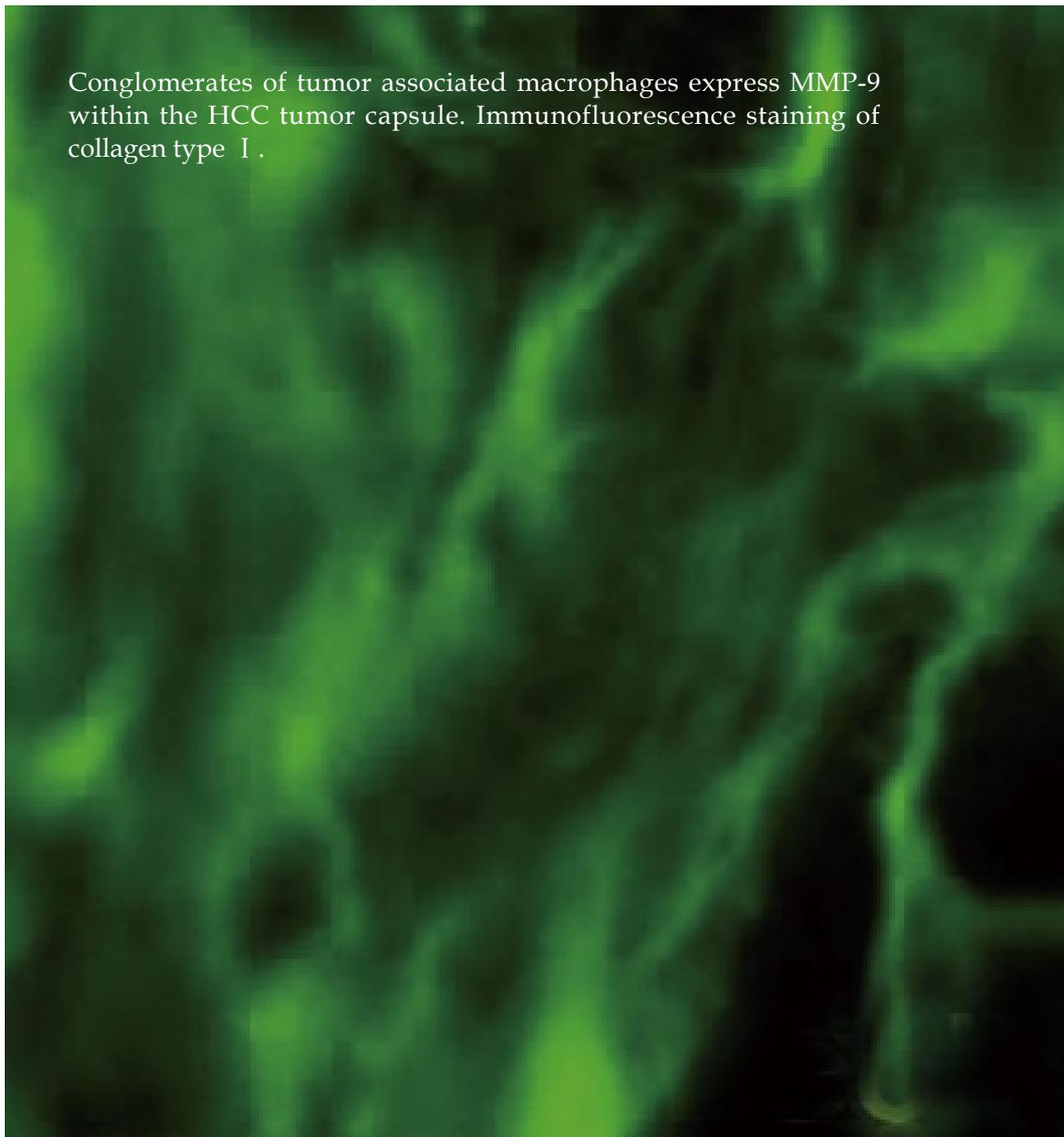


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Hepatocellular carcinoma in thalassemia: A critical review

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Abstract

Due to blood transfusions, thalassemics are often infected with either hepatitis C virus (HCV) or hepatitis B virus and often have hemochromatosis. Hepatocellular carcinoma (HCC) has emerged in thalassemics only recently as a result of the improvement in thalassemia outcomes. In fact, a prospective study estimated an HCC incidence in β -thalassemia of about 2%. Although data are scanty, HCC screening in thalassemics with risk factors for HCC should be carried out. HCV treatments have some efficacy in HCV infected thalassemics despite partial contraindication to ribavirin and iron overload. However, there are no data on how HCV treatment translates into HCC prevention. Preliminary data suggest that HCC treatment in thalassemics should generally have the same outcomes as in non-thalassemics. Although coexistence of severe comorbidities makes liver transplantation challenging, this therapeutic possibility should not be precluded for well selected HCC β -thalassemia patients. In fact, 2 transfusion dependent adult HCC β -thalassemia patients have recently undergone successful liver transplantation with a good outcome. In conclusion, HCC seems to be a developing issue in thalassemia and HCC screening should be carried out. HCC treatment, including liver transplantation, can be performed in selected patients. A multidisciplinary effort is needed for management.

Key words: Thalassemia; Hepatocellular carcinoma; Hemochromatosis; Screening; Complication; Liver transplantation

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INTRODUCTION

Thalassemias are rare inherited hemoglobin disorders resulting in chronic hemolytic anemia, endemic particularly in the Mediterranean Area and South-East Asia^[1]. Depending on severity, two clinical forms are distinguished: thalassemia major (TM), characterized by severe anemia starting during the first year of life, requiring life-long transfusion therapy for survival, and thalassemia intermedia (TI), characterized by later onset and generally milder anemia, permitting survival without regular transfusion therapy^[2].

Due to blood transfusions, many patients with β -thalassemia are infected with either hepatitis C virus (HCV) or hepatitis B virus (HBV), particularly those who were born before the 1990s^[3]. Moreover, most of the patients have hemochromatosis, which is the main cause of morbidity and mortality. In the past, HCC was not a known complication of thalassemia probably because many patients did not survive long enough to develop the condition and also because there was insufficient attention to the issue. In fact, many patients died at a young age, mainly due to heart failure. Recently, the outcome of thalassemia has improved and chronic organ damage, in

particular end-stage liver disease, has become a frequent complication. Better outcome is related to the improved treatment of iron overload with iron chelating drugs in the management of thalassemia complications, particularly heart disease, and in the prevention of infectious diseases secondary to blood transfusion^[4].

Hepatocellular carcinoma (HCC) is a complication of cirrhosis and it affects prognosis. Occasionally, HCC can develop on a cirrhosis-free liver if other risk factors are present. Nowadays, a range of well-established treatments exist that can improve survival of HCC patients^[5-13]. In this paper, the knowledge of HCC in thalassemia will be updated. This review deals with patients with both TM and TI and excludes other thalassemia syndromes, such as Sickle cell/beta thalassemia that are clinically similar to Sickle Cell Disease and quite different from TM and TI.

EPIDEMIOLOGY OF THALASSEMIA-ASSOCIATED HCC

Awareness of HCC as a clinical complication of thalassemia has developed in the last few years. In the past, many patients did not survive long enough to develop HCC. In fact, many patients died at a young age, mainly due to heart failure^[4,14]. Recently, the outcome of thalassemia has improved and chronic organ damage, in particular end-stage liver disease and HCC, has appeared as a frequent complication^[4]. In fact, a recent multicenter retrospective study on patients with different thalassemia syndromes, identified 22 cases of HCC in Italian patients, 19 of whom were affected by either TM or TI^[15]. Furthermore, a prospective study was conducted on HCC incidence in 108 thalassemia patients (38 with TM and 70 with TI; median age 36.8 yr) screened with liver ultrasound imaging. Seventy-two (31 TM, 41 TI) with one or more risk factors for HCC (iron overload in 72, HCV infection in 46, HBV infection in two, liver cirrhosis in 10) and 33 (4 TM, 29 TI) without risk factors were included in the study; three were excluded (1 with a previous diagnosis of HCC on Child-Pugh class C cirrhosis, and two patients who were under 18 years old). All patients with iron overload were treated with at least one iron chelating drug. Once a liver focal lesion was found at ultrasound, HCC diagnosis was confirmed following current guidelines^[13]. Overall, two of the 72 with at least one HCC risk factor were found to have a newly developed HCC, with an estimated HCC incidence in thalassemia of about 2%^[16].

Moreover, other new cases have been recently discovered, suggesting that HCC is becoming one of the leading clinical problem in thalassemia (unpublished data). Finally, if the future trend confirms an increase in the number of HCC in thalassemia, it would be reasonable to recommend HCC screening of all thalassemia patients with some risk factor since this could allow early treatment, leading to improved outcomes^[17,18].

HCC PREVENTION IN THALASSEMIA PATIENTS

Currently, there are no effective tools for preventing HCC in thalassemia. Since hepatic disease is multifactorial in thalassemia, the theoretical possibility of preventing HCC is based on the possibility of preventing or curing HCC risk factors, namely hemochromatosis and chronic viral hepatitis, mainly due to HCV.

Hemochromatosis is a known risk factor for HCC^[19,20]. Treatment of iron overload with iron chelating drugs is certainly the leading reason why thalassemia outcomes have recently improved dramatically. Since subcutaneous or intravenous deferoxamine was first shown to be an effective chelating agent^[4], new oral drugs have been developed which have proven as effective as deferoxamine with better compliance^[21]. Moreover, as recent evidence suggests, the use of multiple chelating drugs taken either concomitantly or sequentially seems likely to be the future route for iron overload treatment in thalassemia^[22]. However, while thalassemia outcomes have been improving in recent years mainly because of iron chelation, HCC has emerged as a new complication, suggesting perhaps that current chelating regimens alone are not effective enough to prevent HCC.

Treatment of HCV-related chronic hepatitis in thalassemia using different kinds of interferon with or without ribavirin has had variable results in a number of small trials. Overall, most of the studies reported some efficacy. However, in addition to the rather small number of patients treated, there remain concerns about the safety of ribavirin. In fact, ribavirin increases not only the number of responders but also increases the need for blood transfusion^[23-33].

Based on the information currently available, there is no evidence suggesting that HCV treatment can translate into HCC prevention in thalassemia. However, it is plausible that prevention of HCV infection due to blood transfusion will, in future, result in a reduction of HCC in thalassemia.

Potential treatments of thalassemia-associated HCC

Since survival of patients with thalassemia has dramatically improved in recent years^[4], thalassemia should not be considered a contraindication per se to HCC treatments that are effective in non-thalassemia patients^[5-13].

To date, there is little published evidence concerning the treatment of HCC in thalassemia. However, both percutaneous radio frequency thermoablation and ethanol injection^[16], surgical resection^[16,34] and chemoembolization (unpublished data) have been shown effective and safe for the treatment of HCC in selected patients with thalassemia. Recent determinations of the therapeutic efficacy of sorafenib showed some significant improvement in prognosis for earlier stage HCC^[7,8]. It is, as yet, unknown whether sorafenib will be beneficial for the HCC associated with thalassemia.

A special mention should be made of the possibility of

liver transplantation in thalassemia patients. In fact, choice of the best treatment for HCC in thalassemia remains controversial. In particular, the possibility of treating HCC or end-stage liver disease with liver transplantation has long been rejected because of organ shortage and because thalassemia is considered a contraindication. Only one combined heart and liver transplantation has been reported and the outcome is currently not known^[35]. The recent demonstration of improvements in thalassemia outcomes should prompt a reconsideration of this issue^[4]. It is no longer reasonable that thalassemia should be considered a contraindication to liver transplantation, provided there is not any significant co-morbidity, namely heart disease and severe pulmonary hypertension^[14]. In fact, encouraging results in this regard have been achieved in recent studies from the author's team, which showed successful liver transplantation in 2 transfusion dependent thalassemia patients at 2 different Liver Transplantation Units. Post-transplantation outcome has been satisfactory in both cases after 6 mo and 2 years follow-up evaluations, respectively. Of course, a multidisciplinary effort is needed for management.

CONCLUSION

Thalassemia is a rare disease in which the appearance of HCC as a complication is mainly the result of recently improved outcomes in developed countries. Preliminary data suggest an incidence of HCC in thalassemia of about 2%. However, since thalassemia is endemic in many under-developed countries where patients are probably not screened for HCC, it is possible that present knowledge of this issue represents only the tip of an iceberg.

Periodic liver Ultrasound HCC screening should probably be considered for thalassemia patients with risk factors for HCC.

Prevention of HCV infection through blood transfusion is nowadays the only known evidence-based means to prevent HCC in thalassemia.

HCC treatment in thalassemia patients should be the same as for non thalassaemic HCC patients. Although coexistence of severe comorbidities makes the role of liver transplantation challenging, this therapeutic possibility should not be precluded for well selected HCC thalassemia patients. Of course, a multidisciplinary effort is needed for management of transplantation patients.

Many of the considerations reported in this review are extrapolated from scanty data that surely lack comprehensive evidence and they are mainly the personal opinion of the author. Multicenter international studies should be performed to strengthen these data.

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Innovative immunohistochemistry identifies MMP-9 expressing macrophages at the invasive front of murine HCC

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Abstract

AIM: To investigate the proteolytic contribution of tumor-associated macrophages (TAM) in tumor invasion, we analyzed whether TAM at the invasive front of small HCC in *Abcb4*^{-/-} mice show an enhanced expression of MMP-9.

METHODS: Liver cryosections of the hepatocellular carcinoma (HCC) invasive front from 12 mo old *Abcb4*^{-/-}

mice were stained for collagen type I and MMP-9 using Alexa488 and Alexa568 labeled secondary antibodies. Afterwards, the Alexa568 dye was bleached and the macrophage marker F4/80 was visualized using Alexa568 labeled secondary antibodies. Finally, photographs of the invasive tumor front were digitally overlaid and analyzed.

RESULTS: After complete bleaching of the primary dye, specific fluorescence staining of a third antigen, here F4/80, was successfully performed on the same histological section. With this method, we were able to identify conglomerates of matrix metalloproteinase (MMP-9) expressing macrophages within the tumor capsule of HCC.

CONCLUSION: MMP-9 expressing macrophages are involved in matrix remodelling at the invasive tumor front of HCC. The described staining protocol provides a simple yet powerful extension of conventional immuno-histochemistry, facilitating visualization of at least three different antigens plus nuclei in one single histological section.

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Key words: Fluorescence staining; Hepatocellular carcinoma; Matrix metalloproteinase; Tumor associated macrophages

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INTRODUCTION

Abcb4 knockout mice (formerly called *Mdr2*-knockout mice) lack the liver-specific P-glycoprotein responsible for phosphatidylcholine transport across the canalicular membrane. The absence of phospholipids from biliary fluid in *Abcb4*^{-/-}-mice results in bile regurgitation and portal inflammation followed by development of dysplasia and hepatocellular carcinoma (HCC)^[1,2]. Therefore, *Abcb4*^{-/-}-mice represent an ideal animal model mimicking cholangitis-associated carcinogenesis^[1,2].

The initial step in carcinogenesis (i.e. invasive and metastatic cell behavior) is the proteolytic destruction of the extracellular matrix (ECM), including the basement membrane. Numerous studies have demonstrated the pivotal role of matrix metalloproteinases (MMPs) for tumor associated ECM degradation^[3-5]. In particular, the gelatinases MMP-2 and MMP-9 have gained considerable attention: in various studies on tumor invasion and metastasis. MMP-2 and MMP-9 were shown to be able to cleave collagen type IV, the main component of the basement membrane^[6-8].

Previous studies demonstrated abundant expression of MMP-9 in tumor cells of HCC^[6,7]. Furthermore, MMP-9 expression was associated with growth and invasiveness of HCC^[6,9,10]. Apart from the malignant cells themselves, growing attention is being paid to the tumor microenvironment as a mediator of invasive and metastatic behaviour^[11,12]. Large quantities of tumor-associated macrophages (TAM) are associated with poor prognosis in various types of cancer, thereby suggesting relevance of these cells for tumor progression^[13]. A number of studies were able to prove the expression of a broad range of tumor promoting factors including MMP-9 by which TAM may drive tumor angiogenesis^[12,14,15].

Against this background, our aim was to analyze the expression patterns and cellular sources of MMP-9 within the tumor microenvironment of HCCs related to TAM in a mouse model of hepatocarcinogenesis. We developed a modification of the well-known and traditionally conducted immunohistochemistry. This allows the parallel visualization of at least three antigens within one histological section. The described approach may represent a powerful advancement in the analytical capabilities of immunohistochemistry.

MATERIALS AND METHODS

Animals and tissue preparation

FVB/N-*Abcb4*^{tm1bor} gene-targeted mice were crossed back towards the fibrosis-susceptible BALB/cJ strain for ten generations as characterized recently^[16,17]. Mice were killed at the age of twelve months and tissue samples of HCCs were fixed in 1% neutral buffered formalin for 16 h and afterwards embedded in Tissue-Tek (Sakura, Zoeterwoude, Netherlands) for cryopreservation at -80°C. The present study was performed with permission of

the State of Hessen, regional council Giessen, according to section 8 of the German Law for the Protection of Animals and conforms to the *Guide for the Care and Use of Laboratory Animals* (Az: V54-19c20/15cGI20/10).

Immunofluorescence

3 µm frozen sections were blocked for 30 min with 5% bovine serum albumin, 2% goat serum (Biomedica, Foster City, USA) and 0.1% cold fish skin gelatine (Sigma-Aldrich) in PBS with 0.1% Triton (Roth, Karlsruhe, Germany) and 0.05% Tween 20 (Serva, Heidelberg, Germany). Sections were immunostained with goat anti-mouse MMP-9 antibodies, Alexa 568-conjugated donkey anti-goat IgG and afterwards with rabbit anti-mouse collagen type I antibodies and Alexa 488-conjugated goat anti-rabbit IgG. DAPI (4',6-diamidino-2-phenylindole dihydrochloride, Sigma) was used for nucleus staining. The stained probes were covered with Dako Cytomation mounting medium (Glostrup, Denmark) and glass cover slips for microscopic analysis. Fluorescence images of the tumor invasive front were obtained under a fluorescence microscope (Leica DMRB, Wetzlar, Germany; camera: Nikon Coolpix 5400, Düsseldorf, Germany). Afterwards, the Alexa568 dye was completely bleached by illumination with green light under the microscope for 2 h. Immunostaining of macrophage antigens was performed with rat anti-F4/80 primary and goat anti-rat Alexa568 secondary antibodies. The carefully adjusted selection of antibodies derived from different host-species was crucial to avoid unintended cross-stainings. Finally, photomicrographs of the invasive tumor front were taken and analyzed digitally using Photoshop software version 9.0.2. (Adobe, München, Germany).

Primary antibodies were diluted 1:50 and secondary antibodies 1:1000 prior to use. Specificity of immunohistological stainings was verified by the use of isotype control-IgGs instead of primary antibodies. MMP-9 IgGs were purchased from R&D-Systems (#AF909, Wiesbaden, Germany), collagen type I IgGs from Biodesign (#T40777R, Freiburg, Germany), and F4/80 IgGs from Dianova (#T-2006, Hamburg, Germany). Alexa 468- and Alexa 488-conjugated secondary antibodies were purchased from Molecular Probes (Eugene, USA).

RESULTS

Within one year all BALB/c-*Abcb4*^{-/-}-mice (*n* = 17) had developed an HCC. On average 7 ± 4 tumors were detected per mouse, with a mean tumor size of 4.5 ± 2.5 mm. Figure 1 illustrates the macroscopic and histological aspect of HCC in *Abcb4*^{-/-}-mice.

Following the modifications of conventional immunohistochemistry as described above, we successfully identified three antigens (MMP-9, collagen I and F4/80) plus nuclei within the same histological section (Figure 2).

Immunofluorescence co-staining of collagen type I and MMP-9 visualized conglomerates of MMP-9 expressing cells within the tumor capsule of HCC. After photobleaching of the red fluorescent dye, staining with

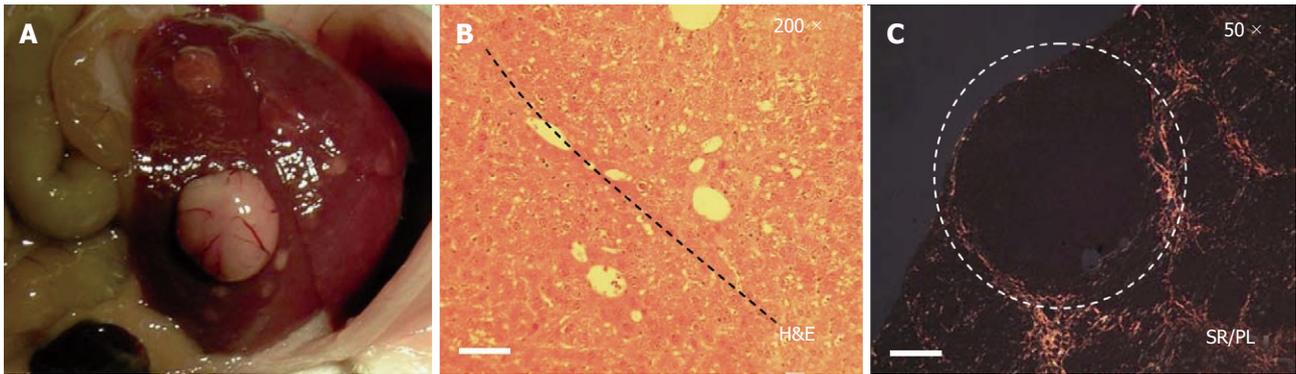


Figure 1 Macroscopic and histological aspect of HCC in BALB/c-*Abcb4*^{-/-} mice. A: Representative macroscopic view of BALB/c-*Abcb4*^{-/-} mouse liver at one year of age. Typically, one prominent tumor and a number of smaller tumors are visible; B: Hematoxylin and Eosin (H&E) stained liver section of the invasive tumor front (tumor on the upper right, fibrotic parenchyma on the lower left side, dashed line: HCC tumor capsule, bar represents 100 μ m); C: Fifty-fold enlarged Sirius red (SR) stained liver section photographed under polarized light (PL, bar represents 400 μ m). Fibrillar collagens appear in red. The tumor capsule is rich in collagen, whereas tumor stroma appears collagen-free.

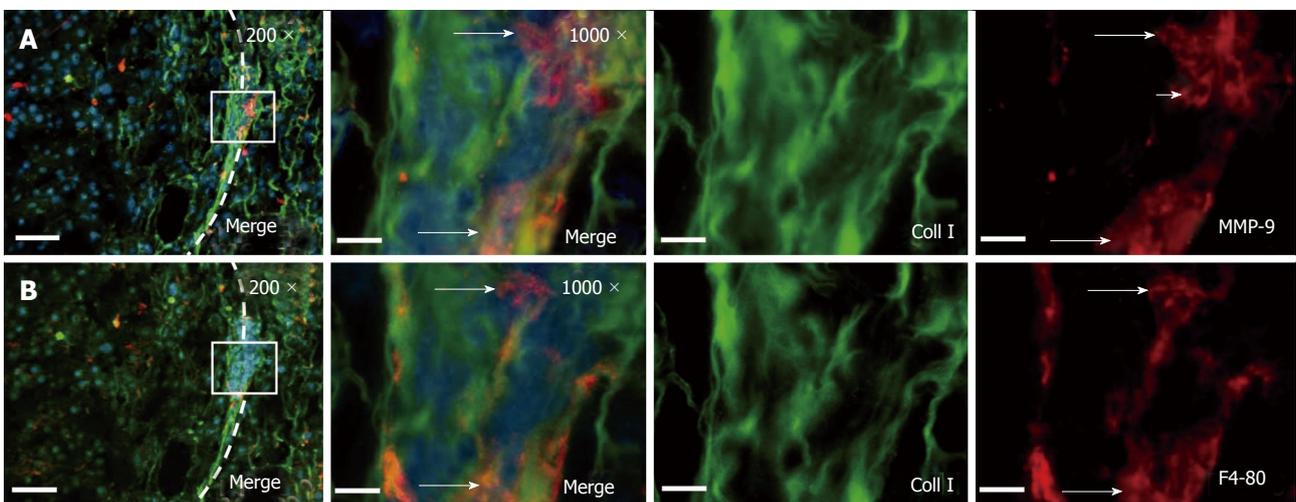


Figure 2 Conglomerates of tumor associated macrophages express MMP-9 within the HCC tumor capsule. A: Immunofluorescence co-staining of collagen type I (green) and MMP-9 (red) displayed clusters of MMP-9 expressing cells in the tumor capsule. B: After photobleaching of the red fluorescence dye, macrophage marker F4/80 was visualized. The long arrows indicate MMP-9 positive macrophages. The shot arrow indicates an MMP-9 expressing cell which is negative for F4/80. DAPI was used for nucleus staining. The left panels provide a section overview with 200-fold magnification and bars represent 100 μ m (left side of the panel: Tumor; dashed line: Tumor capsule; box: This is enlarged to 1000 \times magnification in the corresponding panel, bars represent 20 μ m). Original green and red channel micrographs (1000 \times) were shown. Images were derived by conventional wide field fluorescence microscopy. Representative digital overlays are shown.

macrophage marker F4/80 identified macrophages to be the main source of MMP-9 expressing cells at the invasive tumor front. Representative results of the modified immunohistochemistry are shown in Figure 2.

DISCUSSION

Previous studies reported an enhanced expression of MMP-9 in HCC by the tumor cells themselves^[6,7]. Furthermore, MMP-9 expression was associated with growth and invasiveness of HCC^[6,9,10]. However, growing evidence exists that, apart from malignant cells, the tumor environment may play a pivotal role with regard to invasive and metastatic cell behaviour^[11,12]. Accordingly, a number of *in vitro* and *in vivo* studies demonstrated that cancer cells were able to induce MMP expression in noncancerous cells thereby eventually promoting tumor

progression^[8,18-20]. Representing a part of the tumor microenvironment, TAM have been shown to be essential for both tumor growth and metastasis^[21-23]. A recent study by Tsagozis and coworkers in prostate cancer revealed that the expression of MMP-9 by TAM is necessary to maintain their tumor promoting phenotype^[24]. In murine studies on cervical cancer, Giraudo *et al*^[14] showed that enhanced expression of MMP-9 contributed to cervical carcinogenesis by promoting tumor angiogenesis.

Herein, we analyzed the expression patterns and cellular sources of MMP-9 at the invasive front of HCC with respect to tumor associated macrophages by means of immunohistochemistry. Traditionally, double immunofluorescence staining with green and red fluorescent dyes allows minimal overlapping fluorescence emission spectra. The application of the blue fluorescence channel for antigen visualization in wide field

fluorescence microscopy is not feasible yet, because of low fluorescence quantum yields of the correspondingly labeled secondary antibodies. Therefore, conventional immunohistochemistry was limited to the visualization of only two antigens plus nuclei within the same section.

To achieve within one histological section the combined illustration of the cellular sources of MMP-9 at the invasive front of HCC together with type I collagen representing the main component of fibrotic ECM the visualization of more than two antigens was essential.

We therefore developed a modification of conventional immunohistochemistry, facilitating the parallel visualization of three antigens using two fluorescent dyes. Essential for this approach is the complete wipeout of one dye (Alexa568) by green light. In contrast, green fluorescent Alexa488 dyes withstand the photobleaching without any loss of fluorescence intensity. Following the bleaching procedure, Alexa568 was re-used to tag the third antigen, the macrophage marker F4/80. Owing to the photo stability of Alexa488 towards green light, the defined region of interest was easily reproducible after the second staining.

Applying this approach, we have clearly visualized conglomerates of MMP-9 expressing macrophages within the capsule of HCCs. In accordance with data on TAM and MMP-9 expression for tumor invasion and angiogenesis in the literature, our results suggest an etiologic role for TAM and MMP-9 in the development of HCC^[12,14,15,21-24].

However, whether MMP-9 expression in TAM within the HCC capsule leads to increased tumor invasiveness or eventually promotes angiogenesis remains unclear.

In summary, the staining procedure described herein has been shown to be a simple, feasible yet powerful advancement of the widely used technique of immunohistochemistry. This may be of the utmost interest for those lacking access to spectral imaging via confocal laser microscopy^[25].

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COMMENTS

Background

The initial step in carcinogenesis (i.e. invasive and metastatic cell behavior) is the proteolytic destruction of the extracellular matrix (ECM), including the basement membrane. Numerous reports were able to demonstrate the pivotal role of matrix-metalloproteinases (MMPs) for tumor associated ECM degradation. In particular, the gelatinases MMP-2 and MMP-9 have gained considerable attention: in various studies on tumor invasion and metastasis MMP-2 and MMP-9 were able to cleave collagen type IV, the main component of the basement membrane. Previous studies demonstrated abundant expression of MMP-9 in tumor cells of hepatocellular carcinoma (HCC). Furthermore, MMP-9 expression was associated with growth and invasiveness of HCC. Apart from the malignant cells themselves, growing attention is paid to the tumor microenvironment as a mediator of invasive and metastatic behaviour. Large quantities of tumor-associated macrophages (TAM) are related to poor prognosis in various entities of cancer, thereby suggesting

relevance of these cells for tumor progression. A number of studies was able to prove the expression of a broad range of tumor promoting factors including MMP-9 by which TAM may drive tumor angiogenesis.

Research frontiers

It is unclear, whether MMP expression in TAM within the HCC may lead to increased tumor invasiveness or eventually promotes angiogenesis.

Innovations and breakthroughs

Representing a part of the tumor microenvironment, TAM have been shown to be essential for both, tumor growth and metastasis. A recent study by Tsagozis and coworkers in prostate cancer revealed that the expression of MMP-9 by TAM is necessary to maintain their tumor promoting phenotype. In murine studies on cervical cancer, Giraudo *et al.* showed that enhanced expression of MMP-9 contributed to cervical carcinogenesis by promoting tumor angiogenesis.

Applications

The staining procedure described herein proves to be a simple, feasible yet powerful advancement of the widely used technique of immunohistochemistry. This may be of utmost interest for those lacking access to spectral imaging via confocal laser microscopy. Applying this approach, we have clearly visualized conglomerates of MMP-9 expressing macrophages within the capsule of HCCs. In accordance with data on TAM and MMP-9 expression for tumor invasion and angiogenesis in the literature, our results suggest an etiologic role of TAM and MMP-9 for development of HCC.

Peer review

The rationale of this work is adequate. The methods employed clearly let the authors to reach the aims, and the results are appropriate to sustain a discussion closer to the results showing well knowledge of the subject. This work is simple, direct, and interesting, and brings awareness to the subject.

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Safety of pegylated interferon and ribavirin therapy for chronic hepatitis C in patients with sickle cell anemia

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Abstract

AIM: To evaluate the safety and efficacy of combined pegylated interferon and ribavirin for the treatment of chronic hepatitis C (HCV) in patients with sickle cell anemia (SCA).

METHODS: Fifty-two patients with SCA and HCV were treated over a period of 7 years from June 2002 to July 2009. Their medical records were reviewed for: age at treatment, sex, body mass index, Hb level at the start of therapy and on follow-up, hemoglobin electrophoresis, liver function tests, G6PD level, LDH, bilirubin, HCV-RNA viral load, HCV genotype, liver biopsy, duration of treatment, and side effects. All were treated with pegylated interferon and a standard dose of ribavirin. The treatment was continued for 24 wk for those with genotype 2 and 3 and for 48 wk for those with genotype 1 and 4.

RESULTS: Fifty-two patients (30 females and 22 males) were treated. Their mean age was 29.5 years (range 15-54 years). HCV genotype was determined in 48 and 15 had liver biopsy. Their mean pre-treatment HCV-RNA viral load was 986330 IU/mL (range 12762-3329282 IU/mL). The liver biopsy showed grade I in 6 and grade II in 9 and stage I in 13 and stage II in 2. Only 8 were receiving hydroxyurea at the time of treatment.

All tolerated the treatment well and none experienced a decrease in their Hb which required blood transfusion pre, during or after therapy. There were no hematological side effects attributable to ribavirin at the usual recommended dose. Thirty-seven (71.2%) achieved SVR at 6 mo after the end of treatment. The remaining 15 were non-responders. Two of them showed an ETR but had a relapse. The remaining 13 had a relatively significant HCV-RNA viral load with a mean HCV-RNA viral load of 1829741.2 IU/mL (900000-3329282 IU/mL) and eight of them had HCV genotype 1, four had HCV genotype 4, and one had HCV genotype 5.

CONCLUSION: Patients with SCA and HCV can be treated with pegylated interferon and ribavirin at the usual recommended dose. This is even so in those who are not receiving hydroxyurea. The treatment is safe and effective and the response rate is comparable to those without SCA.

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Key words: Sickle cell anemia; Chronic hepatitis C; Treatment

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INTRODUCTION

Sickle cell anemia (SCA) is one of the common hemo-

globinopathies in the world. In the Eastern Province of Saudi Arabia, SCA is common with a reported sickle cell trait frequency up to 25% and a sickle cell anemia frequency of about 2%^[1,2]. It is well known that SCA can affect any part of the body and one of the common organs to be affected is the hepatobiliary system. This can manifest in several different ways including cholelithiasis, choledocholithiasis, hepatic crisis, hepatic sequestration and cholestatic jaundice as well as transfusion related hepatitis B and C^[3-8]. The exact frequency of hepatitis B and C in patients with SCA in Saudi Arabia is not known but the average annual incidence of seropositivity per 100 000 population was 104.6 for HBV and 78.4 for HCV^[9]. An 18.2% prevalence of antibodies to HCV was reported among sickle cell patients in the Central region of Saudi Arabia^[10]. The life expectancy of patients with SCA has improved considerably, attributable to better understanding of the disease, improved medical and surgical management and the use of hydroxyurea^[11-13]. However, this will put SCA patients with chronic hepatitis B and C at risk of developing liver cirrhosis, hepatocellular carcinoma and liver failure if their hepatitis B and C are not treated. In spite of the markedly improved results in the treatment of chronic hepatitis C using the combined therapy of pegylated interferon and ribavirin, patients with SCA and chronic hepatitis C have not been considered suitable for such treatment because of the believed risk that ribavirin will induce hemolysis and severe anemia which can aggravate their already existing anemia^[14-16]. Recently however there have been three publications reporting success of such treatment in patients with SCA and HCV C although the number of patients reported was small^[17-20]. This report describes our experience in the management of 52 SCA patients with HCV using pegylated interferon and ribavirin at the usual recommended dose.

MATERIALS AND METHODS

In June 2002, we started treating SCA patients with HCV using pegylated interferon and ribavirin at Qatif Central Hospital, Saudi Arabia. This was approved by the ethical and research committee and an informed written consent was obtained from each patient explaining the type and duration of treatment as well as the possible side effects. This was a retrospective study and the following information was noted: age at treatment, sex, body mass index, Hb level at the start of treatment and on follow-up, hemoglobin electrophoresis, liver function tests, G6PD level, LDH, bilirubin, HCV-RNA viral load, HCV genotype, liver biopsy, duration of treatment, and side effects. HCV serology screening was done using a third generation ELISA test and confirmatory RIBA test. HCV-RNA was detected and assayed using an automated extraction system (Cobas Ampliprep). HCV Quantification and Detection was performed using Abbott Real Time M2000 RT instrument and the COBAS TaqMan HCV Test (RT-PCR technology) on the COBAS

AmpliPrep system (Roche) These assays detect and quantitate genotypes (1-6). An Internal Control is included in the assays to monitor any possible amplification inhibitors.

Prior to treatment, all patients had complete blood count, liver function tests, thyroid function tests, alpha fetoprotein, autoimmune profile, HBV and HIV screening and an abdominal ultrasound as part of their work-up. None of our patients had co-infection with HIV or HBV. The Hb level, LDH and bilirubin levels were monitored and checked initially, and during follow-up at 2 wk, 4 wk, 8 wk, 12 wk, 16 wk, 24 wk, 32 wk, 40 wk and 48 wk. The METAVIR fibrosis grading scale was used to grade and stage liver fibrosis^[21]. Treatment was initiated within two to three weeks post liver biopsy. Only 48 out of 52 patients were HCV genotyped as this facility was not available in our hospital at the beginning of the study. All our patients were treated with pegylated interferon alpha-2a (Pegasys 180 micrograms pre-filled syringes, Roche) 180 microgram subcutaneously, once per week or pegylated interferon alpha-2b (Peginterone, Schering-plough) 1 ug/kg subcutaneously, once per week and ribavirin. Ribavirin Rebetol (200 mg capsules, Schering-Plough) or Copegus 200 mg tablets, Roche) was used in a dose of 400 mg twice daily for those with HCV genotype 2 and 3 as well as those in whom HCV genotype was not known and 600 mg twice daily for those with HCV genotype 1, 4 and 5. The treatment was continued for 24 wk for those with genotype 2, 3 and those in whom HCV genotype was not known and for 48 wk for those with genotype 1, 4 and 5. Early virological response (EVR), end of treatment response (ETR) and sustained viral response (SVR) were documented when HCV RNA was undetectable or < 2 log, in comparison with baseline viral load at 12 wk from treatment (EVR), undetectable HCV-RNA at the end of treatment (ETR) and at 6 mo following completion of treatment respectively (SVR).

RESULTS

Fifty-two patients with SCA and HCV (Table 1) were treated, comprising 30 females and 22 males. Their mean age at the time of treatment was 29.5 years (range 15-54 years). Their mean HbS was 74.4% (66%-89%) and their mean HbF was 22.5% (9.6%-33.6%). Twelve (23.1%) had G6PD deficiency. HCV genotype was done in only 48. This indicated genotype 2 in 20; genotype 1 in 13, genotype 4 in 8, genotype 3 in 6 and one had genotype 5. Their mean pretreatment quantitative HCV- RNA level was 986330 IU/mL (range 12762-3329282 IU/mL). Fifteen underwent liver biopsy. This showed grade I in 6 and grade II in 9 and stage I in 13 and stage II in 2. Only 8 were receiving hydroxyurea at the time of treatment. All patients completed their therapy and in none of them was the treatment reduced or discontinued because of major side effects, hemolysis or bone marrow suppression. None of our patients suffered Hb drop which required blood transfusion pre, during or after

Table 1 Clinical data of SCA patients with chronic hepatitis

Total No. of patients	52
Sex	30 F: 22 M
Mean age (range)	29.5 (15-54) years
Mean HbS (range)	74.4% (66%-89%)
Mean HbF (range)	22.5% (9.6%-33.6%)
G6PD deficiency	12 (23.1%)
Mean HCV-RNA viral load (range)	986330 IU/mL (range 12762-3329282 IU/mL)
HCV genotype	
genotype 1	13 (8 non-responders)
genotype 2	20 (2 non-responders)
genotype 3	6
genotype 4	8 (4 non-responders)
genotype 5	1 (1 non-responder)
not available	4
Mean Hb level	
At start of treatment	10.2 g/dL (7.5-11.5 g/dL)
At 3 mo of treatment	10.4 g/dL (8.9-11.2 g/dL)
At 6 mo of treatment	10.35 g/dL (8.5-11.9 g/dL)
At the end of treatment	10.3 g/dL (8.2-11.7 g/dL)
SVR at 6 mo after the end of treatment	37 (71.2%)

SCA: Sickle cell anemia; SVR: Sustained viral response; HCV: Chronic hepatitis C.

therapy. Their mean hemoglobin level at the start of treatment was 10.2 g/dL (7.5-11.5 g/dL). There was no significant change in hemoglobin, LDH and bilirubin levels during or at the end of treatment. Their mean Hb level at 3 mo was 10.4 g/dL (8.9-11.2 g/dL), at 6 mo was 10.35 g/dL (8.5-11.9 g/dL) and at the end of treatment was 10.3 g/dL (8.2-11.7 g/dL). The lowest Hb level at the beginning of treatment was 7.5 g/dL. This patient did not receive pre-treatment blood transfusion and his Hb improved during treatment and at the end of treatment. His mean Hb during treatment was 9.6 g/dL (7.5-11.2 g/dL). The main adverse effects encountered in some of our patients were flu-like symptoms, headache, and loss of appetite and generalized body ache. These adverse effects were transient and tolerated by all patients. There were no more episodes of acute vaso-occlusive crisis, acute chest syndrome or other complications of sickle cell anemia while undergoing therapy. Thirty-seven (71.2%) of our patients achieved SVR at 6 mo after the end of treatment. The remaining 15 were non-responders. Two of them with an HCV genotype 2 showed an end of treatment response but had a relapse. The remaining 13 non responders had relatively significant HCV viral load with a mean HCV-RNA viral load 1829741.2 IU/mL (900000-3329282 IU/mL). Eight of them had HCV genotype 1, four had HCV genotype 4, and one had HCV genotype 5.

DISCUSSION

Patients with SCA frequently need blood transfusions to treat the various complications including acute and chronic anemia, splenic and hepatic sequestration crisis, acute chest syndrome, and priapism and central

nervous system crisis. The need for blood transfusion in these patients starts early especially in those with hemolytic crisis, splenic sequestration crisis and acute chest syndrome. This puts them at potential risk for alloimmunization, iron overload and chronic viral hepatitis B and C^[7,8]. Hepatitis C is one of the most common blood-borne infections worldwide. This is more so in those who receive frequent blood transfusions, especially patients with thalassemia and SCA^[10]. It is also well known that patients with HCV if left untreated are susceptible to liver damage, liver cirrhosis, hepatocellular carcinoma and liver failure. The combination of iron overload and HCV can lead to more rapidly progressive liver disease. The life expectancy of patients with SCA is known to have been shorter than the general population and they usually die of SCA-related complications. This however is not the case nowadays and as a result of better understanding of SCA and its complications, the use of hydroxyurea as well as improved care, SCA patients are living longer^[11-13]. In the past there was no well proven treatment for HCV, but recent progress has made the treatment and cure of HCV possible^[14-16]. The standard treatment for HCV is a combined therapy using pegylated interferon and ribavirin. Treatment response depends on several factors including HCV genotype^[14-16]. A sustained virological response in up to 60% of HCV patients with genotype 1 and 4 and up to 90% in those with genotype 2 and 3 has been reported^[15]. For many years, patients with SCA and HCV were considered to be unsuitable for such treatment. One reason for this is that ribavirin may induce hemolysis and severe anemia which can further aggravate patients' already existing anemia. Swaim *et al*^[17] were the first to report successful treatment of two patients with SCA and HCV using interferon alpha-2b and ribavirin. Ancel *et al*^[18] treated ten patients with HCV, five with thalassemia and five with SCA. Eight received pegylated interferon plus ribavirin while the other two were treated with pegylated interferon as a monotherapy. Nine out of 10 (90%) achieved a virological response and 6 (60%) went on to achieve sustained virological response after treatment was completed. Ayyub *et al*^[20] reported eight patients with SCA and HCV who were treated with pegylated interferon alpha-2a and ribavirin for one year. All eight patients had a complete early virological response and five of them maintained a sustained virological response when assessed 6 mo after the end of treatment. Our series is the largest to be reported so far. We treated 52 SCA patients with HCV and 37 (71.2%) of them showed SVR 6 months after the end of treatment. This is comparable to the results reported by others and to those for non SCA patients treated for HCV^[15,16,18-20]. It is also important to note that the dose of ribavirin was not reduced in any of our patients and the non-responders had a relatively high viral load which negatively predicts the response to treatment^[22]. Eight of them also had HCV genotype 1; four had HCV genotype 4, 2 had HCV genotype 2 and one had HCV genotype 5.

In all the few reports of patients with thalassemia and HCV who were treated with interferon and ribavirin, there was an increase in their transfusion requirements^[18,23-26]. Hamidah *et al*^[24] reported a significant increase in the transfusion requirements of one patient with thalassemia and HCV who was treated with pegylated interferon alpha-2b and ribavirin. Telfer *et al*^[25] treated 11 thalassemic patients with interferon and ribavirin and reported sustained virological response in 5 of them. There was however an increase in their transfusion requirements. Ancel *et al*^[18] reported a 22% increase in the transfusion requirements of five patients with thalassemia and HCV treated with pegylated interferon and ribavirin. Li *et al*^[26] reported a 30% increase in the transfusion requirement of a group of patients with thalassemia treated for HCV with interferon alpha-2b and ribavirin. In general, this was due to ribavirin-induced hemolysis. This however was not the case for patients with SCA where it was shown that either there was no change in their Hb level or in some of them there was actually an increase in their Hb level^[17-20]. This was also shown in our series where none of our patients required blood transfusion during or at the end of treatment and in none of them there was a need to decrease the dose of ribavirin or discontinue treatment. The reason for this difference between patients with thalassemia and SCA in this regard is not known. One contributing factor is hydroxyurea which is currently used to ameliorate the severity of SCA by increasing patients HbF levels^[11,12]. It was postulated that an increase in the HbF level in these patients may decrease the chance of ribavirin-induced hemolytic anemia. Because of this it was recommended to start and maintain these patients on hydroxyurea prior to therapy with ribavirin. SCA patients from the Eastern Province of Saudi Arabia are known to have high levels of HbF which is a contributing factor for the mildness of SCA in our patients^[27]. Our patients had a high mean HbF level of 22.5% (9.6%-33.6%). Only 8 of them were on hydroxyurea and their mean HbF level was 19.8% (11.5%-28.2%). These were started on hydroxyurea treatment prior to the decision to treat them for HCV and none of the remaining patients were intentionally started on hydroxyurea. There was no difference between those who were on hydroxyurea and those who did not receive hydroxyurea in terms of response to treatment or complications. We feel that the use of hydroxyurea prior to or during therapy for HCV in patients with SCA, although beneficial is not a necessity or a prerequisite for the treatment of HCV. This is especially so in our setting where our patients are already having high levels of HbF. It is also well known that ribavirin causes a dose-dependent hemolytic anemia. Another measure to decrease or avoid ribavirin-induced hemolysis was to start these patients at a lower dose of ribavirin as suggested by Ancel *et al*^[18]. Their patients received pegylated interferon at full dose while ribavirin was started at the lower dose of 200 mg twice daily and increased gradually until the full recommended dose was reached, usually within 4-8 wk from the beginning of therapy. This was also the

case for all the patients reported by Ayyub *et al*^[20]. Their patients were treated with pegylated interferon alpha-2a, 180 micrograms subcutaneously once per week and an initial dose of ribavirin 200 mg twice daily, increased gradually to 400 mg twice daily over a period of 4-8 wk. All our patients with HCV genotype 2 and 3 and those in whom the genotype was not available were started on 400 mg ribavirin twice daily and those with HCV genotype 1, 4 and 5 received 600 mg of ribavirin twice daily. None of our patients developed hemolysis, anemia or bone marrow suppression. We feel that patients with SCA and HCV can tolerate from the start the full dose of 400 mg of ribavirin twice daily for those with HCV genotypes 2 and 3 and up to 600 mg twice daily for those with genotypes 1 and 4 without subjecting them to additional major side effects.

In conclusion, patients with SCA and HCV can be treated with pegylated interferon and the usual recommended dose of ribavirin. This is even so for those who are not receiving hydroxyurea. The treatment is safe and effective and the response rate is comparable to those without SCA. There were no hematological side effects attributable to ribavirin even at the usual recommended dose. These patients however need to be followed-up closely while on therapy by a hematologist and a gastroenterologist. Further studies are important in this regard to establish definite guidelines for the treatment of SCA patients with HCV.

COMMENTS

Background

Sickle cell anemia (SCA) is one of the common hemoglobinopathies in the world. It is well known that SCA can affect any part of the body and that patients are liable to transfusion related hepatitis B and C. The life expectancy of patients with SCA has improved considerably and this is attributed to better understanding of the disease, improved medical and surgical management and the use of hydroxyurea. Improved life expectancy will put SCA patients with chronic hepatitis B and C at risk of developing liver cirrhosis, hepatocellular carcinoma and liver failure if their hepatitis B and C are not treated. In spite of the markedly improved results in the treatment of chronic hepatitis C using the combined therapy of pegylated interferon and ribavirin, patients with SCA and chronic hepatitis C were not previously considered suitable for such treatment because of the believed risk that ribavirin would induce hemolysis and severe anemia which could aggravate their already existing anemia.

Research frontiers

There have been three publications reporting success of such treatment in patients with SCA and HCV C although the number of patients reported was small and dose of ribavirin was not the standard.

Innovations and breakthroughs

The standard treatment for HCV is a combined therapy using pegylated interferon and ribavirin. Treatment response depends on several factors including HCV genotype. Patients with SCA and HCV were, for many years, considered to be unsuitable for such treatment. One reason for this is that ribavirin may induce hemolysis and severe anemia which can further aggravate already existing anemia. It is also well known that ribavirin causes a dose-dependent hemolytic anemia. In the previous studies patients were started at a lower dose of ribavirin as a measure to decrease or avoid ribavirin-induced hemolytic anemia. These patients received pegylated interferon at full dose while ribavirin was started at a lower dose of 200 mg twice daily and increased gradually until the full recommended dose was reached, usually within 4-8 wk from the beginning of therapy. This may have an impact on the RVR or SVR. All our patients with HCV genotype 2 and 3 and those in whom the genotype was not available were started on 400 mg ribavirin twice daily while those with HCV genotype 1, 4 and 5 received 600 mg of ribavirin twice daily. None of our

patients developed hemolysis, anemia or bone marrow suppression. Our series is the largest to be reported so far. We treated 52 SCA patients with HCV and 37 (71.2%) of them showed SVR. It was also postulated that an increase in the HbF level in these patients may decrease the chance of ribavirin induced hemolytic anemia. Because of this it was recommended to start and maintain these patients on hydroxyurea prior to therapy with ribavirin. Our patients had a high mean HbF level of 22.5% (9.6%-33.6%). We feel that the use of hydroxyurea prior to or during therapy for HCV in patients with SCA, although beneficial, is not a necessity or a prerequisite for the treatment of HCV. This is especially so in our setting where our patients already have high levels of HbF.

Applications

Patients with SCA and HCV can be treated safely with pegylated interferon in the usual recommended dose of ribavirin without subjecting them to additional major side effects. We feel that patients with SCA and HCV can tolerate from the start the full dose of 400 mg of ribavirin twice daily for those with HCV genotypes 2 and 3 and up to 600 mg twice daily for those with genotypes 1 and 4. This is even so for those who are not receiving hydroxyurea. The treatment is safe and effective and the response rate is comparable to those without SCA.

Peer review

The author presents a nice-sized case series of sickle-cell anemia patients treated for hepatitis C successfully. The paper adds to the literature and provides evidence that this population can be safely given ribavirin.

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Construction of non-covalent single-chain Fv dimers for hepatocellular carcinoma and their biological functions

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Abstract

AIM: To create new diabodies with improved binding activity to antigen of the variable light - variable heavy (VH-VL) oriented single-chain Fv dimers genes (scFv).

METHODS: The linker between VH and VL genes was shortened to 3-5 amino acid residues and cloned into the vector pCANTAB5E. The recombinant plasmids were transformed into TG1 cells and sequenced. The positive transformed cells were infected by M13K07 helper phage to form human recombinant phage antibodies. Expressed products were identified by SDS-PAGE, Western blotting, size exclusion gel chromatography (SEC), ELISA and immunohistochemistry.

RESULTS: Three scFv (scFv-3, scFv-4, scFv-5) were constructed successfully with binding ability to hepato-

cellular carcinoma 3.5-6 fold greater than their parental scFv. The single-chain Fv dimer (scFv-5, termed BDM3) with the best binding ability was successfully expressed in Yeast pichlia, as shown by SDS-PAGE and Western blotting. SEC results suggested the molecular weight of the expressed products was about 61 kDa. Expressed products showed significantly stronger binding to hepatocellular carcinoma cells than scFv, still having 50% binding activity even after 16 h incubation at 37°C. The purified dimers were bound specifically to the tumor antigen of HCC.

CONCLUSION: we have generated scFv dimers by shortening a series of linkers to 3-5 amino acid residues in VH-linker-VL orientation, resulting in highly stable and affinity-improved dimeric molecules. These will become an attractive targeting moiety in immunotherapeutic and diagnostic applications for HCC.

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Key words: Diabody; Antibody-targeted; Specificity; Affinity; Stability

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INTRODUCTION

Hepatocellular carcinoma (HCC) is the fifth most common cancer worldwide and the third most common cause of cancer-related death. The incidence of HCC is rising around the world. The diagnosis and treatment

of HCC have undergone major changes over the past decade. Unfortunately, there is no completely safe and effective treatment available for those who have developed HCC. Antibodies are a powerful tool for use in diagnosis and therapy of the tumor, because they bind with high specificity and high affinity to a variety of molecules, notably proteins and peptides. Whole antibodies provide high target binding specificity but their use in rapid tumor targeting and in vivo imaging is limited by slow tissue penetration, long circulating half-lives and often undesirable functions^[1]. Small monovalent antibody fragments (scFv and Fab) exhibit good tissue penetration but their lack of avidity results in faster off-rates and rapid clearance. Diabodies are noncovalent single-chain Fv (scFv) dimers formed by producing scFv with short (3-5 AA) linkers between their variable light (VL) and variable heavy (VH) chains. This prevents the VH and VL chains in a single molecule from associating with each other in a scFv orientation. Instead, the VL from one molecule associates with the VH from a second molecule, to form a dimer that binds antigen divalently^[2]. Several studies have demonstrated that genetically engineered antitumor diabody molecules can be used as effective vehicles for radioimmunotherapy, or can effectively reduce the growth rate of human tumor xenografts^[3-8]. Diabodies thus represent an improved strategy for selective tumor targeting as compared with scFv, Fab, or monoclonal antibody molecules.

In previous studies we have described the construction, screening and humanization of the scFv (termed HDM) against human hepatocellular carcinoma using phage display technology^[9-11]. HDM has binding specificity to hepatocellular carcinoma and is potentially effective in tumor imaging and therapeutics. To improve its antigen-binding avidity, we constructed the scFv dimers by shorting the linkers to 3-5 amino acid residues in same orientation as their parental scFv and assayed their biological functions, anticipating good behavior in antibody-targeted immunotherapeutic and diagnostic applications.

MATERIALS AND METHODS

Construction of scFv dimers

The pCANAB5E-HDM encoding the anti-HCC scFv HDM was previously constructed in our laboratory. This scFv, which has specificity for human hepatocellular carcinoma, is in the VH-linker-VL format, where the linker consists of the sequence (Gly4Ser)₃. The construct is cloned into the pCANAB 5E phagemid vector immediately downstream of the pIII leader sequence, which directs expression to the periplasm. The gene encoding the scFv is fused, *via* an amber codon, to the pIII gene of a filamentous phage. In *supE* strains of *E. coli* (TG1), this allows expression of the scFv on the surface of phage, as a fusion with the minor coat protein pIII, while in *supE-* strains (HB2151) translation is terminated at the amber codon, producing soluble scFv. A series of bivalent dimers (termed BDMs) was constructed by shortening

Table 1 Primer sequence

scFv-5 residues	
VH	Sense: 5'-TATGGCCCAGCCGGCCATGG-3'
	Antisense: 5'-AGAACCACCACCACCTGAGGAGACGGTGACC GT-3'
VL	Sense: 5'-TCAGGTGGTGGTGGTTCGACATTGAGCTCACCCA GTCTCCA-3'
	Antisense: 5'-TATGCGGCCGCCGTTTCA-3'
scFv-4 residues	
VH	Sense: 5'-TATGGCCCAGCCGGCCATGG-3'
	Antisense: 5'-AGAACCACCACCACCTGAGGAGACGGTGACCGT-3'
VL	Sense: 5'-TCCTCAGGTGGTGGTTCGACATTGAGCTCACCCA GTCTCCA-3'
	Antisense: 5'-TATGCGGCCGCCGTTTCA-3'
scFv-3 residues	
VH	Sense: 5'-TATGGCCCAGCCGGCCATGG-3'
	Antisense: 5'-AGAACCACCTGAGGAGACGGTGACCGT-3'
VL	Sense: 5'-GICTCTCAGGTGGTTCGACATTGAGCTCACCCA GTCTCCA-3'
	Antisense: 5'-TATGCGGCCGCCGTTTCA-3'
Sequencing primers	
	Sense: 5'-CAA CGT GAA AAA ATTATT CGC-3'
	Antisense: 5'-GTA AAT GAA TTT TCT GTA TGA GG-3'

the 15 amino acid linker between the VH and VL domain to 3-5 (Gly2Ser, Gly3Ser, Gly4Ser) residues. To obtain scFv dimers, BDM, the VH domain of scFv HDM, was PCR amplified with flanking restriction sites Sft I and the partial linker sequence. The VL domain of scFv HBM was PCR amplified with flanking restriction sites Not I and the partial linker sequence. Gene splicing by overlap extension was then performed using the VH sense and VL antisense primers. All primer sequences are showed in Table 1.

The PCR reactions were performed using standard conditions with 100 ng of template DNA, 200 mmol/L dNTP (TaKaRa, Japan) and 2.5 U PrimerStar polymerase (TaKaRa, Japan) with 35 cycles at 94°C for 30 sec, 64°C for 30 sec and 72°C for 1 min, using an Eppendorf Thermal Cycler (Eppendorf, Germany). The amplified products were then separated by agarose gel electrophoresis and gel purified using a Gel Extraction Kit (Omega, USA). Equimolar amounts of the two products were then mixed and used in a secondary SOE PCR. For the first five cycles no primers were added, and then the two 'external' primers (VH sense and VL antisense) were added for a further 30 cycles. The resulting full length product was then gel purified as described above, cut with the restriction enzymes SfiI and NotI (TaKaRa, Japan) and ligated onto the pCANAB5E vector molecules at their C-terminal ends with a hexahistidine epitope tag for detection and purification purposes by overnight reaction at 16°C using 0.1 unit T4 ligase (TaKaRa, Japan). The products were transformed into TG1 cells. 7-8 single colonies were picked from the each agar plate and grown overnight at 37°C in shaking culture in 2 × YT media containing 2% glucose and 100 µg/mL ampicillin. The sequence of positive colonies was determined with the ABI Perkin Elmer 373A auto-mated DNA sequencer (Applied

Biosystems, Forster City, CA), using the sequencing primers. Each sequence was determined at least 2 times.

Expression, purification and biophysical analyses of soluble dimers

The positive colonies were grown at 37°C in shaking culture in 2 × YT media containing 2% glucose and 100 µg/mL ampicillin to an OD 600 of ~0.5 and 1 × 10¹⁰/mL M13K07 helper phage (Pharmacia, Amersham.) was added. The bacteria were incubated for 1 h at 37°C before being centrifuged at 1500 g for 20 min and resuspended in 2 × YT containing 100 µg/mL ampicillin and 50 µg/mL kanamycin. Following overnight culture at 30°C the cultures were clarified by centrifugation and the phage precipitated using 2.5 mol/L NaCl/20% PEG 8000.

Cultures of *E. coli* HB2151(*supE*-) were infected with phage containing the relevant constructs. They were then diluted 1/100 and grown at 37°C in 2 × YT medium containing 2% glucose and 100 µg/mL ampicillin to an OD 600 of ~0.5. The bacteria were pelleted and resuspended in 2 × YT medium containing 100 µg/mL ampicillin and 1 mmol/L isopropyl-β-D-thiogalactopyranoside (IPTG). The cells were then grown for 12 h at 30°C^[9,10], centrifuged at 1500 g for 20 min, extracted and harvested for soluble diabodies from the periplasm. The soluble diabodies was purified with a HiTrap Anti-E Tag antibody column (Amersham Pharmacia), utilizing the hexahistidine epitope tag on their C-terminal ends.

The relative molecular mass of each affinity purified scFv dimer was compared by size exclusion gel chromatography on a Superdex 200 HR10/30 column (Amersham Pharmacia) run in PBS at a flow rate of 0.5 mL/min, calibrated with Biorad Gel Filtration Standard proteins^[12,13]. The purity of size-fractionated antibodies was monitored by sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) under reducing conditions after staining with Simply Blue Safe Stain (Invitrogen, Carlsbad, CA). The specificity of eluted fractions was determined by Western blot analysis using an anti His-Tag-peroxidase conjugated mAb (Amersham Pharmacia) followed by chemiluminescent detection (ECL Plus, Amersham Pharmacia).

ELISA analysis of binding activity of soluble scFv dimers to human hepatocellular carcinoma cell lines

The binding of soluble scFv dimers molecules was determined by ELISA. The mixing suspension of Bel-7402, SMMC-7721 and HepG-2 cells (Cell Bank of Wuhan University, China) were harvested and coated onto microtitre plates. Unoccupied sites on the plates were blocked using PBS-milk and the samples, diluted in PBS-milk from 1:1 to 1:256, were added and incubated for 2 h at room temperature. The plates were washed 5 times with PBS-milk and 5 times with PBS. The soluble scFv dimers were detected using the anti His-Tag monoclonal antibody, labeled with horseradish peroxidase (HRP)

conjugate (Pharmacia). The assays were developed using o-phenylenediamine (Dako) and absorbance was read at 490 nm wavelength in Model 680 Microplate Reader (Biorad, USA) with the parental scFv fragment HBM as control.

Immunohistochemistry

To determine the antigen-binding specificity of scFv dimers, immunohistochemistry was performed. Human hepatocellular carcinoma tissue and non-hepatocellular carcinoma tissue (donated by Professor Zhong, China) sections were heated at 56°C for 2 h, washed successively with dimethylbenzene twice for 20 min, 95% alcohol twice for 2 min, 80% alcohol once for 1 min, distilled water once for 1 min, PBS twice for 1 min. After washing, tissue sections were incubated with 3% H₂O₂ for 5 min in room temperature and the above washing steps were repeated once. Subsequently, sera (1:10) from BALB/c mice were added on the surfaces of tissue sections in a humidified atmosphere at room temperature for 10 min and the surplus sera were discarded. Purified scFv dimers were then added to the tissue sections. The tissue sections were kept in a humidified atmosphere at 4°C overnight and then washed with PBS for 5 min. The HRP-Anti-E Tag Conjugate was added and reacted under the above conditions overnight followed by washing with PBS for 3 min. Finally, 0.05% H₂O₂/DAB substrate was added to the tissue sections for 30 min. The specimens were photographed using the Leica Photo System (Qwin).

Biophysical stability analysis

At a concentration of 10 µg/mL, soluble scFv dimers were incubated at 37°C for up to 7 d. Samples were taken at different time points (at intervals of 12 h) and frozen at -20°C until the end of the experiment. Samples were subsequently analyzed for binding activity to human hepatocellular carcinoma cells.

RESULTS

Construction of scFv dimers

Three BDMs with short linkers, scFv-3, scFv-4 and scFv-5, were constructed in the pCANAB5E vector as described in Materials and Methods. The pictures of PCR reaction products and digested vector on agarose gel electrophoresis are shown in Figure 1. The DNA sequence of each scFv dimer construct was confirmed in both orientations to ensure that the correct sequence had been inserted between the VH and VL domains (gene sequences of BDMs not shown).

Expression, purification and biophysical analyses of soluble scFv dimers

The constructs were transferred to the HB2151 strain of *E. coli*. The bacteria were induced for 12 h at 30°C with 1 mmol/L IPTG. The soluble scFv fragments was extracted and purified as described above. Affinity purified scFv

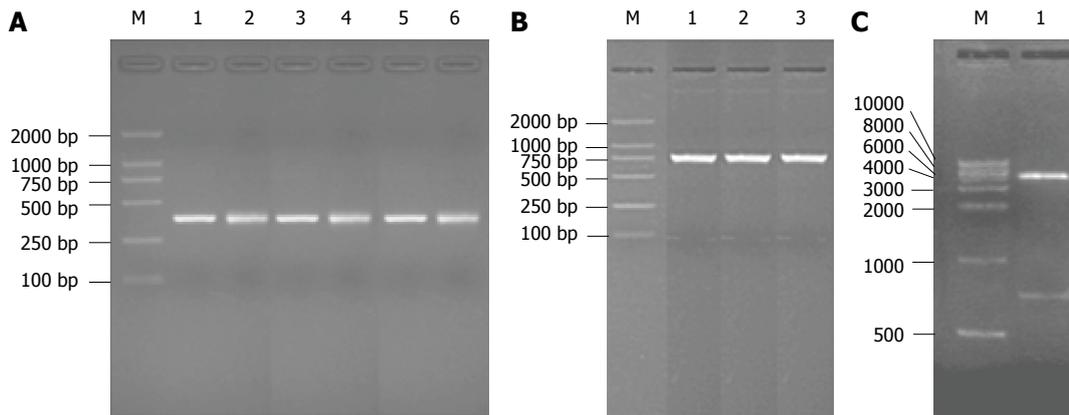


Figure 1 Agarose gel electrophoresis of PCR products. A: PCR products of VH and VL of scFv-3, scFv-4 and scFv-5. M: DNA marker DL2000; Lane 1: PCR products of the VH gene of scFv-3; Lane 2: PCR products of the VL gene of scFv-3. Lane 3: PCR products of the VH gene of scFv-4; Lane 4: PCR products of the VL gene of scFv-4. Lane 5: PCR products of the VH gene of scFv-5; Lane 6: PCR products of the VL gene of scFv-5; B: PCR products of scFv-3, scFv-4 and scFv-5. M: DNA marker DL2000; Lane 1: PCR products of the scFv-3 gene; Lane 2: PCR products of the scFv-4 gene; Lane 3: PCR products of the scFv-5 gene; C: Digested pCANAB5E vector. M: DNA marker; Lane 1: Digested pCANAB5E vector.

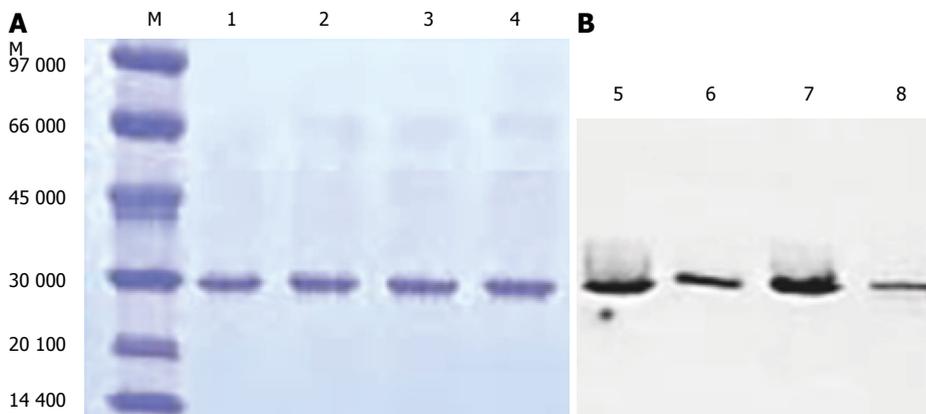


Figure 2 SDS-PAGE and Western blot. A: SDS gel electrophoresis of affinity-purified scFv fragments under reducing conditions. M: Low molecular weight markers (kDa); Lane 1: scFv-3; Lane 2: scFv-4; Lane 3: scFv-5; Lane 4: parental scFv HBM; B: Western blot. Lane 5: scFv-3; Lane 6: scFv-4; Lane 7: scFv-5; Lane 8: parental scFv HBM.

fragments and parental scFv HBM migrated as single bands with the expected molecular mass (30 kDa) on a reducing SDS-PAGE gel (Figure 2A). Western blot analysis (Figure 2B) was used to detect the C-terminal his-tag with an scFv-specific probe. The result indicated that the scFv dimers had been expressed successfully, the bands of dimers and monovalent scFv were all at 30 kDa because of the reducing conditions.

To examine the dimerization behavior of the scFv fragments, samples of scFv-3, scFv-4, scFv-5 were individually subjected to analysis by size exclusion gel chromatography on a calibrated Superdex 200 HR 10/30 column (Figure 3A). The major peaks for each of the scFv fragments showed the following distribution: ScFv-3 eluted in two peaks (25 and 28 min) corresponding to trimers (92 kDa) and dimers (61 kDa), respectively (Figure 3); ScFv-4 showed a predominant peak at an elution time of 28 min corresponding to a dimer (61 kDa) and an additional small peak indicating trimers (Figure 3); ScFv-5 was similar to ScFv-4 (not shown).

ELISA analysis of binding of soluble dimers to human hepatocellular carcinoma cell lines

Affinity constants of the scFv dimers for binding to human hepatocellular carcinoma cell lines were deter-

mined using a Model 680 Microplate Reader (Biorad) at 490 nm wavelength. All three scFv dimers showed improved binding activity compared to parental HDM (Figure 3B). Fitting the data from the equilibrium-binding curves into the non-linear regression model according to the Levenberg-Marquard method revealed a 3.5, 5.0 and 6.0-fold higher apparent affinity compared to the parental HDM for scFv-3, scFv-4 and scFv-5 respectively. The result also showed binding affinity of three dimers in a rank of scFv-5, scFv-4, scFv-3, with the binding activity decreasing according to diluted concentration.

Immunohistochemistry

The results of Immunohistochemistry were showed in Figure 4. As expected, scFv dimers react with human hepatocellular carcinoma tissue but not with non-hepatocellular carcinoma tissue. This suggests that scFv dimers have ideal specificity for binding to human hepatocellular carcinoma tissue. Therefore, the potential application of scFv dimers in clinical antibody-targeted diagnosis and therapy for HCC appears highly promising.

Stability analysis of scFv dimers

The thermostability of the dimeric molecules was analyzed. scFv-3 lost 50% of its initial binding activity

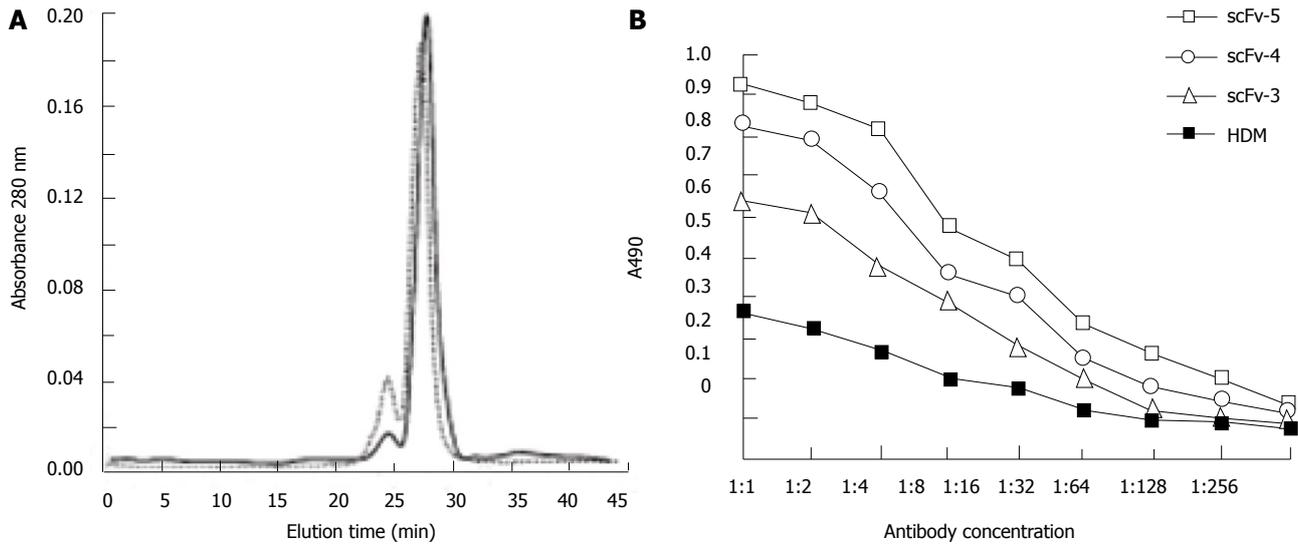


Figure 3 Analysis of soluble scFv dimers. A: Size exclusion chromatography of affinity purified scFv-3, scFv-4, scFv-5. Superimposed are elution profiles from a Superdex 200 gel filtration column of scFv-4 (solid line) and scFv-3 (dashed line). Elution peaks and molecular weight of calibration reference proteins are Gel Filtration Standard proteins (Biorad); B: Equilibrium binding curves of scFv dimers. Binding activity to hepatocellular carcinoma cell lines: ScFv-5 > scFv-4 > scFv-3 > HDM, with the tendency of decreasing binding activity going by diluted concentration. Determined By Model 680 Microplate Reader (Biorad).

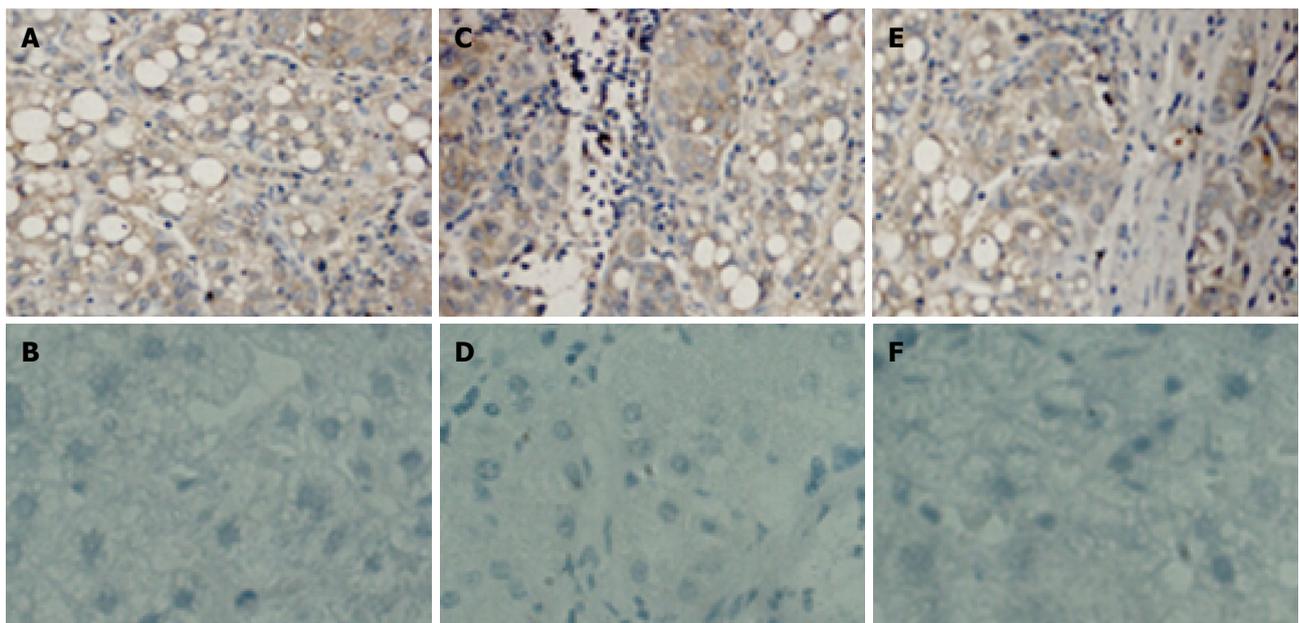


Figure 4 Immunohistochemistry of human hepatocellular carcinoma tissue. A: Reaction of scFv-3 dimer and human hepatocellular carcinoma tissue; B: Reaction of scFv-3 dimer and non- hepatocellular carcinoma tissue; C: Reaction of scFv-4 dimer and human hepatocellular carcinoma tissue; D: Reaction of scFv-4 dimer and non- hepatocellular carcinoma tissue; E: Reaction of scFv-5 dimer and human hepatocellular carcinoma tissue; F: Reaction of scFv-5 dimer and non- hepatocellular carcinoma tissue. Bound scFv dimers were detected with an antibody directed to the hexahistidine tag (Dianova) and peroxidase-conjugated rabbit anti-mouse immunoglobulins (Dako). Staining was performed with diaminobenzidine (DAB)/H₂O₂; nuclei were counterstained with hematoxylin. As expected, scFv dimers react with human hepatocellular carcinoma tissue but not with non- hepatocellular carcinoma tissue. The image was captured at 20 × magnification.

after only 18 hrs incubation at 37°C and completely lost binding activity after 70 hrs incubation. In contrast, scFv-4 was similar to scFv-5, with a 16 hr half life and totally lost binding activity after 60 hrs incubation.

DISCUSSION

Engineered antibodies possess considerable potential for immunotherapeutic and diagnostic applications and some

engineered antibodies have already been approved by FDA in the USA for clinical uses. Antitumor antibodies must bind to tumor antigens with high affinity to achieve durable tumor retention. This has spurred efforts to generate high affinity antibodies for use in cancer therapy. In our previous work, we constructed and biopanned a recombinant phage scFv library to obtain a scFv which has specificity for human hepatocellular carcinoma. This was in the VH-linker-VL format, with the linker

consisting of the sequence (Gly4Ser)₃. We then mutated and humanized the scFv to get scFv dimers (BDMs), which possess reduced immunogenicity^[9-11]. The purpose of this study was to generate and characterize bivalent scFv antibody derivatives from HDMs. The most straightforward method to generate a bivalent scFv is to shorten the variable domain connecting linker peptide, thereby allowing for the non-covalent association of multiple polypeptide chains to a dimeric molecule^[2,12-17], and to form a 'double-headed' fragment with two antigen binding sites that point away from each other^[18]. This compactness contributes not only to low immunogenicity and high tumor penetration but also to rapid clearance from the circulation^[15,19,20]. In addition to the linker length, the orientation of the variable domains was shown to impact the multimerization behavior and affinity of the polypeptide chains^[13,14,21,22]. The distance between the carboxyl terminus end of VL and the amino terminus of VH is greater than that for the opposite orientation. It has, therefore, been suggested that VL-VH orientated scFvs are more constrained than VH-VL oriented fragments when connected by the same linker and therefore tend to exhibit a higher tendency to form higher molecular weight oligomers^[13,14,23].

In this regard, we made several scFv dimers variants differing in linker length with a VH-linker-VL orientation. The sequence analysis showed that the sequences of VH-Gly2Ser-VL, VH-Gly3Ser-VL, VH-Gly4Ser-VL were exactly correct. Based upon size exclusion chromatographic profiles on Superdex 200, we conclude that the scFv-3, scFv-4 and scFv-5 predominantly formed stable dimers. This result is consistent with reports of other scFvs generated in the same domain orientation and with the same 3-5 amino acid linker^[12,21].

The most important advantage of multivalent scFvs over monovalent scFv and Fab fragments is the gain in functional binding affinity (avidity) to target antigens. High avidity requires that scFv multimers are capable of binding simultaneously to target antigens^[24]. The functional affinity of the BDMs was 3.5-6 fold that determined for the parent HDM, reflecting the gain in avidity due to dimerization and the capability of the molecule for simultaneous binding to two epitopes. The findings in immunohistochemistry and thermostability analysis have indicated that scFv dimers have improved specificity and stability and will become the paradigm for high-affinity antibody-based therapeutic and diagnostic reagents for HCC.

In conclusion, we have generated scFv dimers by shortening a series linkers to 3-5 amino acid residues in VH-linker-VL orientation, resulting in a set of highly stable and affinity-improved dimeric molecules. This work is crucial in making the antibody an attractive targeting moiety for HCC.

and the second most common cause of cancer-related death in our country. Traditional therapies, such as resection, chemotherapy and radiotherapy do not have satisfactory efficacy. Single chain variable fragments (scFv) can be a good vector in targeted therapy for HCC as a result of their low molecular weight, strong tumor tissue penetration, weak immunogenicity and short half-lives in blood. They also possess the same binding ability and specificity as the whole antibody.

Research frontiers

This team previously obtained a scFv against hepatocellular carcinoma (scFv 4-16, GenBank: DQ640759) through the phage display antibody library technology. After reconstruction of affinity maturation and humanization of scFv, we attained a scFv with high affinity to HCC (scFvDM) and a humanized scFv (scFvHDM).

Innovations and breakthroughs

scFv is a monovalent antibody with the shortcoming of low affinity, poor stability, too rapid clearance from the circulation and low-level expression in bacterial expression systems. This study set out to improve its binding activity to antigen by shortening the linker of the VH-VL oriented scFv to 3-5 residues. We examined the expression and biological functions of the resulting diabodies

Applications

These diabodies may become an attractive targeting moiety in immunotherapeutic and diagnostic applications for HCC.

Peer review

The authors described an incremental improvement on the genetic engineering of HDM for targeting HCC, which remains to be one of most difficult solid tumor to manage. I recommend accepting this manuscript.

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COMMENTS

Background

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A successful treatment by hepatic arterial infusion therapy for advanced, unresectable biliary tract cancer

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Abstract

Biliary tract cancers (BTC) are relatively rare tumors, and the prognosis is extremely poor. There has been no standard chemotherapy for advanced BTC. However, recently, gemcitabine (GEM) have been used against BTC as the most active agent, and promising response rates and overall survival times with tolerable drug toxicities have been observed. In this article, two cases of advanced intrahepatic cholangiocarcinoma and unresectable metastatic gallbladder (GB) cancer are reported. They were treated with hepatic arterial infusion (HAI) chemotherapy using a combination of GEM and cisplatin, along with the systemic administration of GEM. As a consequence, multiple liver tumors, the GB cancer and metastatic lymph nodes regressed without severe drug toxicities, and favorable results (the overall survival times were 16 and 14 mo, respectively) were achieved. In conclusion, HAI therapy using GEM combined with cisplatin may be a useful and well-tolerated option for advanced BTC, especially in cases where multiple liver metastases are detected.

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Key words: Hepatic arterial infusion; Gemcitabine; Cisplatin; Intrahepatic cholangiocarcinoma; Gallbladder cancer

INTRODUCTION

Biliary tract cancers (BTC) are relatively rare tumors, and their prognosis is extremely poor. The median survival period of gallbladder (GB) cancer is 6 mo, and cholangiocarcinoma (CC) also has an unfavorable prognosis, with a median survival time of 3-9 mo. In particular, intrahepatic cholangiocarcinoma (ICC) generally presents at a more advanced stage than extrahepatic CC. Therefore, its prognosis is even worse. In the past, the efficacy of conventional systemic chemotherapy in advanced, unresectable BTC was negligible. However, after the use of gemcitabine (GEM), a novel nucleoside analog, encouraging results for tumor control rates and survival time have been recently reported for BTC^[1-4]. Furthermore, a chemotherapeutic regimen using GEM combined with cisplatin has been demonstrated to be superior to GEM alone and/or a combination of fluoropyrimidines and cisplatin^[5-9]. Chemotherapy using hepatic arterial infusion (HAI) is a promising option for advanced and multiple hepatocellular carcinoma (HCC) and unresectable metastatic liver tumors of various origins, and this therapeutic modality has resulted in superior tumor control rates compared to systemic chemotherapy^[10-13]. However, there have been few reports

on HAI therapy using GEM together with cisplatin for advanced BTC.

In this article, two cases of advanced ICC and unresectable GB cancer are reported, which showed favorable outcomes when treated with HAI therapy using a combination of GEM and cisplatin.

CASE REPORT

Case 1

A 71-year-old man was admitted to our hospital with multiple hepatic tumors. A palpable liver was found in the right upper abdominal quadrant during physical examination, whereas no other relevant pathological findings were evident. The laboratory findings showed liver dysfunction. Among the serum tumor markers, α -fetoprotein (AFP) and carbohydrate antigen (CA) 19-9 were elevated (124.3 ng/mL, normal < 10; and 62 U/mL, normal < 37, respectively), but the level of carcinoembryonic antigen (CEA) was normal. Contrast-enhanced (CE) computed tomography (CT) of the abdomen revealed a huge liver tumor measuring 11 cm in diameter in the medial segment, and multiple small-sized tumors mainly located in the anterior and medial segment of the liver, which presented with ring enhancement in the arterial phase and central necrosis (Figure 1A and B). Multiple swollen lymph nodes (LNs) were noted in liver hilus and para-aorta, accompanied by ascites. No other tumors presented as primary foci in any other tissue outside the liver. Biopsied specimens from one of the hepatic tumors revealed a moderately-differentiated adenocarcinoma with widespread necrosis, which was not accompanied by hepatic cell components (Figure 2A). Additional histological examinations disclosed positive immunostaining for cytokeratin 7, and immuno-positive AFP in the adenocarcinoma cells (Figure 2B and C), suggesting that the hepatic tumors were AFP-producing cholangiocellular carcinomas. Under the diagnosis that the current case was an ICC of UICC TMN stage III C with an Eastern Cooperative Oncology Group (ECOG) performance status (PS) of 0, intensive chemotherapy was implemented, using both HAI along and intravenous (i.v.) administration. Following informed consent was given concerning the side effects of the therapy including interventional procedures and chemotherapeutic agents, 600 mg/m² of GEM on d 1 and 8, and 10 mg/body of cisplatin on d 1 to 3 and 8 to 10 were infused via a subcutaneously implanted port which was connected to a catheter placed in the proper hepatic artery. Concurrently, 400 mg/m² of GEM was administered intravenously on d 1 and 8, every 3 wk. The patient received 9 cycles of chemotherapy for 26 wk. Following completion of 2 cycles of chemotherapy, the huge tumor in the medial segment showed a reduction in diameter, and the ascitic fluid disappeared completely. However, an abnormal uptake on FDG-PET CT was observed in the hepatic tumors in both lobes, as well as the LNs in the liver

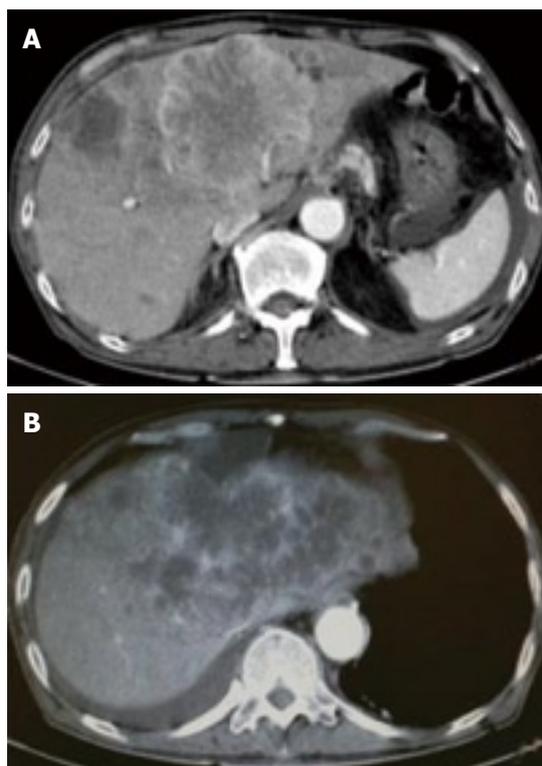


Figure 1 Contrast-enhanced (CE) computed tomography of the abdomen revealed a huge tumor, measuring 11 cm in diameter, which presented with ring enhancement in the arterial phase and central necrosis. A: In the medial segment; B: Multiple small-sized tumors mainly located in the anterior and medial segments of the liver.

hilus and para-aorta (Figure 3). After completion of 7 cycles of chemotherapy, almost all of the hepatic tumors had shrunk in size, and the swollen LNs were almost absent upon a CE-CT of the abdomen (Figure 4A). Meanwhile, a PET-CT revealed abnormal uptake only in the tumors in the medial segment, and no uptake in tumors in the right lobe and LNs (Figure 4B), suggesting that the tumor response was a partial response (PR), according to the Response Evaluation Criteria in Solid Tumors (RECIST) criteria. The elevated serum levels of CA19-9 and AFP were also decreased to almost the normal range after 8 cycles of chemotherapy (Figure 5). Following 9 cycles of chemotherapy, hematological toxicities such as severe thrombocytopenia (NCI-CTC, grade 3) and leukopenia (grade 2) were observed. Therefore, a modified chemotherapeutic regimen (80% of the initial dose of GEM on d 1, and 10 mg cisplatin on d 1 and 2) was carried out biweekly thereafter. Although a greater reduction of the abnormal uptake from the tumor in the medial segment was revealed on the PET-CT after the end of 11 cycles of chemotherapy, new abnormal uptake was found in the LN in the liver hilus. Therefore, the present case was diagnosed as a progressive disease (PD). A new chemotherapy regimen using GEM (480 mg/m² for HAI and 320 mg/m² for i.v. administration on d 2) and oral S-1 (80 mg/body for 7 to 14 d, consecutively) every 3 wk was started.

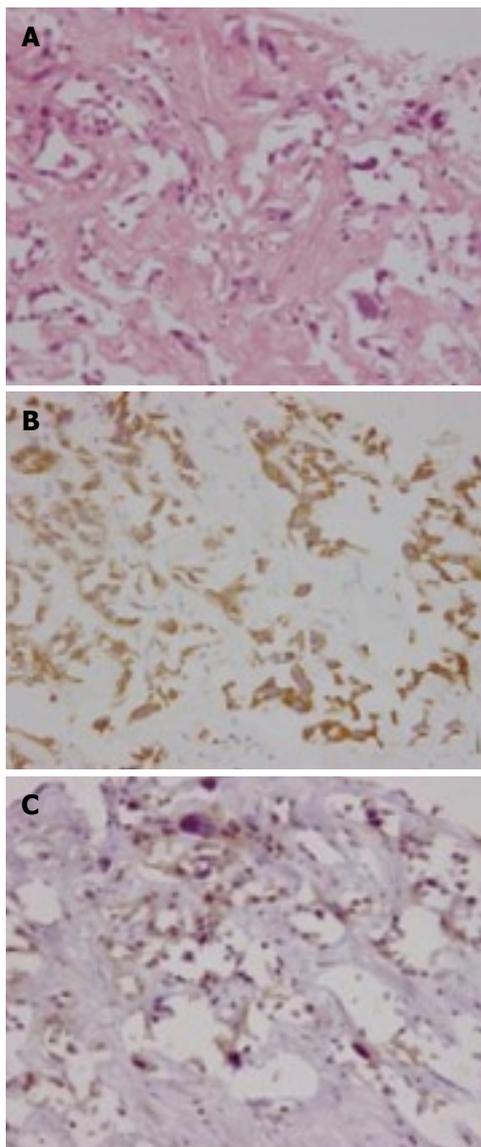


Figure 2 Biopsied specimens from one of the hepatic tumors. A: A moderately-differentiated adenocarcinoma with wide-spread necrosis, which were not accompanied by hepatic cell components (HE $\times 200$); B: Positive immunostaining for cytokeratin 7 ($\times 200$) in the adenocarcinoma cells; C: Immuno-positive AFP ($\times 200$) in the adenocarcinoma cells.

Following the end of 4 cycles of the new regimen, the liver tumors and metastatic LNs became larger. An ECF regimen using epirubicin 40 mg/m² and cisplatin 48 mg/m² for HAI on d 1, and 5-FU 160 mg/m² for i.v. administration for 4 consecutive days, was then carried out every 3 wk. However, systemic metastases (peritoneum, bone and lung) and massive ascites were soon observed. At 16 mo after the diagnosis, the patient died of multiple organ failure.

Case 2

An 83-year-old woman presented to our hospital with abdominal discomfort and appetite loss. Physical examination revealed a swollen gallbladder but no other pathological findings. The laboratory data demonstrated liver dysfunction and an elevation in the serum levels

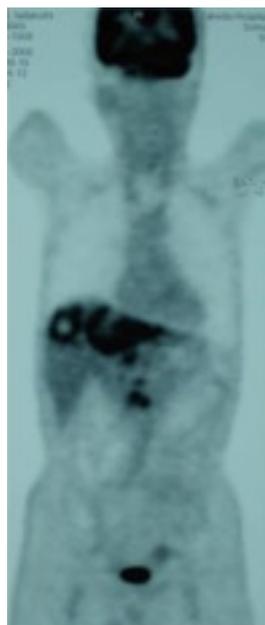


Figure 3 FDG-PET CT after the end of 2 cycles of chemotherapy revealed abnormal uptake in the hepatic tumors in both lobes and the LNs in the liver hilus and para-aorta.

of CA 19-9 and CEA (82.0 U/mL, normal < 37; 61.3 ng/mL, normal < 5, respectively). Ultrasonography and CE-CT of the abdomen revealed gall stones and a GB tumor, which invaded the adjacent liver directly. Multiple liver tumors and swollen LNs in the liver hilus were also observed. The patient's PS was 1 and, based on the diagnosis that the current case was an advanced GB carcinoma (UICC TNM stage IV), chemotherapy using a 4 wk cycle of i.v. administration of 1000 mg/m² of GEM on d 1, 8, 15 was started. After completion of 4 cycles of chemotherapy, the tumor increased in size in the GB, liver and LNs in the liver hilus (Figure 6A and B), and ascitic fluid appeared on CE-CT (RECIST, PD). The serum levels of CA 19-9 and CEA elevated gradually (Figure 7). Furthermore, metastasis in the lumbar vertebra was detected. Immediately thereafter, radiation therapy to the lumbar vertebrae was started. After informed consent was given, a heparin-coated catheter was placed in the common hepatic artery to supply all of the tumor vessels, including the GB and liver. Thereafter, the catheter was connected to the injection port. The gastroduodenal artery was occluded by steel coils to prevent gastroduodenal injury from anticancer drugs. Next, a new chemotherapeutic regimen comprising HAI (600 mg/m² of GEM on d 1 and 8, and 10 mg/body of cisplatin on d 1 to 3 and 8 to 10) and the i.v. administration of GEM (400 mg/m², d 1 and 8), was carried out every 3 wk for 3 cycles. After hematological toxicity (grade 2) occurred at the end of 3 cycles of chemotherapy, a modified regimen in which cisplatin administration was shortened to 2 d was instigated for 2 cycles. At the end of 3 cycles of chemotherapy with HAI, the tumors in the GB and liver decreased in size (Figure 8A and B), and the swollen LNs in the liver hilus regressed on the abdominal CT (RECIST, PR). However, following the 2 cycles of modified HAI chemotherapy, a perforation in the duodenum which was invaded directly

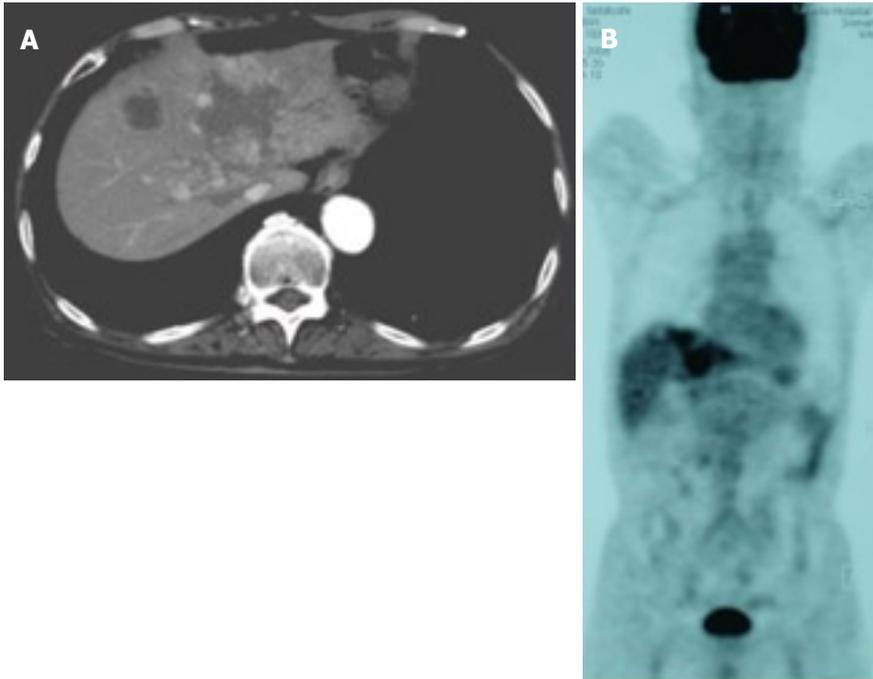


Figure 4 Images after the end of 7 cycles of chemotherapy. A: CE-CT showed a reduction in the size of a huge tumor seen in the medial segment; B: PET-CT revealed abnormal uptake only in the tumors in the medial segment, and no uptake was observed in tumors from the right lobe and the LNs in the liver hilum and para-aorta.

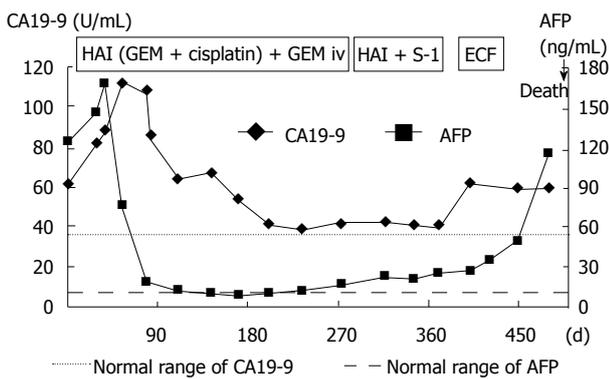


Figure 5 Clinical course and changes in the serum levels of CA19-9 and AFP.

by the GB cancer occurred. The chemotherapy was then discontinued, and 14 mo after the initial diagnosis, the patient was died of multiple organ failure.

DISCUSSION

In the past, there was no standard chemotherapy for advanced BTC i.e. GB cancer and CC. However, GEM has recently been used against BTC as the most active agent, and promising response rates and overall survival times with tolerable drug toxicities were demonstrated^[1-4]. GEM has a synergistic and cytotoxic effect *in vivo* and *in vitro*, in combination with cisplatin^[14,15] and capecitabine^[16]. More recently, the superiority of combination chemotherapy using GEM plus cisplatin, as well as the combination of GEM with capecitabine or oxaliplatin, has been demonstrated among the GEM-only, GEM-based and 5-FU-based chemotherapeutic regimens for BTC^[4,5,8,9,17-19]. The intra-arterial (i.a.) administration of GEM was used for the treatment of advanced pancreatic

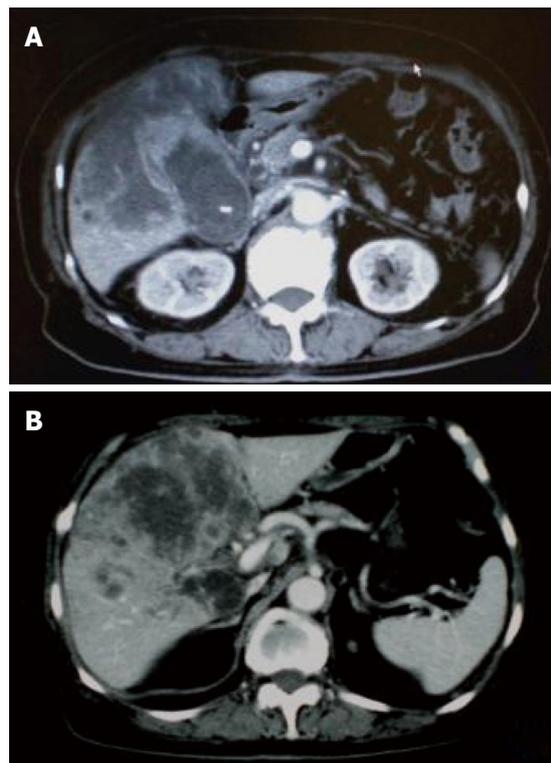


Figure 6 CE-CT after the end of 4 cycles of i.v. administration chemotherapy showed that the tumor increased in size. A: The GB tumor; B: Liver tumor.

carcinoma by Shamseddine et. al. in 2005^[20]. They reported that GEM in i.a. administration could be safely escalated to 1400 mg/m² within a tolerable toxicity. On the other hand, in low dose FP (5-FU and cisplatin) therapy for advanced and multiple HCCs, cisplatin is administered intra-arterially at a dose of 10 mg/body for 5 consecutive days^[13]. Based on the previous reports mentioned above and our judgment that the control of

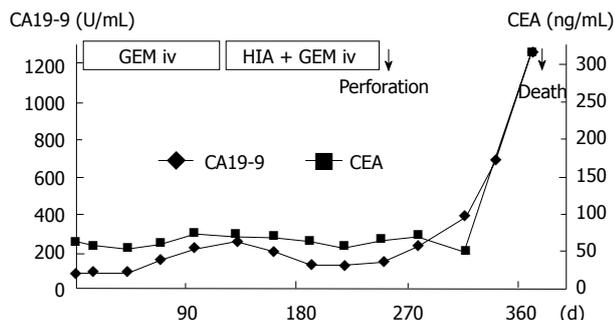


Figure 7 Clinical course and changes in the serum levels of CA19-9 and CEA.

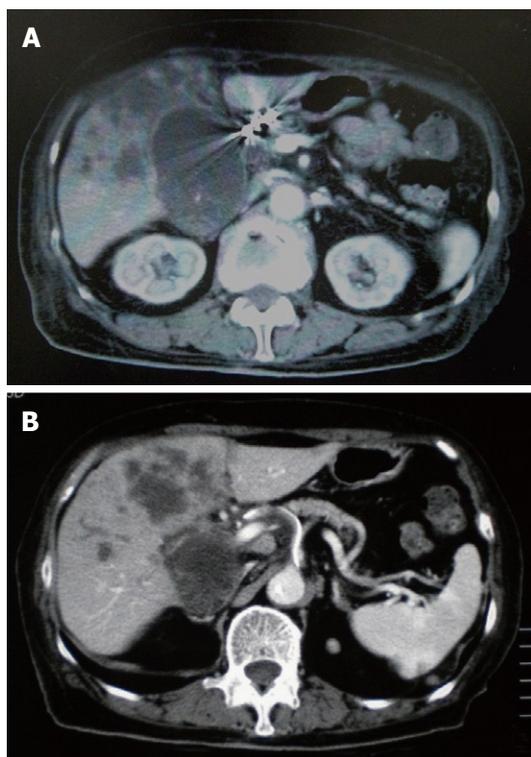


Figure 8 CE-CT after the end of 3 cycles of HAI chemotherapy showed a decrease in the size of tumor. A: The GB tumor; B: Liver tumor.

multiple liver tumors might decide the patient's survival time, HAI therapy using GEM combined with cisplatin was performed. Following consideration of treatment efficacy against distant metastases (LNs and bone), the systemic administration of GEM was used together with HAI in both cases. In case 1, after the end of 7 cycles of chemotherapy, the liver tumors had regressed, and the swollen LNs and ascitic fluid disappeared completely. In case 2, because of the PD after 4 cycles of the i.v. administration of GEM, HAI therapy was started again, and consequently favorable tumor responses were achieved. The survival times of the current two cases were also similar.

A chemotherapeutic regimen using the i.a. administration of GEM together with hepatic embolization has been performed against advanced BTC, and res-

pectable results were achieved^[21,22]; the survival time for the group treated by GEM with microspheres was 20.2 mo, whereas that of the TACE group using a combination of GEM plus cisplatin was 13.8 mo. Arterial administration can deliver anti-cancer agents at high concentrations into liver tumors, and has a longer lasting cytotoxic effect. A previous report demonstrated that the peak plasma concentration of GEM after i.a. administration was reduced to ~1/7th of that observed using the systemic i.v. route, and that no grade 3 or 4 toxicity was documented after i.a. administration of up to 1400 mg/m² of GEM^[20]. Vogl *et al*^[21] noted that the use of GEM doses (~1800 mg/m²) higher than the recommended 1000 mg/m² was well tolerated if microspheres were used. Moreover, Gusani *et al*^[22] showed that the median survival in unresectable CC treated with GEM-based TACE was not significantly different in patients with liver disease only, as compared to those with extra-hepatic disease. In 2008, HAI therapy using a combination of GEM and mitomycin against advanced CC revealed a poor tumor control rate, as compared to that against metastatic breast cancer and colorectal carcinoma^[12]. Therefore, HAI therapy using GEM combined with cisplatin may be a useful and well-tolerated option for advanced BTC, especially in which multiple liver metastases are detected.

In recent years, targeted therapy has been carried out for advanced BTC as a second-line chemotherapy^[23] and in phase II trials^[24]. Chemotherapeutic regimens using cetuximab or bevacizumab combined with GEM and oxaliplatin have also been used against advanced BTC and, at present, passable tumor-responses have been achieved^[23,24]. In conclusion, HAI therapy with GEM plus cisplatin might be an effective and well-tolerated option in advanced BTC, and in the future we recommend that clinical trials of HAI therapy using GEM-based platinum or capecitabine, with or without targeted agents, should be performed for advanced BTC.

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Intraoperative intracardiac thrombosis in a liver transplant patient

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Abstract

A 66-year-old female with cryptogenic cirrhosis complicated by ascites, hepatic encephalopathy, variceal bleeding and malnutrition with MELD of 34 underwent orthotopic deceased donor liver transplantation performed with piggyback technique. Extensive eversion thromboendovenectomy was performed for a portal vein thrombus which resulted in an excellent portal vein flow. The liver graft was recirculated without any hemodynamic instability. Subsequently, the patient became hypotensive progressing to asystole. She was resuscitated and a transesophageal probe was inserted which revealed a mobile right atrial thrombus and an underfilled poorly contractile right ventricle. The patient was noted to be coagulopathic at the time. She became progressively more stable with a TEE showing complete resolution of the intracardiac thrombus.

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Key words: Liver transplantation; Cardiac thrombus; Reperfusion injury; Coagulopathy; Transesophageal echocardiogram

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INTRODUCTION

Intracardiac thrombosis (ICT) during orthotopic liver transplantation is a catastrophic complication that carries a very high mortality. The exact cause of this condition is not known but it appears that it has multiple etiologies including administration of antifibrinolytic drugs intraoperatively and venovenous bypass.

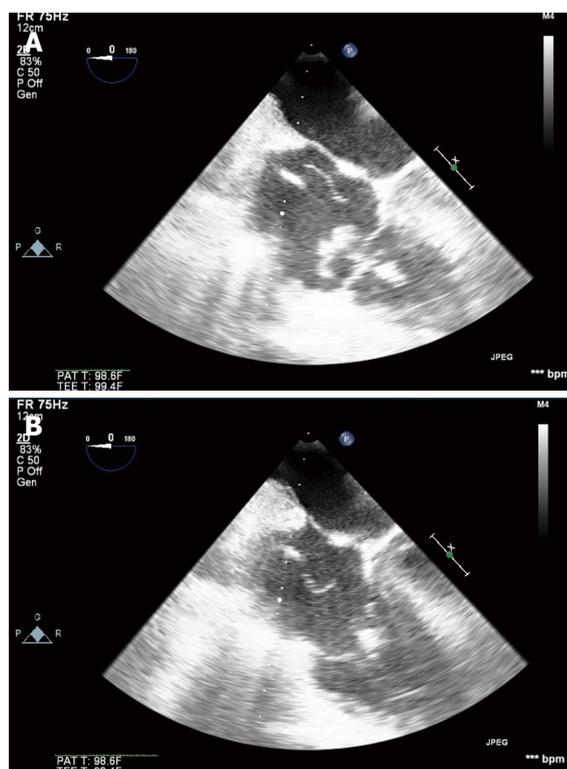
In this case report we describe a patient who developed ICT during orthotopic liver transplantation. No risk factors could be identified.

CASE REPORT

A 66-year-old female with cryptogenic cirrhosis was admitted to our Intensive Care Unit (ICU) with acutely worsening liver function. She had had multiple operations in the past including open cholecystectomy, total abdominal hysterectomy with salpingo-oophorectomy and appendectomy. She was noted to have a nodular liver while undergoing laparotomy for lysis of adhesions in 2007. Her liver disease was complicated by ascites, hepatic

Table 1 Patient coagulation parameters at the beginning of the case, at the time of the intracardiac thrombus diagnosis and at the end of the case

	Platelets / μ L	INR	Fibrinogen (mg/dL)
Beginning of case	44 000	1.8	210
Detection of clot	13 000	6.4	< 25
End of case	55 000	1.7	196

**Figure 1** Transesophageal echocardiography images. A: Large thrombus in right atrium in region of tricuspid valve and a second mass in right ventricle; B: empty right atrium with small mass on tricuspid valve; right ventricle is clear.

encephalopathy, variceal bleeding and malnutrition. She was on continuous venovenous hemofiltration for renal dysfunction. Her MELD was 34. A contrast-enhanced CT revealed a cirrhotic liver with thrombus in the portal vein extending into the superior mesenteric and splenic veins. The patient did not have any known hypercoagulable disorder and the thrombus in the portal vein was attributed to her liver disease. She was treated in the ICU for 15 d when an organ became available. The orthotopic deceased donor liver transplantation was performed with piggyback technique. Extensive eversion thromboendovenectomy resulted in an excellent portal vein flow. The liver graft was recirculated after the hepatic and portal vein anastomoses were completed. The patient tolerated recirculation well without any hemodynamic instability. While the hepatic artery was dissected and mobilized for the arterial anastomosis, the patient became acutely hypotensive progressing to asystole. She was resuscitated with external cardiac massage and inotropic support. A transesophageal echocardiography (TEE)

probe was emergently inserted and showed mobile right atrial thrombus and an underfilled, poorly contractile right ventricle (Figure 1A). At the time of resuscitation, INR was 6.4, fibrinogen < 25 mg/dL and platelets 13 000/ μ L (Table 1). No antifibrinolytic therapy was administered. Continued monitoring by TEE showed the clot adherent to the tricuspid valve (Figure 1B) and later was not seen on four chamber view indicating either spontaneous thrombolysis or propagation into the pulmonary vasculature. The right ventricular function and filling improved and the patient became hemodynamically more stable, allowing successful completion of the liver transplantation. The patient was transferred to the ICU for postoperative care on dopamine drip (10 mcg/kg per minute). The patient was weaned off pressors on POD zero and was successfully extubated on POD two. An echocardiogram performed on POD two demonstrated normal right and left ventricular size and function with ejection fraction of 57%. There was no evidence of intracardiac thrombus. The postoperative ultrasound of the liver graft revealed widely patent hepatic veins, hepatic artery and portal vein with normal flows. Ultrasound of the extremities was negative for deep vein thrombosis. The patient currently has normal liver function and continues to make good progress.

DISCUSSION

Intracardiac thrombosis (ICT) during liver transplantation is a devastating complication with poor outcomes. Warnaar *et al*^[1] reported a mortality rate of 68% with a majority of patients dying in the operating room. This complication is extremely rare (1%-1.5%) and unfortunately the exact etiology is not known. It has been associated with intraoperative administration of anti-fibrinolytics including ϵ -aminocaproic acid, aprotinin and clotting factors used to decrease intraoperative bleeding and transfusion requirement^[2]. Other potential risk factors described are the presence of venovenous bypass, pulmonary artery catheter, continuous venovenous hemofiltration, migration of preexisting thrombi, sepsis and disseminated intravascular coagulation. Empiric administration of antifibrinolytics and clotting factors should be strongly discouraged since the risk of ICT formation and pulmonary embolus at times outweighs the risk of intraoperative bleeding.

Systemic hypotension, increase in central venous pressure, pulmonary artery pressure, decrease in cardiac output and cardiac arrest are frequently encountered signs^[2]. The definitive diagnosis is made on transesophageal echocardiography.

There is no consensus for treatment since the condition is so rare. Administration of tissue plasminogen activator into the right atrium^[2,3] and heparin boluses^[4] have been described although they are associated with potentially massive hemorrhage. Suction embolectomy is another treatment option.

The source of the ICT in our patient is uncertain.

It is unclear whether this clot was preformed or newly developed. After recirculation, our patient became profoundly coagulopathic. We believe that the fibrinolysis that occurred after reperfusion of the liver graft led to the lysis of the thrombus and subsequently to a successful outcome.

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Huge extrahepatic portal vein aneurysm as a late complication of liver transplantation

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Abstract

A 60-year-old male underwent orthotopic liver transplantation because of hepatitis C virus related cirrhosis. After 12 d, the patient underwent re-transplantation due to primary graft non function. One year later the patient developed a thrombosis of the main portal vein needing a surgical revision. After 11 years the patient was operated on because of a clinical picture of intestinal occlusion. As an incidental finding, a large aneurysm of the main portal vein was diagnosed. The incidence of intra- and extrahepatic Portal vein aneurysms (PVAs) is not clear. To the best of our knowledge, only one case of intrahepatic PVA in a liver transplant has been reported in the literature. In addition, we have found no documented cases of extrahepatic PVAs in liver transplanted patients.

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Key words: Portal vein aneurysm; Late complication; Liver transplantation; Portal hypertension; Doppler ultrasound

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TO THE EDITOR

A 60-year-old male underwent orthotopic liver transplantation in Cambridge, England in 1993, because of hepatitis C virus related cirrhosis. After 12 d, the patient underwent re-transplantation due to primary graft non function. In 1994, a thrombosis of the main portal vein (MPV) was detected. As a result, the patient underwent surgical reconstruction (there were no medical records or details of the previous surgery in the patient's possession). From 1994 to 2003, the patient showed no liver function alterations. However, liver ultrasound (US) performed during follow-up showed a slow dimensional increase of the MPV (16 mm in December 1994, 18 mm in 1996, 25 mm in 2002, and 30 mm in 2003).

The patient did not undergo any other radiological examinations at any time. In June 2003, the patient was admitted to another hospital for acute intestinal occlusion, with abdominal pain and vomiting. The patient underwent explorative laparotomy with adhesiolysis. At that time no aneurysm of the portal vein was diagnosed or detected.

In July 2003, the patient was admitted to our center with a clinical picture of intestinal occlusion. 64



Figure 1 Multi intensity projection shows a large aneurysm (maximum diameter 6.3 cm) of the MPV that arose 1 cm after the spleno-mesenteric confluence.

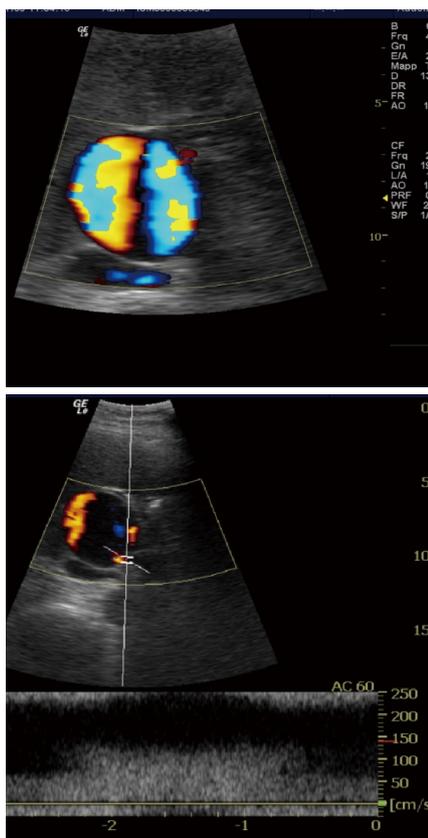


Figure 2 Color-Doppler US shows turbulent flow inside the aneurysm (red and blue color). An increase velocity of flow is detected at the origin of the aneurysm.

Multi Detector CT scan (Lightspeed VCT, GE) performed with and without contrast media showed an overdilatation of the small bowel and right colon in the presence of a severe neoplastic lesion occluding the transverse colon. A colonoscopy confirmed the lesion, which was also biopsy proven. As an incidental finding, a large aneurysm of the MPV, 6.5 cm in diameter and arising 1 cm after the spleno-mesenteric confluence, was found (Figure 1). The patient underwent surgery for

colon resection; the presence of tenacious adhesences made it impossible to check the portal vein situation. A mild amount of ascites and splenomegaly was detected (longitudinal diameter of 16.3 cm) with no other radiological signs of portal hypertension. At that time liver function tests were normal. Doppler US showed increased velocity near the origin of the aneurysm, and turbulent flow inside. Monophasic flow with normal velocity was detected in the intrahepatic portal branches (Figure 2).

Extrahepatic portal vein aneurysms, defined as MPV greater than 1.9 cm, are rare in non-transplanted patients and in cirrhotic patients, and only 67 cases have been documented, in the English literature. Portal vein aneurysms (PVAs) can be congenital or acquired, and even if the etiology remains unclear, there are some related conditions, such as portal hypertension, pancreatitis or trauma.

PVAs are usually asymptomatic^[1-3] if small, while large PVAs can cause biliary tract obstruction, chronic portal hypertension, portal vein thrombosis, acute portal hypertension, abdominal discomfort and, though rarely, adjacent structure compression and rupture.

Our patient showed no symptoms or complications related to the aneurysm that justified surgical treatment. The patient was followed with Doppler US every three months.

The incidence of intra- and extrahepatic PVAs is not clear. To the best of our knowledge, only one case of intrahepatic PVA in a liver transplant has been reported in the literature^[4]. In addition, we have found no documented cases of extrahepatic PVAs in liver transplanted patients. We believe that in our case, one or more of the three surgical interventions on the portal vein, created a weak point in the portal vein which was responsible, over a period of years, for the development of the aneurysm. In addition, at the time of the colon resection surgery, the intestinal occlusion could have increased the intra-abdominal pressure and consequently the portal pressure, leading to the enlargement of the portal aneurysm.

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Meetings

Events Calendar 2010

January 25-26
 Tamilnadu, India
 International Conference on Medical
 Negligence and Litigation in Medical
 Practice

January 25-29
 Waikoloa, HI, United States
 Selected Topics in Internal Medicine

January 26-27
 Dubai, United Arab Emirates
 2nd Middle East Gastroenterology
 Conference

March 04-06
 Bethesda, MD, United States
 8th International Symposium on
 Targeted Anticancer Therapies

March 05-07
 Peshawar, Pakistan
 26th Pakistan Society of
 Gastroenterology & Endoscopy
 Meeting

March 12-14
 Bhubaneswar, India
 18th Annual Meeting of Indian
 National Association for Study of
 the Liver

March 25-28
 Beijing, China
 The 20th Conference of the Asian
 Pacific Association for the Study of
 the Liver

March 27-28
 San Diego, California, United States
 25th Annual New Treatments in
 Chronic Liver Disease

April 07-09
 Dubai, United Arab Emirates
 The 6th Emirates Gastroenterology
 and Hepatology Conference, EGHC
 2010

April 14-18
 Vienna, Austria
 The International Liver Congress™
 2010

May 01-05
 New Orleans, LA, United States
 Digestive Disease Week Annual
 Meeting

May 06-08
 Munich, Germany
 The Power of Programming:
 International Conference on
 Developmental Origins of Health
 and Disease

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 Chicago, IL, United States
 American Society of Clinical
 Oncologists Annual Meeting

June 16-19
 Hong Kong, China
 ILTS: International Liver
 Transplantation Society ILTS Annual
 International Congress

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 International Liver Association's
 Fourth Annual Conference

September 12-15
 Boston, MA, United States
 ICAAC: Interscience Conference
 on Antimicrobial Agents and
 Chemotherapy Annual Meeting

September 16-18
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 Prague Hepatology Meeting 2010

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J Gastroenterol 2007; **13**: 6356-6364 [PMID: 18081224
DOI: 10.3748/wjg.13.6356]

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In press

- 3 **Tian D**, Araki H, Stahl E, Bergelson J, Kreitman M. Signature of balancing selection in Arabidopsis. *Proc Natl Acad Sci USA* 2006; In press

Organization as author

- 4 **Diabetes Prevention Program Research Group**. Hypertension, insulin, and proinsulin in participants with impaired glucose tolerance. *Hypertension* 2002; **40**: 679-686 [PMID: 12411462 PMCID:2516377 DOI:10.1161/01.HYP.0000035706.28494.09]

Both personal authors and an organization as author

- 5 **Vallancien G**, Emberton M, Harving N, van Moorselaar RJ; Alf-One Study Group. Sexual dysfunction in 1, 274 European men suffering from lower urinary tract symptoms. *J Urol* 2003; **169**: 2257-2261 [PMID: 12771764 DOI:10.1097/01.ju.0000067940.76090.73]

No author given

- 6 21st century heart solution may have a sting in the tail. *BMJ* 2002; **325**: 184 [PMID: 12142303 DOI:10.1136/bmj.325.7357.184]

Volume with supplement

- 7 **Geraud G**, Spierings EL, Keywood C. Tolerability and safety of frovatriptan with short- and long-term use for treatment of migraine and in comparison with sumatriptan. *Headache* 2002; **42** Suppl 2: S93-99 [PMID: 12028325 DOI:10.1046/j.1526-4610.42.s2.7.x]

Issue with no volume

- 8 **Banit DM**, Kaufer H, Hartford JM. Intraoperative frozen section analysis in revision total joint arthroplasty. *Clin Orthop Relat Res* 2002; (**401**): 230-238 [PMID: 12151900 DOI:10.1097/00003086-200208000-00026]

No volume or issue

- 9 Outreach: Bringing HIV-positive individuals into care. *HRS A Careaction* 2002; 1-6 [PMID: 12154804]

Books

Personal author(s)

- 10 **Sherlock S**, Dooley J. Diseases of the liver and biliary system. 9th ed. Oxford: Blackwell Sci Pub, 1993: 258-296

Chapter in a book (list all authors)

- 11 **Lam SK**. Academic investigator's perspectives of medical treatment for peptic ulcer. In: Swabb EA, Azabo S. Ulcer disease: investigation and basis for therapy. New York: Marcel Dekker, 1991: 431-450

Author(s) and editor(s)

- 12 **Breedlove GK**, Schorfheide AM. Adolescent pregnancy. 2nd ed. Wiczorek RR, editor. White Plains (NY): March of Dimes Education Services, 2001: 20-34

Conference proceedings

- 13 **Harnden P**, Joffe JK, Jones WG, editors. Germ cell tumours V. Proceedings of the 5th Germ cell tumours Conference; 2001 Sep 13-15; Leeds, UK. New York: Springer, 2002: 30-56

Conference paper

- 14 **Christensen S**, Oppacher F. An analysis of Koza's computational effort statistic for genetic programming. In: Foster JA, Lutton E, Miller J, Ryan C, Tettamanzi AG, editors.

Genetic programming. EuroGP 2002: Proceedings of the 5th European Conference on Genetic Programming; 2002 Apr 3-5; Kinsdale, Ireland. Berlin: Springer, 2002: 182-191

Electronic journal (list all authors)

- 15 Morse SS. Factors in the emergence of infectious diseases. *Emerg Infect Dis serial online*, 1995-01-03, cited 1996-06-05; 1(1): 24 screens. Available from: URL: <http://www.cdc.gov/ncidod/EID/eid.htm>

Patent (list all authors)

- 16 **Pagedas AC**, inventor; Ancel Surgical R&D Inc., assignee. Flexible endoscopic grasping and cutting device and positioning tool assembly. United States patent US 20020103498. 2002 Aug 1

Statistical data

Write as mean \pm SD or mean \pm SE.

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Express *t* test as *t* (in italics), *F* test as *F* (in italics), chi square test as χ^2 (in Greek), related coefficient as *r* (in italics), degree of freedom as ν (in Greek), sample number as *n* (in italics), and probability as *P* (in italics).

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