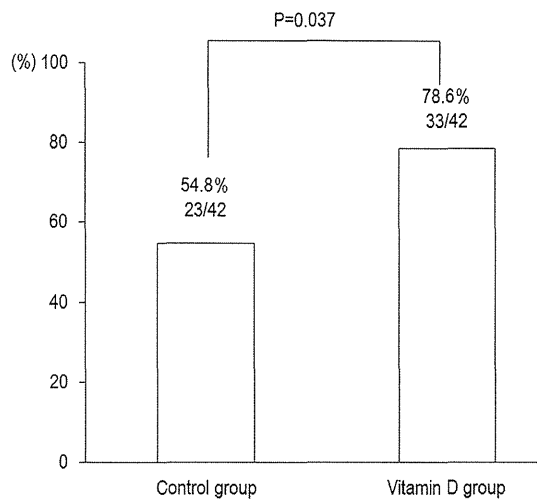


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REVIEW

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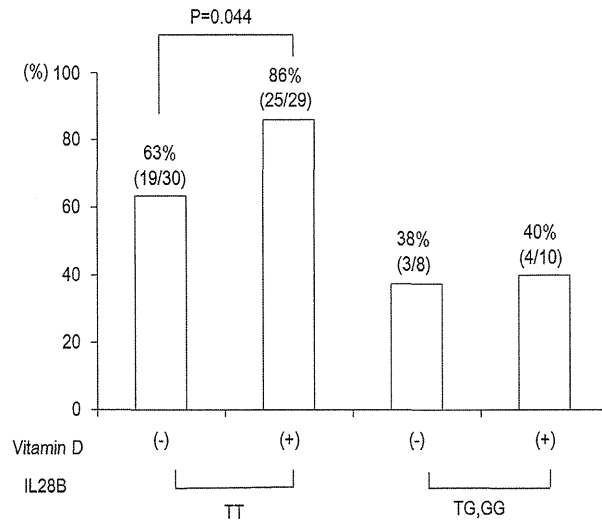
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Figure. 4

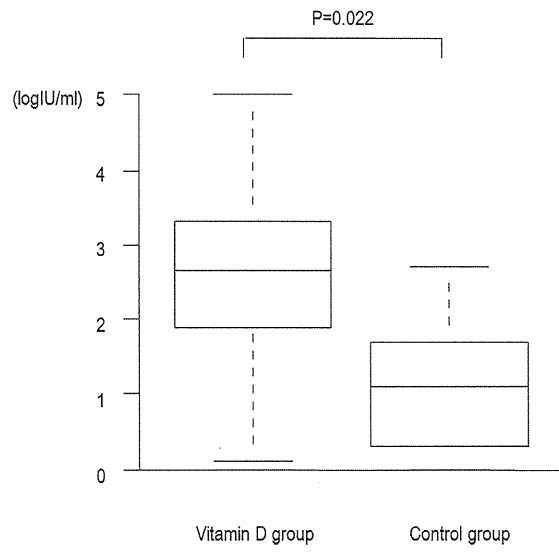


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Figure. 5



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Review

Risk Factors for Long-Term Persistence of Serum Hepatitis B Surface Antigen Following Acute Hepatitis B Virus Infection in Japanese Adults

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The proportion of patients who progress to chronicity following acute hepatitis B (AHB) varies widely worldwide. Moreover, the association between viral persistence after AHB and hepatitis B virus (HBV) genotypes in adults remains unclear. A nationwide multicenter study was conducted throughout Japan to evaluate the influence of clinical and virological factors on chronic outcomes in patients with AHB. For comparing factors between AHB patients with viral persistence and those with self-limited infection, 212 AHB patients without human immunodeficiency virus (HIV) coinfection were observed in 38 liver centers until serum hepatitis B surface antigen (HBsAg) disappeared or a minimum of 6 months in cases where HBsAg persisted. The time to disappearance of HBsAg was significantly longer for genotype A patients than that of patients infected with non-A genotypes. When chronicity was defined as the persistence of HBsAg positivity for more than 6 or 12 months, the rate of progression to chronicity was higher in patients with genotype A, although many cases caused by genotype A were prolonged cases of AHB, rather than chronic infection. Multivariate logistic regression analysis revealed only genotype A was independently associated with viral persistence following AHB. A higher peak level of HBV DNA and a lower peak of alanine aminotransferase (ALT) levels were characteristics of AHB caused by genotype A. Treatment with nucleotide analogs (NAs) did not prevent progression to chronic infection following AHB overall. Subanalysis suggested early NA initiation may enhance the viral clearance. **Conclusion:** Genotype A was an independent risk factor for progression to chronic infection following AHB. Our data will be useful in elucidating the association between viral persistence after AHB, host genetic factors, and treatment with NAs in future studies. (HEPATOLOGY 2014;59:89-97)

Abbreviations: AHB, acute hepatitis B; ALT, alanine aminotransferase; anti-HBc, antibody to hepatitis B core antigen; anti-HBs, antibody to HBsAg; HBeAg, hepatitis B e-antigen; CLIA, chemiluminescent enzyme immunoassay; ELA, enzyme immunoassay; HBsAg, hepatitis B surface antigen; HBV, hepatitis B virus; HIV, human immunodeficiency virus; IgM, immunoglobulin M; anti-HBe, antibody to HBeAg; NAs, nucleotide analogs; RPHA, reverse passive hemagglutination.

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Hepatitis B virus (HBV) infection is one of the most common persistent viral infections in humans. Approximately 2 billion people worldwide have been exposed to HBV and 350 million of them remain persistently infected.¹ The incidence of HBV infection and the patterns of transmission vary greatly worldwide among different population subgroups.² In Western countries, chronic HBV infection is relatively rare and acquired primarily in adulthood by way of sexual transmission or the use of injectable drugs. Meanwhile, in Asia and most of Africa, the majority of infections are the result of transmission from an infected mother to her newborn. However, very few studies of acute hepatitis B (AHB) in adults have been reported.

HBV genomic sequences vary worldwide and have been classified into at least eight genotypes (A through H) based on an intergroup divergence of 8% or more over the complete nucleotide sequence.^{3,4} These genotypes have distinct geographic distributions.⁵⁻⁷ In particular, genotype A is predominant in Northwestern Europe, the United States, Central Africa, and India.^{8,9} The Japanese have been infected with genotypes B and C since prehistoric times.¹⁰ Recently, many lines of evidence have revealed among the Japanese an increase in acute infection with HBV genotype A following sexual transmission.^{11,12} As a result of this increasing transmission of genotype A, the distribution of HBV genotypes in Japan clearly differs among patients with acute and chronic infections.¹³ Moreover, recent studies suggest that acute infection with HBV genotype A may be associated with an increased risk of progression to persistent infection.¹⁵ Indeed, the prevalence of HBV genotype A in chronic hepatitis B patients doubled in Japan between 2000-2001 and 2005-2006 (1.7% versus 3.5%).¹¹

Human immunodeficiency virus (HIV)-1 infection results in an immunodeficient state, with the virus sharing routes of transmission with HBV. HIV-related immunodepletion influences the natural history of HBV infection, and epidemiological studies have

revealed that HIV-positive patients are more likely to have a prolonged acute illness following HBV infection and lower rates of hepatitis B e-antigen (HBeAg) clearance.¹⁶ Therefore, in this study patients with coinfection of HIV were excluded to examine the influence of HBV genotype directly without the confounding influence of HIV.

From 2005 to 2010, a multicenter cohort study was conducted throughout Japan on 212 patients with AHB. The aim of this cohort study was to assess the influence of clinical and virological factors, including HBV genotypes and treatment with nucleotide analogs (NAs), on AHB patients who became persistently infected.

Patients and Methods

Patients With AHB. The multiple-source cohort included 212 randomly selected AHB patients without coinfection of HIV. From 2005 through 2010, the study participants were recruited from 38 liver centers throughout Japan. The cohort included patients who were admitted to the hospital because of AHB and who visited the hospital every month after being discharged. The diagnosis of AHB was contingent on the rapid onset of clinical symptoms accompanied by elevated serum alanine aminotransferase (ALT) levels, the detection of serum hepatitis B surface antigen (HBsAg), and a high-titer antibody to hepatitis B core antigen (anti-HBc) of the immunoglobulin M (IgM) class. Patients with initial high-titer anti-HBc (>10.0 S/CO) were diagnosed as having an exacerbation of chronic hepatitis B and were excluded. If the patient had been tested previously, the absence of serum HBsAg and anti-HBc before admission was verified from the medical record to discriminate a new infection from an acute exacerbation of a persistent infection. Patients with acute hepatitis A, hepatitis C, and drug- or alcohol-induced acute hepatitis were also excluded; hepatitis D virus infection was not determined because of its extreme rarity in Japan. The study protocol conformed to the 1975 Declaration of

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Potential conflict of interest: Nothing to report.

Additional Supporting Information may be found in the online version of this article.

Table 1. Characteristics of Patients With Genotype A or a Non-A Genotype Acutely Infected With Hepatitis B Virus

Features	Genotype A (n = 107)	Non-A Genotypes (n = 105)*	P Value
Age (years)	36.3 ± 12.0	40.7 ± 14.3	0.032
Male sex	102 (95.3)	75 (71.4)	<0.001
HBeAg positive	104 (97.2)	79 (75.2)	<0.001
ALT (IU/L)	1210 ± 646	2225 ± 2851	0.045
Total bilirubin (mg/dL)	9.9 ± 9.4	7.5 ± 6.7	0.115
HBV DNA (log copies/mL)	7.0 ± 1.5	5.8 ± 1.5	<0.0001
Duration until disappearance of HBsAg (month)	6.7 ± 8.5	3.4 ± 6.5	<0.0001
Persistence of HBsAg positivity more than 6 months	25 (23.4)	9 (8.6)	0.003
Persistence of HBsAg positivity more than 12 months	8 (7.5)	1 [†] (0.9)	0.018
Sexual transmission	81/84 (96.4) [‡]	71/79 (89.9) [§]	0.095
Treatment with NAs	61 (57.0)	42 (40.0)	0.013

Data are presented as n (%), mean ± standard deviation. HBV, hepatitis B virus; HBeAg, hepatitis B e-antigen; ALT, alanine aminotransferase; NAs, nucleotide analogs.

*Non-A genotypes include genotypes B, C, D, F and H (n = 25, 77, 1, 1, and 1, respectively).

[†]One patient had genotype C.

[‡]Transmission routes were unknown for 23 patients.

[§]Transmission routes were unknown for 26 patients.

Helsinki and was approved by the Ethics Committees of the institutions involved. Every patient gave informed consent for this study.

Serological Markers of HBV Infection. HBsAg, HBeAg, antibodies to HBsAg (anti-HBs), HBeAg (anti-HBe), and HBcAg, and anti-HBc of the IgM class were tested by a chemiluminescent enzyme immunoassay (CLIA) by ARCHITECT (Abbott Japan, Tokyo, Japan). HBV DNA measurements were performed using a real-time polymerase chain reaction (PCR) assay (Cobas TaqMan HBV Auto; Roche Diagnostics, Tokyo, Japan).

Genotyping of HBV. The six major HBV genotypes (A through F) were determined serologically by enzyme immunoassay (EIA) using commercial kits (HBV GENOTYPE EIA; Institute of Immunology, Tokyo, Japan). This method is based on the pattern of detection by monoclonal antibodies of a combination of epitopes on preS2-region products, which is specific for each genotype.^{17,18} Samples for which EIA could not determine the genotype were examined by direct sequencing of the pre-S2/S gene, followed by phylogenetic analysis.

Treatment With NAs. Treatments with NAs were performed using lamivudine or entecavir for more than 3 months. The individual clinicians determined if NAs were administered to patients, and when the treatment was to be started. The time to onset of treatment with NAs was measured in days from onset of AHB.

Statistical Analysis. Categorical variables were compared between groups by the chi-squared test and noncategorical variables by the Mann-Whitney *U* test.

A *P* value less than 0.05 was considered significant. Multivariate analysis was performed using a backward stepwise logistic regression model to determine independent factors for viral persistence following AHB. Variables in the multivariate analysis were selected based on variables that were marginally significant with *P* < 0.1 in univariate analysis. Maintenance of HBsAg positivity was analyzed using the Kaplan-Meier method and significance was tested with the log-rank test. STATA Software (StataCorp, College Station, TX) v. 11.0 was used for analyses.

Results

Comparison of Characteristics Between Genotype A and Non-A Genotype AHB Patients. A total of 107 AHB patients (50.5%) were infected with genotype A while 105 AHB patients (49.5%) were infected with non-A genotypes, including genotypes B (25 [11.8%]), C (76 [35.8%]), D (1 [0.5%]), F (1 [0.5%]), and H (1 [0.5%]). Compared to those infected with non-A genotypes, genotype A patients were significantly younger (36.3 ± 12.0 versus 40.7 ± 14.3 years, *P* = 0.032), predominantly men (95.3% versus 71.4%, *P* < 0.001), and more frequently positive for HBeAg (97.2% versus 75.2%, *P* < 0.001). Moreover, genotype A patients had a lower peak ALT levels (1,210 ± 646 versus 2,225 ± 2,851 IU/L, *P* = 0.045) and a higher peak level of HBV DNA (6.7 ± 8.5 versus 3.4 ± 6.5 log copies/mL, *P* < 0.0001). A significantly higher percentage of genotype A patients were treated with NAs (57% versus 40%, *P* = 0.013). These data are summarized in Table 1.

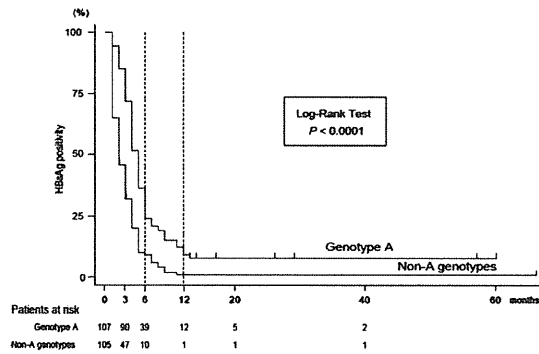


Fig. 1. Comparison of the cumulative proportion of AHB patients maintaining HBsAg positivity between genotype A and non-A genotypes, analyzed using the Kaplan-Meier test. $P < 0.0001$, genotype A: red line, non-A genotypes: blue line.

Cumulative Maintenance of HBsAg Positivity During Follow-up in Patients With Genotype A and Non-A Genotypes.

In the patients infected with genotype A and non-A genotypes, the mean durations of HBsAg positivity maintenance were 6.7 ± 8.5 and 3.4 ± 6.5 months, respectively ($P < 0.0001$; Table 1, Fig. 1). For 6 months after AHB onset, the number of patients with genotype A and non-A genotypes maintaining HBsAg positivity were 39/107 (36.4%) and 10/105 (9.5%), respectively ($P < 0.001$). However, in many patients HBsAg disappeared between 7 and 12 months after AHB onset; that is, HBsAg disappeared in 31/107 (29.0%) of patients with genotype A and in 9/105 (8.6%) of patients with non-A genotypes during this time period. However, in some patients HBsAg never disappeared after persisting for more than 12

months following AHB onset. When chronicity after AHB was defined as the persistence of HBsAg for more than 12 months, chronicity developed in 7.5% (8/107) of patients with genotype A and in 0.9% (1/105) of patients with non-A genotypes ($P = 0.018$).

Comparison of Characteristics Between Patients in Whom HBsAg Persisted More Than 6 or 12 Months and Those With Self-Limited AHB Infection.

Table 2 compares the demographic and clinical characteristics between patients in whom HBsAg disappeared within 6 months and those in whom HBsAg persisted for more than 6 months from AHB. The peak ALT levels ($1,882 \pm 2,331$ versus $1,018 \pm 696$ IU/L, $P = 0.0024$) and peak HBV DNA levels (6.3 ± 1.6 versus 7.4 ± 1.6 mg/dL, $P = 0.0004$) were significantly higher and lower in the former group than in the latter group, respectively. Moreover, marked differences were present in the distribution of genotypes between the two groups. The percentage of the HBV genotype A (46.1% versus 73.5%, $P = 0.003$) was significantly higher among patients in whom HBsAg was persistent for more than 6 months. In addition, we compared the demographic and clinical characteristics between patients in whom HBsAg disappeared within 12 months and those in whom HBsAg persisted for more than 12 months from AHB. Peak ALT ($1,787 \pm 2,118$ versus 775 ± 513 IU/L, $P = 0.0089$) and peak total bilirubin (8.7 ± 8.2 versus 3.8 ± 6.6 mg/dL, $P = 0.0039$) levels were significantly higher in the former group than in the latter group. In contrast, the peak HBV DNA levels (6.4 ± 1.6 versus 7.9 ± 1.4 mg/dL, $P = 0.0046$) were significantly lower

Table 2. Comparison Between Patients With Chronicity Following Acute Hepatitis B and Those With Self-Limited Acute Infections Determined by the Persistence of HBsAg for More Than 6 or 12 Months

Features	Persistence of HBsAg		P Value	persistence of HBsAg for More Than 12 Months		P Value
	Disappearance of HBsAg Within 6 Months (n = 178)	for More Than 6 Months From AHB (n = 34)		Disappearance of HBsAg Within 12 Months (n = 203)	From AHB (n = 9)	
Age (years)	38.2 ± 13.1	40.0 ± 14.5	0.454	38.1 ± 13.2	46.7 ± 14.0	0.061
Male sex	147 (82.6)	30 (88.2)	0.416	169 (83.3)	8 (88.9)	0.677
HBeAg positive	150 (84.3)	32 (94.1)	0.131	175 (86.2)	8 (88.9)	0.815
ALT (IU/L)	1882 ± 2331	1018 ± 696	0.0024	1787 ± 2118	775 ± 513	0.0089
Total bilirubin (mg/dL)	8.6 ± 7.5	8.7 ± 11.3	0.137	8.7 ± 8.2	3.8 ± 6.6	0.0039
HBV DNA (log copies/mL)	6.3 ± 1.6	7.4 ± 1.6	0.0004	6.4 ± 1.6	7.9 ± 1.4	0.0046
HBV genotype						
Non-A	96 (53.9)	9 (26.5)		104 (51.2)	1 (11.1)	
A	82 (46.1)	25 (73.5)	0.003	99 (48.8)	8 (88.9)	0.018
Sexual transmission	128/137 (93.4)*	24/26 (92.3) [†]	0.711	146/157 (93.0) [‡]	6/6 (100.0) [§]	0.356
NAs treatment (+)	82 (46.1)	21 (61.8)	0.093	98 (48.3)	8 (88.9)	0.017

Data are presented as n (%) and mean \pm SD. HBsAg, hepatitis B surface antigen; AHB, acute hepatitis B, HBeAg, hepatitis B e-antigen; ALT, alanine aminotransferase; HBV, hepatitis B virus; NAs, nucleotide analogs.

*Transmission routes of 41 patients were unknown.

[†]Transmission routes of 8 patients were unknown.

[‡]Transmission routes of 46 patients were unknown.

[§]Transmission routes of 3 patients were unknown.

Table 3. Multivariate Analysis of Factors Independently Associated With Persistence of HBsAg Positivity Following Acute Hepatitis B

Factors	Persistence of HBsAg More Than 6 Months From AHB		
	Odds Ratio	95% CI	P Value
ALT (per 1 IU/L increase)	1.000	0.999-1.000	0.035
HBV DNA (per 1 log copy/mL increase)	1.176	0.931-1.484	0.173
Genotypes			
Non-A	1.00		
A	4.224	1.853-9.631	0.001

95% CI, 95% confidence interval; ALT, alanine aminotransferase; HBV, hepatitis B virus.

in the former group than in the latter group. The percentages of HBV genotype A (48.8% versus 88.9%, $P=0.018$) and NAs treatment (+) (48.3% versus 88.9%, $P=0.017$) were significantly higher among patients in whom the HBsAg persisted for more than 12 months.

Factors Independently Associated With Viral Persistence Following AHB. A stepwise logistic regression model was used to perform multivariate analysis which explains relationships between some factors and persistence of HBsAg positivity more than 6 months following AHB. Peak ALT level, peak HBV DNA level, genotype A, and treatment with NAs were retained in the final multivariate logistic model in a backward stepwise manner ($P<0.1$). For predicting the persistence of HBsAg for more than 6 months, only genotype A was independently associated with progression of AHB to the persistence of HBsAg (odds ratio [OR]: 4.224, $P=0.001$, Table 3).

Characteristics of Patients Who Progressed to Chronicity That Was Defined as the Persistence of HBsAg for More Than 12 Months Following Acute Hepatitis B. Table 4 shows the clinical and virological characteristics of nine patients who progressed to

chronicity defined as the persistence of HBsAg for more than 12 months following AHB. Among the nine patients who progressed to chronicity from AHB, eight (88.9%) were men and eight (88.9%) were HBeAg-positive. In general, among the patients who progressed to chronicity following AHB, the peak HBV DNA levels were high, and the peak total bilirubin and ALT levels were low. In eight (88.9%) patients, entecavir was administered; however, the duration until the onset of NA treatment from AHB onset was long (75-570 days).

Early Onset of Treatment With NAs Was Able to Prevent Viral Persistence After AHB Caused by Genotype A. The cumulative proportion maintaining HBsAg positivity during follow-up, expressed in terms of time after AHB onset, were significantly longer in patients with NAs treatment than in those without NAs treatment ($P=0.046$, Fig. 2A). Table 5 shows the percentages of patients in whom HBsAg persisted for more than 6 or 12 months among patients categorized based on the period of time (i.e., duration) until the onset of NAs treatment. For patients in whom the onset of NAs treatment was less than 4 weeks from the onset of AHB, 12.7% of the patients showed persistent HBsAg for more than 6 months, while none showed HBsAg positivity for more than 12 months. For patients in whom the onset of NAs treatment was at 5-8 weeks, 37.5% of the patients showed persistent HBsAg for more than 6 months, whereas none showed persistent HBsAg for more than 12 months. For all groups, the period of HBsAg positivity in patients starting NAs treatment within 8 weeks from AHB onset was significantly shorter than that in patients beginning NAs treatment after more than 8 weeks from AHB onset ($P<0.0001$, Fig. 2B). Patients starting NAs treatment within 8 weeks from AHB onset never progressed to chronicity after AHB caused by genotype A.

Table 4. Characteristics of Patients Who Progressed to Chronicity Following Acute Hepatitis B

Case	Age	Gender	HIV	HBeAg	HBV DNA (log copies/mL)	Total Bilirubin (mg/dL)	ALT (IU/L)	Observation Period (Months)	NAs Treatment	Duration Until		Transmission Routes	Genotype
										NAs Treatment	NAs Treatment (Days)		
1	23	Male	(-)	(+)	7.6	1.7	1271	26	ETV	570	Heterosexual	A	
2	40	Male	(-)	(-)	8.8	1.4	568	13	ETV	240	Heterosexual	A	
3	45	Male	(-)	(+)	7.7	0.9	867	57	ETV	135	Heterosexual	A	
4	37	Male	(-)	(+)	7.6	3.4	384	29	ETV	75	Unknown	A	
5	54	Male	(-)	(+)	9	2	455	17	ETV	155	Homosexual	A	
6	45	Male	(-)	(+)	4.8	21.2	512	60	(-)	(-)	Homosexual	A	
7	61	Male	(-)	(+)	9.1	1.5	804	17	ETV	88	Unknown	A	
8	56	Male	(-)	(+)	9.0	1.1	1820	14	ETV	118	Unknown	A	
9	31	Female	(-)	(+)	7.4	0.8	296	66	ETV	150	Blood transfusion	C	

HIV, human immunodeficiency virus; HBeAg, hepatitis B e-antigen; HBV, hepatitis B virus; ALT, alanine aminotransferase; NAs, nucleotide analogs; ETV, entecavir.

Table 5. Proportion of Patients in Whom HBsAg Persisted for More Than 6 or 12 Months Among Patients Categorized Based on the Number of Weeks Until the Onset of NAs Treatment

Duration Until Onset of NAs Treatment (Weeks)	Persistence of HBsAg for More Than 6 Months	Persistence of HBsAg for More Than 12 Months	Total Patients
<4 weeks (n, %)	9 (12.7)	0 (0)	71
5-8 weeks (n, %)	6 (37.5)	0 (0)	16
9-12 weeks (n, %)	1 (33.3)	1 (33.3)	3
13-16 weeks (n, %)	4 (100)	1 (25.0)	4
>17 weeks (n, %)	9 (100)	6 (66.7)	9
Total	29	8	103

HBsAg, hepatitis B surface antigen; NAs, nucleotide analogs.

Discussion

A multicenter nationwide study was conducted throughout Japan to evaluate the influence of clinical and virological factors on chronic outcomes in Japanese patients who contracted AHB in adulthood. The study was feasible in Japan, where a universal vaccination program for HBV has not been implemented because of the extremely high efficacy of the immunoprophylaxis that is given to babies born to carrier mothers. The implementation of this program has resulted in a decrease in the persistent HBV carrier rate from 1.4% to 0.3%.¹⁹ Selective vaccination means that Japanese are more likely to be infected with HBV by way of horizontal transmission since the percentage of the population possessing anti-HBs is much lower than that in countries in which universal vaccination programs have been established.²⁰ In addition, Japan is faced with the ever-increasing impacts of globalization: as many as 17 million Japanese travel abroad and over 7 million people

visit Japan from overseas each year. This “population mixing” may help to explain the increased prevalence in Japan of AHB due to genotype A, which is transmitted through indiscriminate sexual contact. Consequently, Japan may be the only country in the world where the influences of HBV genotypes, including genotype A (as is predominant in Western countries) and genotypes B and C (as are predominant in Asian countries), on chronic outcomes after AHB can be compared.

Currently, the persistence of HBsAg in serum for more than 6 months is considered to represent a progression to chronic infection.²¹ However, our data showed that HBsAg frequently disappeared between 7 to 12 months after the onset of AHB in patients with genotype A (31/107 [29.0%]) and non-A genotypes (9/105 [8.6%]) (Fig. 1). These patients were considered to exhibit prolonged cases of AHB, rather than persistent infection. This finding reflects the higher sensitivity of the most up-to-date assays for HBsAg as compared with previous methods. In the present study, HBsAg was measured by CLIA, which has been reported to be about 150 times more sensitive in the detection of HBsAg than reverse passive hemagglutination (RPHA)-HBsAg, which has been used for the last 30 years in Japan.²² The use of a more sensitive assay for HBsAg results in a longer period during which HBsAg may be detected. In this study, HBsAg did not disappear in nine patients after remaining continuously detectable for more than 12 months. Therefore, the persistence of HBsAg for more than 12 months, as measured with a highly sensitive method for detecting HBsAg, may be suitable for defining the progression of AHB to chronicity; however, further study is necessary to determine whether this definition is appropriate worldwide.

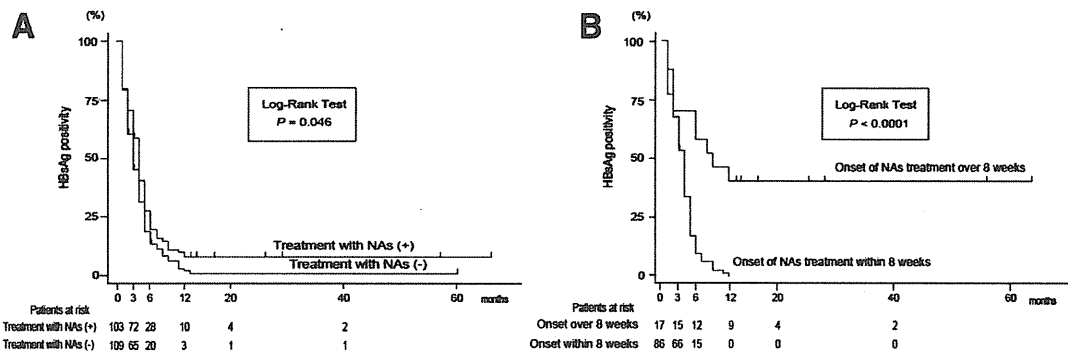


Fig. 2. (A) Comparison of the cumulative proportion of AHB patients maintaining HBsAg positivity between treatment with NAs (+) and treatment with NAs (-), as analyzed using the Kaplan-Meier test. $P = 0.046$, treatment with NAs (+): red line, treatment with NAs (-): blue line. (B) Comparison of the cumulative proportion of AHB patients in genotype A maintaining HBsAg positivity between treatment onset with NAs within 8 weeks and treatment onset with NAs over 8 weeks after onset of AHB, as analyzed using the Kaplan-Meier test. $P < 0.0001$, treatment onset with NAs over 8 weeks: red line, treatment onset with NAs within 8 weeks: blue line.

It has been reported that ~10% of patients who contract HBV as adults do not clear HBsAg from their serum and become carriers.²³ Meanwhile, a wide variation has been seen in the rate of persistence after AHB infection in adults. For example, viral persistence following AHB was seen in 0.2% (1/507) of adults in Greece,²⁴ 7.7% (5/65) of adult Alaskan Eskimos, and 12.1% (7/58) of adults in Germany.²⁵ The difference in the proportion of patients progressing from AHB to chronicity in different regions may be attributable to virological and host factors. In this study, 4.2% (9/212) of patients progressed to chronicity after AHB: 7.5% (8/107) of those infected with genotype A and 0.9% (1/105) of those infected with non-A genotypes. The non-A genotypes included genotypes B, C, D, F, and H (n = 25, 77, 1, 1, and 1, respectively). Genotypes B and C are predominant in eastern Asian countries, where the majority of those infected with HBV acquired the virus during the perinatal period by way of vertical transmission.²⁶ On the other hand, genotype A is predominant in Western countries, where the main route is horizontal transmission later in life.^{26,27} Because HBeAg persists long after the infection in the genotype C as compared to other genotypes, this genotype has been shown to be a risk factor for perinatal and horizontal transmission in newborns and children.²⁸ The predominance of genotype A in Western countries may be attributable to a higher chronicity rate following AHB by way of horizontal transmission in adults.

In this study the characteristics of AHB associated with genotype A were a higher peak level of HBV DNA and a lower peak level of ALT. These findings were similar to those for patients with HBV-HIV coinfection.²⁹ Such characteristics of genotype A or coinfection with HIV are assumed to be attributable to milder hepatitis associated with weaker cellular immune responses. More slowly replicating viruses have been reported to evoke weaker cellular responses, enhancing the likelihood of persistence.³⁰ Indeed, our prior study showed that the replication of genotype A was significantly slower than that of genotype C in immunodeficient, human hepatocyte chimeric mice.³¹ Moreover, variation among genotypes in the expression pattern of HBeAg may affect the progression of AHB to chronicity. Another previous study of ours revealed that a single form of HBeAg was detected by western blot analysis in serum samples from patients infected with genotypes B through D, but that two additional larger forms of HBeAg were detected in patients with genotype A.³² Milich and Liang³³ reported that HBeAg may modulate the host immune response as a

tolerogen to promote chronicity. Therefore, the different expression pattern of HBeAg by genotype A HBV may contribute to chronicity following AHB.

Early NAs initiation appeared to enhance the viral clearance across genotypes, although treatment with NAs did not show any overall benefit in duration of HBsAg. Previous studies examining the efficacies of NAs for preventing progression to chronic infection after AHB have reported conflicting results. Some small-scale studies have suggested the efficacy of lamivudine and entecavir in preventing the progression of AHB to chronic hepatitis.^{34,35} Another study showed a lower seroconversion rate of HBsAg in lamivudine users.³⁶ Further, a randomized placebo-controlled trial showed no significant difference in clinical outcomes.³⁷ However, these previous studies did not mention the prevalence of HBV genotypes in the respective study populations. Although this was a retrospective study, our study included data on the prevalence of HBV genotypes. Additionally, our findings suggested that larger prospective randomized studies for every HBV genotype should be performed to determine whether early treatment with NAs prevented the progression of AHB to a chronic state.

In conclusion, in Japan genotype A was an independent risk factor for progression to chronic infection following AHB in adults. Confirmation of this association in patients with AHB in other countries is desirable and may provide insight into the pathogenetic mechanisms underlying this association. Early NA treatment appeared to reduce the likelihood of chronicity but this potentially important intervention needs to be prospectively studied before recommendations can be made.

Appendix

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References

- Mast EE, Alter MJ, Margolis HS. Strategies to prevent and control hepatitis B and C virus infections: a global perspective. *Vaccine* 1999;17:1730-1733.
- Lavanchy D. Worldwide epidemiology of HBV infection, disease burden, and vaccine prevention. *J Clin Virol* 2005;34(Suppl 1):S1-S3.
- Okamoto H, Tsuda F, Sakugawa H, Sastroewignjo RI, Imai M, Miyakawa Y, et al. Typing hepatitis B virus by homology in nucleotide sequence: comparison of surface antigen subtypes. *J Gen Virol* 1988;69(Pt 10):2575-2583.
- Norder H, Hammas B, Lofdahl S, Courouce AM, Magnus LO. Comparison of the amino acid sequences of nine different serotypes of hepatitis B surface antigen and genomic classification of the corresponding hepatitis B virus strains. *J Gen Virol* 1992;73(Pt 5):1201-1208.
- Miyakawa Y, Mizokami M. Classifying hepatitis B virus genotypes. *Intervirology* 2003;46:329-338.
- Fung SK, Lok AS. Hepatitis B virus genotypes: do they play a role in the outcome of HBV infection? *HEPATOLOGY* 2004;40:790-792.
- Norder H, Courouce AM, Coursaget P, Echevarria JM, Lee SD, Mushahwar IK, et al. Genetic diversity of hepatitis B virus strains derived worldwide: genotypes, subgenotypes, and HBsAg subtypes. *Intervirology* 2004;47:289-309.
- Kurbanov F, Tanaka Y, Mizokami M. Geographical and genetic diversity of the human hepatitis B virus. *Hepatol Res*;40:14-30.
- Kao JH. Hepatitis B viral genotypes: clinical relevance and molecular characteristics. *J Gastroenterol Hepatol* 2002;17:643-650.
- Orito E, Ichida T, Sakugawa H, Sata M, Horiike N, Hino K, et al. Geographic distribution of hepatitis B virus (HBV) genotype in patients with chronic HBV infection in Japan. *HEPATOLOGY* 2001;34:590-594.
- Matsuura K, Tanaka Y, Hige S, Yamada G, Murawaki Y, Komatsu M, et al. Distribution of hepatitis B virus genotypes among patients with chronic infection in Japan shifting toward an increase of genotype A. *J Clin Microbiol* 2009;47:1476-1483.
- Ozasa A, Tanaka Y, Orito E, Sugiyama M, Kang JH, Hige S, et al. Influence of genotypes and precore mutations on fulminant or chronic outcome of acute hepatitis B virus infection. *HEPATOLOGY* 2006;44:326-334.
- Kobayashi M, Ikeda K, Arase Y, Suzuki F, Akuta N, Hosaka T, et al. Change of hepatitis B virus genotypes in acute and chronic infections in Japan. *J Med Virol* 2008;80:1880-1884.
- Mayerat C, Mantegani A, Frei PC. Does hepatitis B virus (HBV) genotype influence the clinical outcome of HBV infection? *J Viral Hepat* 1999;6:299-304.
- Ogawa M, Hasegawa K, Naritomi T, Torii N, Hayashi N. Clinical features and viral sequences of various genotypes of hepatitis B virus compared among patients with acute hepatitis B. *Hepatol Res* 2002;23:167-177.
- Gilson RJ, Hawkins AE, Beecham MR, Ross E, Waite J, Briggs M, et al. Interactions between HIV and hepatitis B virus in homosexual men: effects on the natural history of infection. *AIDS* 1997;11:597-606.
- Usuda S, Okamoto H, Iwanari H, Baba K, Tsuda F, Miyakawa Y, et al. Serological detection of hepatitis B virus genotypes by ELISA with monoclonal antibodies to type-specific epitopes in the preS2-region product. *J Virol Methods* 1999;80:97-112.
- Usuda S, Okamoto H, Tanaka T, Kidd-Ljunggren K, Holland PV, Miyakawa Y, et al. Differentiation of hepatitis B virus genotypes D and E by ELISA using monoclonal antibodies to epitopes on the preS2-region product. *J Virol Methods* 2000;87:81-89.
- Noto H, Terao T, Ryou S, Hirose Y, Yoshida T, Ookubo H, et al. Combined passive and active immunoprophylaxis for preventing perinatal transmission of the hepatitis B virus carrier state in Shizuoka, Japan during 1980-1994. *J Gastroenterol Hepatol* 2003;18:943-949.
- Yoshikawa A, Suzuki K, Abe A, Tanaka T, Yamaguchi K, Ishikawa Y, et al. Effect of selective vaccination on a decrease in the rate of hepatitis B virus-positive Japanese first-time blood donors. *Transfus Med* 2009;19:172-179.
- Lok AS, McMahon BJ. Chronic hepatitis B. *HEPATOLOGY* 2007;45:507-539.
- Sato S, Ohhashi W, Ihara H, Sakaya S, Kato T, Ikeda H. Comparison of the sensitivity of NAT using pooled donor samples for HBV and that of a serologic HBsAg assay. *Transfusion* 2001;41:1107-1113.
- Sherlock SDJ, editor. *Virus hepatitis*. London: Blackwell Scientific; 1997.
- Tassopoulos NC, Papaevangelou GJ, Sjogren MH, Roumeliotou-Karayannis A, Gerin JL, Purcell RH. Natural history of acute hepatitis B surface antigen-positive hepatitis in Greek adults. *Gastroenterology* 1987;92:1844-1850.

25. McMahon BJ, Alward WL, Hall DB, Heyward WL, Bender TR, Francis DP, et al. Acute hepatitis B virus infection: relation of age to the clinical expression of disease and subsequent development of the carrier state. *J Infect Dis* 1985;151:599-603.
26. Kao JH, Chen DS. Global control of hepatitis B virus infection. *Lancet Infect Dis* 2002;2:395-403.
27. Chu CJ, Keeffe EB, Han SH, Perrillo RP, Min AD, Soldevila-Pico C, et al. Hepatitis B virus genotypes in the United States: results of a nationwide study. *Gastroenterology* 2003;125:444-451.
28. Livingston SE, Simonetti JP, Bulkow LR, Homan CE, Snoweball MM, Cagle HH, et al. Clearance of hepatitis B e antigen in patients with chronic hepatitis B and genotypes A, B, C, D, and F. *Gastroenterology* 2007;133:1452-7.
29. Colin JF, Cazals-Hatem D, Lorient MA, Martinot-Peignoux M, Pham BN, Auperin A, et al. Influence of human immunodeficiency virus infection on chronic hepatitis B in homosexual men. *HEPATOLOGY* 1999;29:1306-1310.
30. Bocharov G, Ludewig B, Bertoletti A, Klenerman P, Junt T, Krebs P, et al. Underwhelming the immune response: effect of slow virus growth on CD8⁺-T-lymphocyte responses. *J Virol* 2004;78:2247-2254.
31. Sugiyama M, Tanaka Y, Kato T, Orito E, Ito K, Acharya SK, et al. Influence of hepatitis B virus genotypes on the intra- and extracellular expression of viral DNA and antigens. *HEPATOLOGY* 2006;44:915-924.
32. Ito K, Kim KH, Lok AS, Tong S. Characterization of genotype-specific carboxyl-terminal cleavage sites of hepatitis B virus e antigen precursor and identification of furin as the candidate enzyme. *J Virol* 2009;83:3507-3517.
33. Milich D, Liang TJ. Exploring the biological basis of hepatitis B e antigen in hepatitis B virus infection. *HEPATOLOGY* 2003;38:1075-1086.
34. Lisotti A, Azzaroli F, Buonfiglioli F, Montagnani M, Alessandrelli F, Mazzella G. Lamivudine treatment for severe acute HBV hepatitis. *Int J Med Sci* 2008;5:309-312.
35. Jochum C, Gieseler RK, Gawlista I, Fiedler A, Manka P, Saner FH, et al. Hepatitis B-associated acute liver failure: immediate treatment with entecavir inhibits hepatitis B virus replication and potentially its sequelae. *Digestion* 2009;80:235-240.
36. Yu JW, Sun LJ, Zhao YH, Kang P, Li SC. The study of efficacy of lamivudine in patients with severe acute hepatitis B. *Dig Dis Sci* 2010;55:775-783.
37. Kumar M, Satapathy S, Monga R, Das K, Hissar S, Pande C, et al. A randomized controlled trial of lamivudine to treat acute hepatitis B. *HEPATOLOGY* 2007;45:97-101.

トピックス

KeyWords

ウイルス肝炎に関する国の対策事業，公費助成や受診勧奨など

- ◎ウイルス肝炎
- ◎インターフェロン
- ◎核酸アナログ
- ◎医療費助成制度
- ◎検診事業

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Headline

1. 肝炎に対する抗ウイルス療法には医療費助成制度が設けられており，課税所得に応じて自己負担額の上限が決められる。
2. B型肝炎に対する核酸アナログ製剤は継続内服が基本となるため，医療費助成の更新が認められている。
3. 肝炎検診事業において無料検査が実施されていることを多くの人が認知しておらず，普及啓発に関する広報活動の推進が必要である。

はじめに

昨今の肝炎治療の進歩はめざましく，患者の予後改善が大いに図られているところである。B型肝炎については，2000年にラミブジンが核酸アナログ製剤の一番手として臨床応用され，その後，ラミブジン耐性ウイルスが高率に出現することが問題となったが，そのレスキュー薬剤として2004年アデホビルが承認された。その後，エンテカビルが2006年に承認され，現在における第一選択薬剤として位置づけられている。さらに，治療効果および忍容性において従来のインターフェロン（interferon:IFN）製剤を凌駕するペグインターフェロン（pegylated interferon;PEG-IFN） α 2a製剤が2011年に承認されたことで，B型肝炎の病態コントロールがさらに可能となっている。一方，C型肝炎治療薬として，リバビリン（ribavirin:RBV）が2001年に登場し，その後，PEG-IFN製剤として α 2aが2003年， α 2bが2004年に承認されたことにより，わが国に多い1b型・高ウイルス量の患者においても50%の著効率が期待しうる状況となった。さらに，2011年11月にはプロテアーゼ阻害薬

テラプレビル（telaprevir:TPV）が承認されたことで，治療期間が従来の48～72週間から24週間に短縮されたにもかかわらず，初回治療例における著効率が70%以上と極めて満足すべき効果が得られるようになった。しかし，これら新規薬剤は薬価がかなり高額であることも認識する必要がある。例えば，B型肝炎治療薬のラミブジン・アデホビル併用療法では医療費3割負担の場合でも月額15,742円となり，エンテカビルでは月額8,671円となる。C型肝炎のPEG-IFN+RBV+TPV 3剤併用療法になると，体重60 kg以上の患者では3割負担で月額11万5,000円を超える。しかし，これらの抗ウイルス療法が奏効すれば，患者の生命予後を左右する肝硬変・肝細胞がんへの進展が有意に抑制されることから，適応のある患者を1人でも多く治療すべきであることは言うまでもない。そのためには「自身の感染を知らずに社会に潜在している」¹⁾肝炎ウイルスキャリアを発掘する必要があるわけで，肝炎検診事業の重要性が叫ばれるゆえんもそこにある。

表1 肝炎治療特別促進事業による医療費助成制度の推移

2008年4月 2009年4月	B型・C型肝炎に対するインターフェロン（IFN）治療 IFN医療費助成の運用変更 ①助成期間の延長（72週投与への対応） ②所得階層区分の認定に係る例外的取扱い
2010年4月	肝炎医療費助成の拡充 ①自己負担限度額の引下げ 所得に応じ，1，3，5万円→原則1万円（上位所得階層2万円） ②B型肝炎の核酸アナログ製剤治療への助成開始 ③IFN治療に係る利用回数の制限緩和 ④身体障害者福祉法における肝臓機能障害の追加 （非代償性肝硬変の一部が該当）
2011年7月 2011年9月26日 2011年12月22日 2011年12月26日	C型代償性肝硬変に対するペガシス+コペガス併用療法 B型慢性肝炎に対するペガシス単独療法 C型代償性肝硬変に対するペグイントロン+レボトル併用療法 1b・高ウイルス量のC型慢性肝炎へのペグイントロン+レボトル+テラプレビル3剤併用療法

ウイルス性肝炎に関する公費助成

1. 医療費助成制度の仕組み

先述したように，B型・C型肝炎のIFN療法やB型肝炎の核酸アナログ製剤治療では，かなり高額な医療費が必要であるため，国と自治体は肝炎治療特別促進事業の一環として，IFN治療については2008年度から，核酸アナログ製剤については2010年度から医療費助成制度の拡充に取り組んできた（表1）．自己負担額の月額上限は前年度の課税所得に応じて1万円ないし2万円の2段階に設定されている（2010年4月改定までは1，3，5万円の3段階）．

医療費助成の対象項目として，IFNや核酸アナログ製剤等の肝炎治療薬剤，血液検査，腹部超音波・CT・MRIなどの画像診断，および，抗ウイルス療法に起因すると担当医が判断した有害事象の治療にかかわる医療費等が含まれるとの解釈が一般的である．B型肝炎に対する核酸アナログ製剤の場合，安易な内服中止はむしろ肝炎を増悪させる危険性があるため，基本的には内服を継続する必要がある．そのため，核酸アナログ製剤の医療費助成に関しては1年ごとの更新が認められている．一方，B型・C型肝炎に対するIFN治療

の場合は，治療薬剤の種類によって医療費助成期間に制限があるが，一定の条件を満足する場合には，再治療や期間延長に対しても医療費助成が認可される．これら抗ウイルス療法に対する医療費助成の認定基準を表2に示す．なお，2011年11月に保険収載されたTPV併用療法の取り扱いが極めて複雑であり，過去のPEG-IFN+RBV 2剤併用療法施行の有無やウイルス学的反応性等によって医療費助成制度適用の可否が決まる．詳細は他書²⁾に譲る．

2. 医療費助成交付件数の推移

国と地方自治体が2011年度までに行った交付件数が厚生労働省から公表されている．これによると，IFN医療費助成については初年度が43,536人と最も多く，その後減少したが26,595人，28,797人とほぼ横這いで推移し，2011年度はTPV以降の新規薬剤の登場を見込んで治療導入が手控えられたためか，17,674人であった（図1）．この4年間に約11万7,000人が受給したことになる．また，核酸アナログ製剤医療費助成治療についても2010年度，2011年度における交付件数が公表されており，2011年度には初年度からの更新分36,766人と新規分11,916人を合わせて約4万9,000人が受給している．特に交付件数の

表2 B型・C型慢性肝疾患に対する医療費助成の認定基準

1. B型慢性肝疾患	
(1) インターフェロン (IFN) 治療について	HBe抗原陽性かつHBV DNA陽性のB型慢性活動性肝炎でIFN治療を行う予定、またはIFN治療実施中の者のうち、肝がんの合併のないもの（ただし、PEG-IFN製剤を用いる治療に限っては、HBe抗原陰性のB型慢性活動性肝炎も対象とする。） ※上記において2回目の助成を受けることができるのは、これまでにPEG-IFN製剤による治療を受けたことがない者が同製剤による治療を受ける場合とする。
(2) 核酸アナログ製剤治療について	B型肝炎ウイルスの増殖を伴い肝機能の異常が確認されたB型慢性肝疾患で核酸アナログ製剤治療を行う予定、または核酸アナログ製剤治療実施中の者
2. C型慢性肝疾患	
(1) IFN単剤治療ならびにIFNおよびRBV併用治療について	HCV RNA陽性のC型慢性肝炎およびC型代償性肝硬変でIFN治療を行う予定、またはIFN治療実施中の者のうち、肝がんの合併のないもの。ただし、これまでの治療において、十分量の3剤併用療法（PEG-IFN、RBVおよびTPV）による24週投与が行われた場合を除く。 ※上記において2回目の助成を受けることができるのは、以下の①、②のいずれにも該当しない場合とする。 ①これまでの治療において、十分量のPEG-IFNおよびRBV併用療法による48週投与を行ったが、36週目までにHCV RNAが陰性化しなかったケース ②これまでの治療において、PEG-IFNおよびRBV併用療法による72週投与が行われたケース
(2) PEG-IFN、RBVおよびTPV 3剤併用療法について	HCV RNA陽性のC型慢性肝炎で、PEG-IFN、RBVおよびTPVによる3剤併用療法を行う予定、または実施中の者のうち、これまでに3剤併用療法を受けたことがなく、かつ肝がんの合併のないもの
※1上記については、1回のみ助成とする。ただし、2. (1)に係る治療歴の有無を問わない。 ※2 3剤併用療法の実施は、日本皮膚科学会皮膚科専門医（日本皮膚科学会が認定する専門医主研修施設または研修施設に勤務する者に限る。）と連携し、日本肝臓学会肝臓専門医が常勤する医療機関に限る。	

IFN:interferon, RBV:ribavirin, TPV:telaprevir
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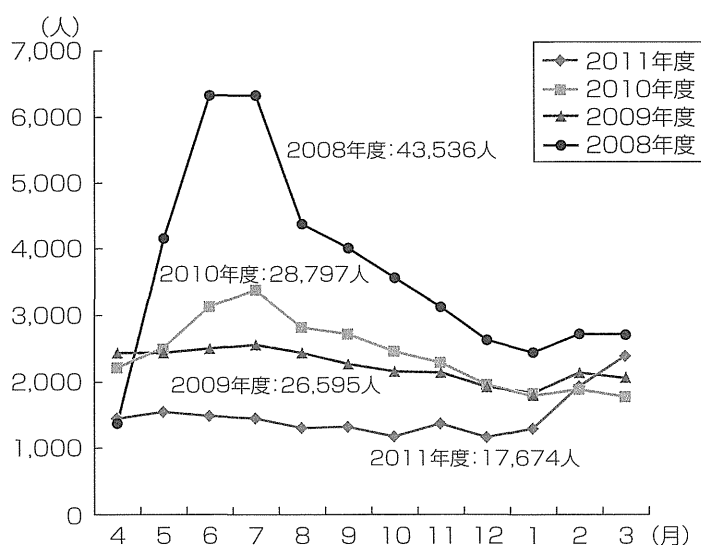


図1 肝炎IFN治療受給者証交付件数の推移
2008~2011年度の月ごとの交付件数を示す。
(http://www.mhlw.go.jp/bunya/kenkou/kekkaku-kansenshou09/080328_josei.htmより抜粋)