

level and 3% of individuals with an ALT level of ≥ 201 IU/l have ongoing subclinical infection of various HEV strains, suggesting the frequent occurrence of subclinical HEV infection, although clinical HEV infection is rarely reported. A large study of individuals who do not have an elevated ALT level is needed to assess the exact frequency of subclinical HEV infection, taking into consideration the geographic region in Japan.

REFERENCES

- Abe T, Aikawa T, Akahane Y, Arai M, Asahina Y, Atarashi Y, Chayama K, Harada H, Hashimoto N, Hori A, Ichida T, Ikeda H, Ishikawa A, Ito T, Kang JH, Karino Y, Kato H, Kato M, Kawakami M, Kitajima N, Kitamura T, Masaki N, Matsubayashi K, Matsuda H, Matsui A, Michitaka K, Mihara H, Miyaji K, Miyakawa H, Mizuo H, Mochida S, Moriyama M, Nishiguchi S, Okada K, Saito H, Sakugawa H, Shibata M, Suzuki K, Takahashi K, Yamada G, Yamamoto K, Yamanaka T, Yamato H, Yano K, Mishiro S. 2006. Demographic, epidemiological, and virological characteristics of hepatitis E virus infections in Japan based 254 human cases collected nationwide. *Kanzo* 47:384-391.
- Aggarwal R, Kamili S, Spelbring J, Krawczynski K. 2001. Experimental studies on subclinical hepatitis E virus infection in cynomolgus macaques. *J Infect Dis* 184:1380-1385.
- Amon JJ, Drobeniuc J, Bower WA, Magana JC, Escobedo MA, Williams IT, Bell BP, Armstrong GL. 2006. Locally acquired hepatitis E virus infection, El Paso, Texas. *J Med Virol* 78:741-746.
- Buisson Y, Grandadam M, Nicand E, Cheval P, van Cayck-Gandre H, Innis B, Rehel P, Coursaget P, Teyssou R, Tsarev S. 2000. Identification of a novel hepatitis E virus in Nigeria. *J Gen Virol* 81:903-909.
- Caudill JD, Malik IA, Tsarev SA. 1994. Evidence for human hepatitis E virus (HEV) infection without acute antibody response. *Am J Trop Hyg* 51:201.
- Clayson ET, Bruce L, Innis BL, Myint KSA, Narupth S, Vaughn DW, Giri S, Ranabhat P, Shrestha MP. 1995. Detection of hepatitis E virus infections among domestic swine in the Kathmandu valley of Nepal. *Am J Med Hyg* 53:228-232.
- Emerson SU, Purcell RH. 2003. Hepatitis E virus. *Rev Med Virol* 13:145-154.
- Emerson SU, Anderson D, Arankalle A, Meng XJ, Purdy M, Schlauder GG, Tsarev SA. 2004. Hepatitis E virus. In: Fauquet CM, Mayo MA, Maniloff J, Desselberger U, Ball LA, editors. *Virus taxonomy*, The eighth report of the International Committee on Taxonomy of Viruses. Elsevier/Academic Press, London: pp 851-855.
- Felsenstein J. 1985. Confidence limits on phylogenies: An approach using the bootstrap. *Evolution* 39:783-791.
- Fukuda S, Sunaga J, Saito N, Fujimura K, Itoh Y, Sasaki M, Tsuda F, Takahashi M, Nishizawa T, Okamoto H. 2004. Prevalence of antibodies to hepatitis E virus among Japanese blood donors: Identification of three blood donors infected with a genotype 3 hepatitis E virus. *J Med Virol* 73:554-561.
- Harrison TJ. 1999. Hepatitis E virus—an update. *Liver* 19:171-176.
- Ijaz S, Arnold E, Banks M, Bendall RP, Cramp ME, Cunningham R, Dalton HR, Harrison TJ, Hill SF, Macfarlane L, Meigh RE, Shafi S, Sheppard MJ, Smithson J, Wilson MP, Teo CG. 2005. Non-travel-associated hepatitis E in England and Wales: Demographic, clinical, and molecular epidemiological characteristics. *J Infect Dis* 192:1166-1172.
- Ina Y. 1994. ODEN: A program package for molecular evolutionary analysis and database search of DNA and amino acid sequences. *Comput Appl Biosci* 10:11-12.
- Inoue J, Nishizawa T, Takahashi M, Aikawa T, Mizuo H, Suzuki K, Shimosegawa T, Okamoto H. 2006. Analysis of the full-length genome of genotype 4 hepatitis E virus isolates from patients with fulminant or acute self-limited hepatitis E. *J Med Virol* 78:476-484.
- Koizumi Y, Isoda N, Sato Y, Iwaki T, Ono K, Ido K, Sugano K, Takahashi M, Nishizawa T, Okamoto H. 2004. Infection of a Japanese patient by genotype 4 hepatitis E virus while traveling in Vietnam. *J Clin Microbiol* 42:3883-3885.
- Koonin EV, Gorbalenya AE, Purdy MA, Rozanov MN, Reyes GR, Bradley DW. 1992. Computer-assisted assignment of functional domains in the nonstructural polyprotein of hepatitis E virus: Delineation of an additional group of positive-strand RNA plant and animal viruses. *Proc Natl Acad Sci USA* 89:8259-8263.
- Kwo PY, Schlauder GG, Carpenter HA, Murphy PJ, Rosenblatt JE, Dawson GJ, Mast EE, Krawczynski K, Balan V. 1997. Acute hepatitis E by a new isolate acquired in the United States. *Mayo Clin Proc* 72:1133-1136.
- Li TC, Zhang J, Shinzawa H, Ishibashi M, Sata M, Mast EE, Kim K, Miyamura T, Takeda N. 2000. Empty virus-like particle-based enzyme-linked immunosorbent assay for antibodies to hepatitis E virus. *J Med Virol* 62:327-333.
- Lu L, Li C, Hagedorn CH. 2006. Phylogenetic analysis of global hepatitis E virus sequences: Genetic diversity, subtypes and zoonosis. *Rev Med Virol* 16:5-36.
- Mansuy JM, Peron JM, Abravanel F, Poirson H, Dubois M, Miedouge M, Vischi F, Alric L, Vinel JP, Izopet J. 2004. Hepatitis E in the south west of France in individuals who have never visited an endemic area. *J Med Virol* 74:419-424.
- Matsubayashi K, Nagaoka Y, Sakata H, Sato S, Fukai K, Kato T, Takahashi K, Mishiro S, Imai M, Takeda N, Ikeda H. 2004. Transfusion-transmitted hepatitis E caused by apparently indigenous hepatitis E virus strain in Hokkaido, Japan. *Transfusion* 44:934-940.
- Matsuda H, Okada K, Takahashi K, Mishiro S. 2003. Severe hepatitis E virus infection after ingestion of uncooked liver from a wild boar. *J Infect Dis* 188:944.
- Meng XJ. 2005. Hepatitis E as a zoonotic disease. In: Thomas HC, Lemon S, Zuckerman AJ, editors. *Viral hepatitis 3rd ed*. Blackwell Publishing, Malden, MA: p 611-623.
- Mine H, Emura H, Miyamoto M, Tomono T, Minegishi K, Murokawa H, Yamanaka R, Yoshikawa A, Nishioka K, Japanese Red Cross NAT Research Group. 2003. High throughput screening of 16 million serologically negative blood donors for hepatitis B virus, hepatitis C virus and human immunodeficiency virus type-1 by nucleic acid amplification testing with specific and sensitive multiplex reagent in Japan. *J Virol Methods* 112:145-151.
- Mitsui T, Tsukamoto Y, Yamazaki C, Masuko K, Tsuda F, Takahashi M, Nishizawa T, Okamoto H. 2004. Prevalence of hepatitis E virus infection among hemodialysis patients in Japan: Evidence for infection with a genotype 3 HEV by blood transfusion. *J Med Virol* 74:563-572.
- Mitsui T, Tsukamoto Y, Suzuki S, Yamazaki C, Masuko K, Tsuda F, Takahashi M, Tsatsralt-Od B, Nishizawa T, Okamoto H. 2005. Serological and molecular studies on subclinical hepatitis E virus infection using periodic serum samples obtained from healthy individuals. *J Med Virol* 76:526-533.
- Mizuo H, Suzuki K, Takikawa Y, Sugai Y, Tokita H, Akahane Y, Itoh K, Gotanda Y, Takahashi M, Nishizawa T, Okamoto H. 2002. Polyphyletic strains of hepatitis E virus are responsible for sporadic cases of acute hepatitis in Japan. *J Clin Microbiol* 40:3209-3218.
- Nakamura M, Takahashi K, Taira K, Taira M, Ohno A, Sakugawa H, Arai M, Mishiro S. 2006. Hepatitis E virus infection in wild mongooses of Okinawa, Japan: Demonstration of anti-HEV antibodies and a full-genome nucleotide sequence. *Hepato Res* 34:137-140.
- Nicand E, Grandadam M, Teyssou R, Rey JL, Buisson Y. 2001. Viraemia and faecal shedding of HEV in symptom-free carriers. *Lancet* 357:68-69.
- Okamoto H, Takahashi M, Nishizawa T. 2003. Features of hepatitis E virus infection in Japan. *Intern Med* 42:1065-1071.
- Preiss JC, Plentz A, Engelmann E, Schneider T, Jilg W, Zeitz M, Duchmann R. 2006. Autochthonous hepatitis E virus infection in Germany with sequence similarities to other European isolates. *Infection* 34:173-175.
- Purcell RH, Emerson SU. 2001. Hepatitis E virus. In: Knipe DM, Howley PM, Griffin DE, Martin MA, Lamb RA, Roizman B, Straus SE, editors. *Fields virology 4th ed*. Lippincott Williams and Wilkins, Philadelphia, PA: pp 3051-3061.
- Sadler GJ, Mells GF, Shah NH, Chesner IM, Walt RP. 2006. UK acquired hepatitis E—An emerging problem? *J Med Virol* 78:473-475.
- Saitou N, Nei M. 1987. The neighbor-joining method: A new method for reconstructing phylogenetic trees. *Mol Biol Evol* 4:406-425.
- Schlauder GG, Mushahwar IK. 2001. Genetic heterogeneity of hepatitis E virus. *J Med Virol* 65:282-292.
- Smith JL. 2001. A review of hepatitis E virus. *J Food Prot* 64:572-586.
- Takahashi K, Iwata K, Watanabe N, Hatahara T, Ohta Y, Baba K, Mishiro S. 2001. Full-genome nucleotide sequence of a hepatitis E virus strain that may be indigenous to Japan. *Virology* 287:9-12.

- Takahashi K, Kang JH, Ohnishi S, Hino K, Mishiro S. 2002. Genetic heterogeneity of hepatitis E virus recovered from Japanese patients with acute sporadic hepatitis. *J Infect Dis* 185:1342–1345.
- Takahashi K, Kang JH, Ohnishi S, Hino K, Miyakawa H, Miyakawa Y, Maekubo H, Mishiro S. 2003a. Full-length sequences of six hepatitis E virus isolates of genotypes III and IV from patients with sporadic acute or fulminant hepatitis in Japan. *Intervirology* 46:308–318.
- Takahashi M, Nishizawa T, Miyajima H, Gotanda Y, Iita T, Tsuda F, Okamoto H. 2003b. Swine hepatitis E virus strains in Japan form four phylogenetic clusters comparable with those of Japanese isolates of human hepatitis E virus. *J Gen Virol* 84:851–862.
- Takahashi K, Kitajima N, Abe N, Mishiro S. 2004. Complete or near-complete nucleotide sequences of hepatitis E virus genome recovered from a wild boar, a deer, and four patients who ate the deer. *Virology* 330:501–505.
- Takahashi M, Kusakai S, Mizuo H, Suzuki K, Fujimura K, Masuko K, Sugai Y, Aikawa T, Nishizawa T, Okamoto H. 2005. Simultaneous detection of immunoglobulin A (IgA) and IgM antibodies against hepatitis E virus (HEV) is highly specific for diagnosis of acute HEV infection. *J Clin Microbiol* 43:49–56.
- Tam AW, Smith MM, Guerra ME, Huang C, Bradley DW, Fry KE, Reyes GR. 1991. Hepatitis E virus (HEV): Molecular cloning and sequence of the full-length viral genome. *Virology* 185:120–130.
- Tanaka E, Takeda N, Li TC, Orii K, Ichijo T, Matsumoto A, Yoshizawa K, Iijima T, Takayama T, Miyamura T, Kiyosawa K. 2001. Seroepidemiological study of hepatitis E virus infection in Japan using a newly developed antibody assay. *J Gastroenterol* 36:317–321.
- Tanaka E, Matsumoto A, Takeda N, Li TC, Umemura T, Yoshizawa K, Miyakawa Y, Miyamura T, Kiyosawa K. 2005. Age-specific antibody to hepatitis E virus has remained constant during the past 20 years in Japan. *J Viral Hepat* 12:439–442.
- Tei S, Kitajima N, Takahashi K, Mishiro S. 2003. Zoonotic transmission of hepatitis E virus from deer to human beings. *Lancet* 362:371–373.
- Thompson JD, Higgins DG, Gibson TJ. 1994. CLUSTAL W: Improving the sensitivity of progressive multiple sequence alignment through sequence weighting, position-specific gap penalties and weight matrix choice. *Nucl Acids Res* 22:4673–4680.
- Velazquez O, Stetler HC, Avila C, Ornelas G, Alvarez C, Hadler SC, Bradley DW, Sepulveda J. 1990. Epidemic transmission of enterically transmitted non-A, non-B hepatitis in Mexico, 1986–1987. *JAMA* 263:3281–3285.
- Waar K, Herremans MM, Vennema H, Koopmans MP, Benne CA. 2005. Hepatitis E is a cause of unexplained hepatitis in The Netherlands. *J Clin Virol* 33:145–149.
- Yazaki Y, Mizuo H, Takahashi M, Nishizawa T, Sasaki N, Gotanda Y, Okamoto H. 2003. Sporadic acute or fulminant hepatitis E in Hokkaido, Japan, may be food-borne, as suggested by the presence of hepatitis E virus in pig liver as food. *J Gen Virol* 84:2351–2357.
- Zafrullah M, Ozdener MH, Oanda SK, Jameel S. 1997. The ORF3 protein of hepatitis E virus is a phosphoprotein that associates with cytoskeleton. *J Virol* 71:9045–9053.

医薬品
 医薬部外品 研究報告 調査報告書
 化粧品

識別番号・報告回数		報告日		第一報入手日 2007年5月7日	新医薬品等の区分 該当なし	厚生労働省処理欄
一般的名称	人ハプトグロビン	研究報告の 公表状況	Medical Molecular Morphology 2007; 40 (1): 23-28		公表国 日本	
販売名 (企業名)	ハプトグロビン注-ヨシトミ(ベネシス)					
研究報告の概要	<p>GBV-C 及び HGV は、非 B 非 C 型肝炎の原因として関与する新たなウイルスとして紹介されたが、これらウイルス粒子の形態は依然不明であるとともに、非 A-E 肝炎の殆どの症例は、それらの感染と関係がない。我々は、ALT が高く、HCV 抗体と B 型肝炎表面抗原が陰性であるドナーからの血漿サンプル中のウイルス様粒子を視覚的に捉えようと試み、このウイルス様粒子と非経口的に感染する GBV-C/HGV の遺伝子との関係を調査した。</p> <p>HBV 及び HCV どちらにも感染していない高 ALT レベルの血漿 13 検体及び正常 ALT レベルの血漿 10 検体の計 23 検体を 40-60% ショ糖密度勾配遠心分離法によって分離し、ウイルス様粒子 (VLPs) を電子顕微鏡で観察した。血漿中の GBV-C/HGV RNA を検査した。ウイルス様粒子は、高 ALT レベルの血漿 13 検体のうち 12 検体 (92.3%) 及び正常対照血漿 10 検体のうち 1 検体 (10%) の 1.15-1.16g/mL の密度のフラクションから見つかった。VLPs の超微細構造形態は、大きさ、外観とも多形性であった (大部分の VLPs は 35-45nm の内核と表面に長さ 9-12nm のスパイク様突起を有する 50-80nm の球状粒子)。直径 50-70nm、長さ 110-160nm の棒状の VLPs も同じ検体で観察された。循環血液中の VLPs の検出率は、有意に高 ALT レベルと関係 (P<0.001) していたが、VLPs を含む血漿のいずれにも、GBV-C/HGV RNA は検出されなかった。HBV 及び HCV ともに陰性の血漿中に高 ALT との有意な関係が認められた球状の VLPs が確認され、それらが非 B 非 C 型肝炎に関係していることが示唆された。</p>					使用上の注意記載状況・その他参考事項等
	報告企業の意見					今後の対応
<p>非 B 非 C 型肝炎との関連が示唆されるウイルス様粒子を発見したとの報告である。</p> <p>非 B 非 C 型肝炎との関連が示唆されるウイルス様粒子は、GBV-C/HGV と関連していないこと以外の性状等に関する情報は不足しているため、現時点において安全性評価は困難であり、今後も情報収集に努める。</p>					<p>ウイルス様粒子に関する追加情報の入手に努める。</p>	

使用上の注意記載状況・その他参考事項等

2. 重要な基本的注意

(1) 本剤の原材料となる献血者の血液については、HBs 抗原、抗 HCV 抗体、抗 HIV-1 抗体、抗 HIV-2 抗体、抗 HTLV-I 抗体陰性で、かつ ALT (GPT) 値でスクリーニングを実施している。更に、プールした試験血漿については、HIV-1、HBV 及び HCV について核酸増幅検査 (NAT) を実施し、適合した血漿を本剤の製造に使用しているが、当該 NAT の検出限界以下のウイルスが混入している可能性が常に存在する。本剤は、以上の検査に適合した血漿を原料として、Cohn の低温エタノール分画で得た画分から人ハプトグロビンを濃縮・精製した製剤であり、ウイルス不活化・除去を目的として、製造工程において 60℃、10 時間の液状加熱処理及び濾過膜処理 (ナノフィルトレーション) を施しているが、投与に際しては、次の点に十分注意すること。

3

4

ORIGINAL PAPER

Masahiko Kaito · Hideaki Tanaka · Shinichiro Horiike
Naoki Fujita · Motoh Iwasa · Yoshinao Kobayashi
Esteban Cesar Gabazza · Yukihiko Adachi
Masayoshi Konishi · Shozo Watanabe



Unidentified virus-like particles are detected in plasmas with elevated ALT levels: are they significant of etiological agent(s) of non-B, non-C hepatitis?

Received: May 9, 2006 / Accepted: June 22, 2006

Abstract GB virus C (GBV-C) and hepatitis G virus (HGV) have been proposed as new viruses etiologically implicated in non-B, non-C hepatitis, but the morphology of these particular virus particles is still unknown, and most cases of non-A to E hepatitis do not relate to their infections. We tried to visualize virus-like particles (VLPs) in plasma samples from hepatitis B surface antigen- and antibody to hepatitis C virus (HCV)-negative blood donors with elevated alanine aminotransferase (ALT), and examined the association of the virus-like particles and the genomes of parenterally transmissible GBV-C/HGV. Twenty-three plasma samples, 13 with elevated ALT levels and 10 with normal ALT values, from blood donors without infections of hepatitis B virus (HBV) and HCV, were subjected to a 20%–60% sucrose density gradient centrifugation, and virus-like particles were observed by electron microscopy. GBV-C/HGV RNAs in the plasmas were tested. Virus-like particles were found in the fractions with densities of 1.15–1.16 g/ml from 12 of 13 (92.3%) plasmas with elevated ALT levels and 1 of 10 (10%) normal controls. The ultrastructural morphology of visualized VLPs was pleomorphic in size and appearance; the majority of the VLPs were 50- to 80-nm spherical particles with a 35- to 45-nm inner core and 9- to 12-nm-long surface spikelike projections. Rodlike VLPs 50–70 nm in diameter with a length of 110–160 nm were also observed in the same samples. The incidence of detection of the circulating VLPs was significantly ($P < 0.001$) related to elevated ALT levels, but GBV-C/HGV RNAs were detected in none of the plasmas containing the virus-like particles. Spherical VLPs are detected in HBV- and HCV-

negative plasmas significantly correlated with the elevation of ALT, suggesting that they are implicated in non-B, non-C hepatitis.

Key words Non-B, non-C hepatitis · Virus-like particle · Hepatitis C virus · GB virus C · Electron microscopy

Introduction

The genomes of hepatitis C virus (HCV), GB virus C (GBV-C), and hepatitis G virus (HGV) have been successfully cloned without isolation of the virus particles.^{1–3} The causative role of HCV in bloodborne acute and chronic hepatitis has been well established. Although GBV-C/HGV can be transmitted parenterally, most of their infections are not associated with acute and chronic non-B, non-C hepatitis, and there is some doubt whether GBV-C/HGV replicates in the liver and causes hepatitis.^{4,5} During immunogold electron microscopy^{6–9} of HBV and HCV particles, we have noticed that some plasma samples from blood donors with elevated alanine aminotransferase (ALT) levels contained virus-like particles (VLPs) that did not react positively with antibodies specific to the HCV envelope protein. The VLPs, thus immunologically distinguished from HCV virion, were also different in morphology from the latter, and their etiological implications were not clear. In this article, we have tried to visualize VLPs in hepatitis B surface antigen (HBsAg)- and anti-HCV-negative plasmas and evaluated whether the circulating VLPs were detected with or without a relationship to elevated plasma ALT levels or to GBV-C/HGV RNAs.

Materials and methods

Twenty-three blood donor plasma samples were the subjects in this study. Clinical characteristics of the blood donors are summarized in Table 1. These samples were negative for HBsAg (AUSRIA II-125; Dainabot, Tokyo,

M. Kaito (✉) · H. Tanaka · S. Horiike · N. Fujita · M. Iwasa · Y. Kobayashi · E.C. Gabazza · Y. Adachi
Department of Gastroenterology and Hepatology, Division of Clinical Medicine and Biomedical Science, Institute of Medical Science, Mie University Graduate School of Medicine, 2-174 Edobashi, Tsu, Mie 514-8507, Japan
Tel. +81-59-232-1111; Fax +81-59-231-5223
e-mail: kaitoma@clin.medic.mie-u.ac.jp

M. Konishi · S. Watanabe
Health Administration Center, Mie University, Mie, Japan

Table 1. Summarized data of detection of virus-like particles and nucleic acids of known hepatitis viruses in plasma samples negative for HBsAg and anti-HCV

Plasma no.	ALT (IU/l)	Virus-like particles ^a		HBV DNA ^b	HCV RNA ^c	GBV-C RNA ^d	HGV RNA ^e
		Spherical	Rod-like				
1	91	-	-	-	-	-	-
2	81	2+	-	-	-	ND	ND
3	87	2+	-	-	-	-	-
4	70	3+	2+	-	-	-	-
5	82	3+	2+	-	-	-	-
6	103	3+	1+	-	-	-	-
7	121	3+	3+	-	-	-	-
8	67	2+	1+	-	-	-	-
9	117	2+	-	-	-	-	-
10	72	2+	1+	-	-	-	-
11	103	2+	-	-	-	-	-
12	78	3+	1+	-	-	ND	ND
13	77	1+	-	-	-	-	-
14	4	1+	-	-	-	-	-
15	28	-	-	-	-	-	-
16	6	-	-	-	-	-	-
17	10	-	-	-	-	-	-
18	7	-	-	-	-	-	-
19	11	-	-	-	-	-	-
20	4	-	-	-	-	-	-
21	6	-	-	-	-	-	-
22	6	-	-	-	-	-	-
23	8	-	-	-	-	-	-

ALT, alanine aminotransferase (normal < 35 IU/L); ND; not dated

^aCirculating virus-like particles (VLPs) in a 5-square section of a 300-mesh grid was photographed under electron microscopy. The mean numbers of VLPs in one square was indicated as - (negative), 1+ (0 < + < 10), 2+ (10 ≤ 2+ < 100), and 3+ (100 ≤ 3+)

^bHBV DNA was tested using a Quantiplex HBV-DNA assay kit

^cHCV RNA was tested using an AmpliCor HCV kit

^dGBV-C and ^eHGV RNAs were assayed as described in the text

Japan), antibody to HBV core antigen (CORAB; Dainabot), HBV DNA (Quantiplex HBV-DNA Assay; Chiron, Emeryville, CA, USA), anti-HCV (Ortho HCV Ab IRMA Test III; Ortho-Clinical Diagnostics, Tokyo, Japan), HCV RNA (AmpliCor HCV; Roche Molecular Systems, Branchburg, NJ, USA), and antibodies to human T-cell leukemia virus type I (Determiner HTLV-I Antibody; Kyowa Medex, Tokyo, Japan) and human immunodeficiency virus (GENELAVIA MIXT; Sanofi Diagnostics Pasteur, Marnes la Coquette, France). Thirteen samples were elevated in alanine aminotransferase (ALT) levels (≥35 IU/l), and 10 samples were within a normal level in ALT (<35 IU/l). Total cholesterol and triglyceride levels were normal in all plasma samples.

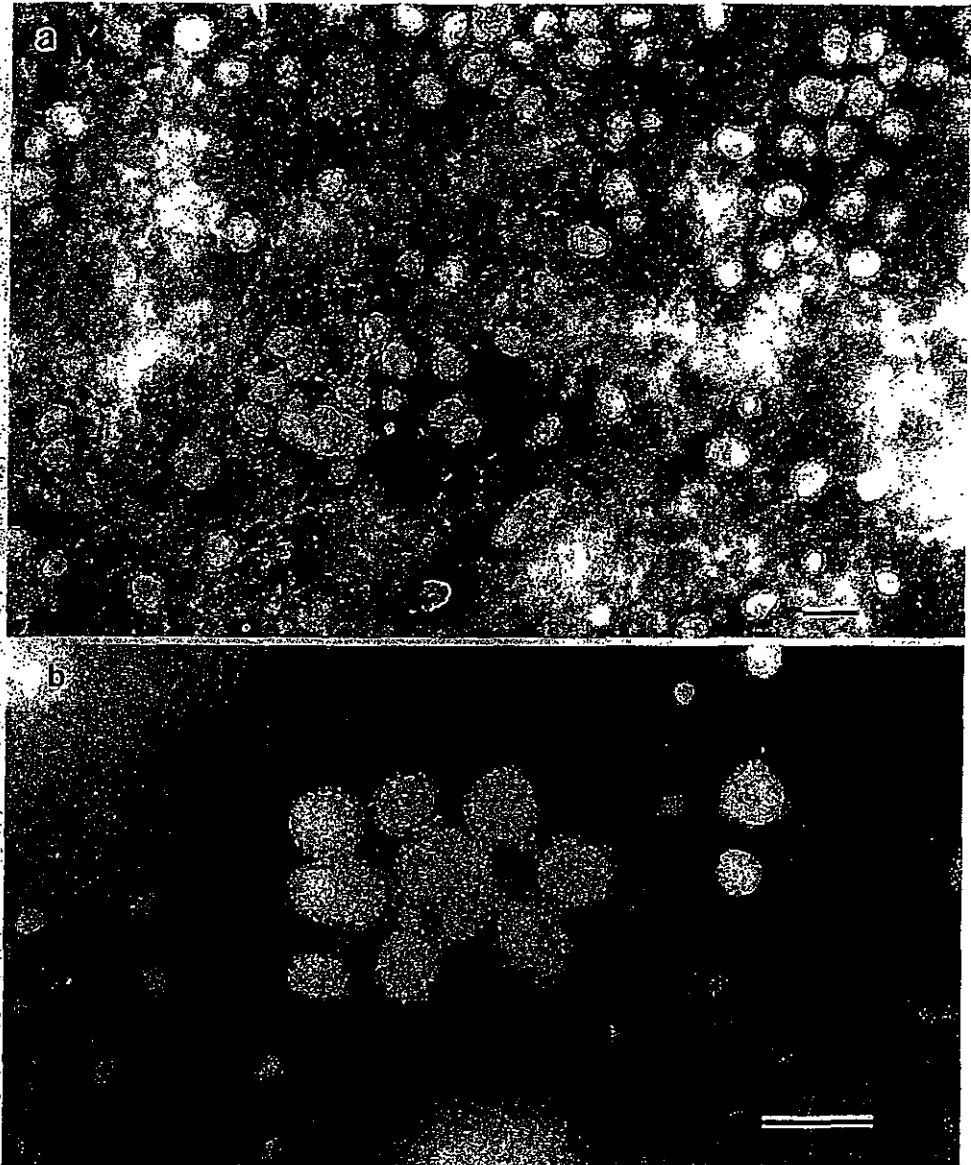
One hundred milliliters of each plasma sample was diluted with two volumes of TEN [100mM Tris-HCl, pH 8.0, 1mM ethylenediaminetetraacetic acid (EDTA), 100mM NaCl] and centrifuged at 75000g for 6h at 4°C. A suspension of the pellet in TEN was centrifuged again at 150000g for 2.5h at 4°C. An approximately 1000-fold-concentrated suspension of the sample in TEN was layered on a 20%–60% (W/W) linear sucrose density gradient in TE (100mM Tris-HCl, pH 8.0, 1mM EDTA) and centrifuged at 100000g for 16h at 4°C, which was followed by fractionation from the bottom of the tube. After measurement of the sucrose concentration of every fraction, each sucrose fraction was

diluted in phosphate-buffered saline (PBS, pH 7.4) and then centrifuged at 150000g for 2.5h at 4°C. The resulting pellet was suspended in 100ml PBS, equivalent to 1:1000 of the original plasma volume, for observing VLPs, and stored at -80°C until use. Two to three microliters of the concentrated specimen was mounted on a Formvar-coated and carbon-vaporized copper grid and examined under a Hitachi H-800 electron microscope operated at 100kV after staining with 2% phosphotungstic acid (pH 6.5). To observe the nucleic acid-containing virus core structure, the specimen was subjected to repeated freezing and thawing several times, and then stained with 0.2% uranyl acetate (pH 4.4).

The GBV-C genome was assayed by reverse transcription-nested polymerase chain reaction (PCR) using primers derived from the NS3/helicase region of GBV-C genome reported by Simons et al.,² and this assay was performed as described elsewhere.¹⁰ The genome of HGV, provisionally designated by Linnen et al.,³ was assayed by using the kit of Boehringer Mannheim (Mannheim, Germany), which contained one primer pair for the NS5A region and a second primer pair for the 5'-untranslated region as well as the corresponding capture probes, according to the manufacturer's instructions.

Fisher's exact probability test was used to assess the significance of differences between the sample groups with or without elevated ALT levels.

Fig. 1. Electron micrograph of negatively stained 50- to 80-nm spherical virus-like particles (VLPs) in hepatitis B surface antigen (HBsAg)- and anti-hepatitis C virus (HCV)-negative plasma with an elevated alanine aminotransferase (ALT) level. VLPs from sample no. 6 (a) and sample no. 3 (b) are shown. Bar 100 nm.



Results

Spherical VLPs were detected in 12 of 13 (92.3%) plasma samples with elevated ALT levels and 1 of 10 (10%) plasma samples with normal ALT values. The incidence of detection of the virus-like particles was significantly ($P < 0.001$) related to the elevation of plasma ALT levels (see Table 1).

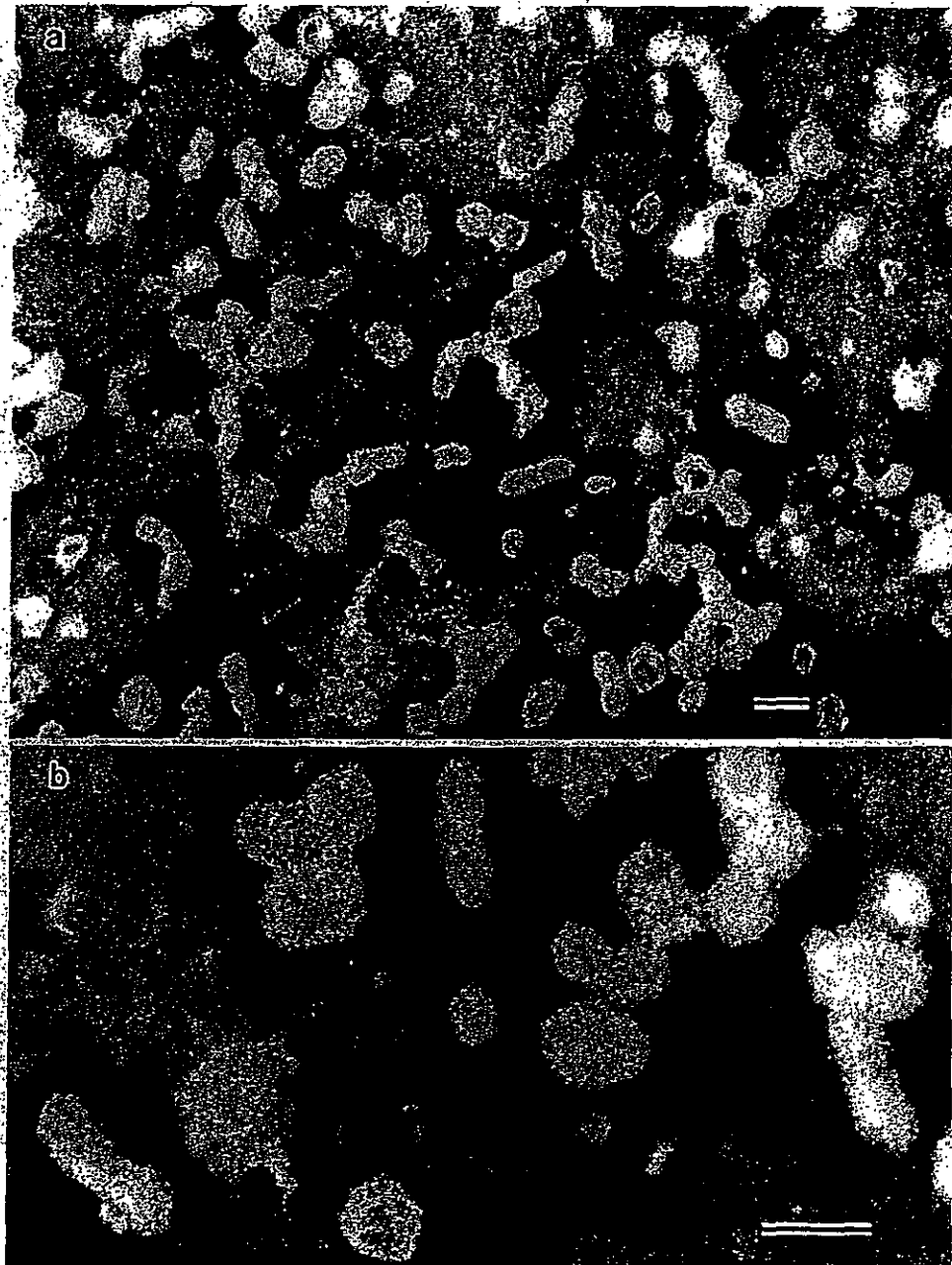
Most the visualized VLPs were spherical particles 50–80 nm in diameter with 9- to 12-nm-long surface spikelike projections (Fig. 1a,b). Some VLPs have a large diameter, more than 100 nm (Fig. 1a). In some specimens were detected more than a moderate number of VLPs, and rodlike VLPs 50–60 nm in diameter with a length of 110–160 nm were also detected that had surface projections similar to those of the 50- to 80-nm spherical particles (Fig. 2). The

diameter of the rodlike VLPs was 50–70 nm and their length was 110–160 nm. In addition, a 35- to 45-nm corelike structure containing the electron-dense material within a spherical VLP (Fig. 3a,b) and two inner cores of a rodlike VLP (Fig. 3c) were visualized by positive staining with uranyl acetate. These VLPs were found to be constantly banded in a sucrose density gradient at around 1.15–1.16 g/ml (range, 1.12–1.18 g/ml). GBV-C/HGV RNAs were not detected in any of the plasma samples (see Table 1).

Discussion

We found VLPs in ALT-elevated plasma samples from blood donors without infections of HBV and HCV that were consistently banded at around 1.15–1.16 g/ml by

Fig. 2. Negatively stained electron micrograph of rodlike VLPs presented in plasma containing a moderate number of the spherical VLPs. Rodlike VLPs from sample no. 7 (a) and sample no. 5 (b) are shown. Bar 100nm



sucrose density gradient centrifugation. The majority were 50- to 80-nm spherical particles with 9- to 12-nm-long surface projections (see Fig. 1), and were, if anything, morphologically resembling togaviruses or coronaviruses. Positive staining revealed that the electron-dense material combined with uranium existed in a 35- to 45-nm internal core structure of the spherical virus-like particle, and indicated that the spherical particles had the nucleic acid therein (see Fig. 3). In addition, rodlike VLPs (Fig. 2) were observed concomitantly in the specimens containing many spherical VLPs, and the surface spikelike projections of rodlike forms looked similar to those of the spherical particles. Interestingly, two internal cores (Fig. 3c) were detected in a rodlike form, suggesting that this form could be a diploid particle

of the virus. Thus, we might consider that both the spherical and the rodlike virus-like particles belong to the same virus species.

In the past three decades, togavirus-like particles have been detected in acute-phase serum from a hemodialysed patient with non-A, non-B hepatitis and in the acute-phase urine of two icteric non-A, non-B hepatitis cases,¹¹ or in the liver of a patient with sporadic non-A, non-B fulminant hepatitis,¹² but the number of those objects was too small, and thus the etiological implications were not developed. Circulating VLPs, as presented here, were highly prevalent (92.3%) in ALT-elevated plasma samples, and were relatively easy to visualize by conventional electron microscopy, whereas in only one of ten normal controls were a few