

## VIRAL HEPATITIS

**Branched-chain amino acids reduce hepatic iron accumulation and oxidative stress in hepatitis C virus polyprotein-expressing mice**Masaaki Korenaga<sup>1,2</sup>, Sohji Nishina<sup>1</sup>, Keiko Korenaga<sup>1</sup>, Yasuyuki Tomiyama<sup>1</sup>, Naoko Yoshioka<sup>1</sup>, Yuichi Hara<sup>1</sup>, Yusuke Sasaki<sup>3</sup>, Yasushi Shimonaka<sup>3</sup> and Keisuke Hino<sup>1</sup>

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**Keywords**

hepatitis C virus – hepatic mitochondrial dysfunction – hepcidin-25 – iron metabolic disorder – reactive oxygen species

**Abbreviations**

BAP, biological antioxidant potential; BCAA, branched-chain amino acids; BTR, ratio of BCAA relative to tyrosine; CHOP, CCAAT/enhancer-binding protein homology protein; CPT I, carnitine palmitoyl transferase I; dROM, derivatives of reactive oxygen metabolites; HCC, hepatocellular carcinoma; HCV, hepatitis C virus; HCVTgM, transgenic mice expressing hepatitis C virus polyprotein; ROS, reactive oxygen species; SOD2, superoxide dismutase 2; SREBP, sterol regulatory element-binding protein.

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Hepatitis C virus (HCV) causes acute and chronic hepatitis, cirrhosis, and hepatocellular carcinoma (HCC) (1). New direct-acting antiviral treatments are expected to

eliminate this virus in about 90% of patients (2), but therapies that could reduce disease progression in chronically infected individuals would be highly beneficial.

**Abstract**

**Background & Aims:** Branched-chain amino acids (BCAA) reduce the incidence of hepatocellular carcinoma (HCC) in patients with cirrhosis. However, the mechanisms that underlie these effects remain unknown. Previously, we reported that oxidative stress in male transgenic mice that expressed hepatitis C virus polyprotein (HCVTgM) caused hepatic iron accumulation by reducing hepcidin transcription, thereby leading to HCC development. This study investigated whether long-term treatment with BCAA reduced hepatic iron accumulation and oxidative stress in iron-overloaded HCVTgM and in patients with HCV-related advanced fibrosis. **Methods:** Male HCVTgM were fed an excess-iron diet that comprised either casein or 3.0% BCAA, or a control diet, for 6 months. **Results:** For HCVTgM, BCAA supplementation increased the serum hepcidin-25 levels and antioxidant status [ratio of biological antioxidant potential (BAP) relative to derivatives of reactive oxygen metabolites (dROM)], decreased the hepatic iron contents, attenuated reactive oxygen species generation, and restored mitochondrial superoxide dismutase expression and mitochondrial complex I activity in the liver compared with mice fed the control diet. After 48 weeks of BCAA supplementation in patients with HCV-related advanced fibrosis, BAP/dROM and serum hepcidin-25 increased and serum ferritin decreased compared with the pretreatment levels. **Conclusions:** BCAA supplementation reduced oxidative stress by restoring mitochondrial function and improved iron metabolism by increasing hepcidin-25 in both iron-overloaded HCVTgM and patients with HCV-related advanced fibrosis. These activities of BCAA may partially account for their inhibitory effects on HCC development in cirrhosis patients.

Valine, leucine and isoleucine are essential branched-chain amino acids (BCAA). A decreased ratio of serum BCAA relative to aromatic amino acids, a hallmark of cirrhosis, is caused by several factors, including reduced nutritional intake and ammonia detoxification in skeletal muscles (3). BCAA supplementation can improve the nutritional status and albumin synthesis by activating the mammalian target of rapamycin signalling cascade (4, 5) and glucose metabolism in skeletal muscles (6, 7). Long-term oral BCAA supplementation decreases the frequency of HCC in male obese patients with cirrhosis (8). BCAA also had an antihepatocarcinogenic activity in an animal model of insulin resistance (9, 10). In addition, glucose intolerance is closely linked to hepatocarcinogenesis. However, the mechanisms that underlie these effects remain unknown.

Hepatic oxidative stress and iron overload have been implicated in liver injury and hepatocarcinogenesis in HCV-associated chronic liver diseases (11, 12). The HCV core protein inhibits mitochondrial complex I and generates reactive oxygen species (ROS) *in vivo* (13). Previously, we reported that HCV-induced ROS increases the hepatic iron concentration by reducing hepcidin transcription in transgenic mice that express HCV polyprotein (14), where even modest iron supplementation in these mice resulted in the development of liver tumours, including HCC, because of mitochondrial injury (15). Thus, hepatic iron overload and oxidative stress via mitochondrial injury are critical during HCC pathogenesis.

In the present study, we examined whether long-term BCAA supplementation could prevent the development of hepatic iron accumulation and oxidative stress in HCV transgenic mice fed an excess-iron diet and in patients with HCV-related advanced fibrosis.

## Materials and methods

### Animals and experimental design

The pAlbSVPA-HCV transgene contains the full-length HCV polyprotein-coding region under the control of the murine albumin promoter/enhancer (16, 17). HCV polyprotein is processed into individual proteins in the liver and expressed at biologically relevant levels in FLN/35 transgenic mice (HCVTgM) (17). In the present study, male HCVTgM (8 weeks old) were fed a normal rodent diet, including carbonyl iron (45 mg/kg; control,  $n = 6$ ), or an excess-iron diet (carbonyl iron, 225 mg/kg) that contained either 3.0% BCAA (BCAA/iron;  $n = 5$ ) or casein (casein/iron;  $n = 7$ ). Six months later, the mice were sacrificed by CO<sub>2</sub> asphyxiation after a 12-h fast, according to the criteria outlined in the Guide for the Care and Use of Laboratory Animals.

### Clinical chemistry tests

The serum concentrations of alanine aminotransferase, aspartate aminotransferase (AST), albumin, glucose,

insulin, BCAA, tyrosine and hepcidin-25 were determined in blood samples collected from the inferior vena cava of sacrificed mice at 12 h after fasting. The blood glucose levels were periodically measured using a glucometer (OneTouch Ultra, Lifescan, Inc., Milpitas, CA, USA). The serum insulin levels were measured using an ultrasensitive mouse insulin ELISA kit (Morinaga Milk, Kanagawa, Japan). The serum hepcidin-25 levels were determined by LC/MS/MS (18).

### Hepatic iron and triglyceride contents

The hepatic iron concentrations were measured by atomic absorption spectrometry, as described previously (15). The liver tissue was homogenized and the lipids were extracted (19), and the triglyceride levels were measured using a TGE-test Wako kit (Wako Pure Chemicals, Tokyo, Japan), according to the manufacturer's instructions. The protein concentrations were determined by the Lowry method (20) using a DC protein assay kit (Bio-Rad Laboratories, Hercules, CA, USA).

### *In situ* ROS detection

*In situ* liver ROS production was assessed by staining with dihydroethidium (Invitrogen Corp., Carlsbad, CA, USA), as described previously (14). Dihydroethidium is oxidized to ethidium bromide in the presence of ROS, which stains the nuclei bright red via DNA intercalation (21). The intensity of the fluorescence was quantified using the NIH Image analysis program in three randomly selected areas of the digital images for each mouse.

### Derivatives of reactive oxygen metabolites and biological antioxidant potential levels

The derivatives of reactive oxygen metabolites (dROM) and biological antioxidant potential (BAP) levels were measured using a Free Radical Elective Evaluator (Wismarll Co. Ltd, Tokyo, Japan) (22, 23). The dROM measurements were determined based on the ability of transition metals to catalyse the formation of coloured free radicals (detection at 505 nm). The results were expressed in Cartelli units (U.CARR), where 1 U.CARR = 0.8 mg/L of H<sub>2</sub>O<sub>2</sub>. To obtain the BAP measurements, the blood samples were added to a solution containing FeCl<sub>3</sub> bound to a chromogenic substrate (AT, a derivative of thiocyanate). Fe<sup>3+</sup> reduction to Fe<sup>2+</sup> caused a chromatic change that was directly proportional to the plasma ROS reduction, which was measured at 505 nm using a photometer. Blood aliquots (10 µl) were mixed with the FeCl<sub>3</sub> solution and incubated for 5 min at 37°C before the photometric analysis.

### Histological staining

Part of each liver sample was snap-frozen immediately in liquid nitrogen to determine the hepatic triglycerides

and iron concentration. The remaining liver tissue was fixed in 4% paraformaldehyde in phosphate-buffered saline and embedded in paraffin for use in the histological analysis. The liver sections were stained with haematoxylin and eosin.

#### Real-time reverse transcriptase-PCR

One-step real-time reverse transcriptase-PCR (RT-PCR) was performed, as described previously (14), where the results were expressed as the hepcidin, interleukin 6 (IL6), BMP6 and superoxide dismutase 2 (SOD2) gene mRNA levels relative to  $\beta$ -actin mRNA.

#### Extraction of nuclear and histone deacetylase activity assay

For isolation of nuclear proteins from mice liver, Nuclear Extraction Kit 1 (Epigentek, Farmingdale, NY, USA) was used. Histone deacetylase (HDAC) activity was assessed using HDAC Activity/Inhibition Direct Assay Kit (Epigentek) according to the manufacturer's instruction.

#### Isolation of mitochondria and complex I activity determination

Liver mitochondria were isolated and the activity of complex I was assayed (at 25°C) as described previously (3, 24).

#### Protein extraction and Western blotting

The liver lysate and mitochondrial lysate proteins were separated by sodium dodecyl sulphate-polyacrylamide gel electrophoresis. These proteins were then transferred to polyvinylidene difluoride membranes (Millipore, Bedford, MA, USA) and blocked overnight at 4°C with 1–3% skim milk and 0.1% Tween 20 in Tris-buffered saline, which was followed by incubation at room temperature for 1 h with a primary antibody. Anti-rabbit carnitine palmitoyl transferase I (CPT I), anti-rabbit CPT II (Alpha Diagnostic International, San Antonio, TX, USA), anti-rabbit SREBP1 (Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA), or anti-bacterially expressed mouse CCAAT/enhancer-binding protein homology protein (CHOP) fusion protein (Abcam, Cambridge, England) were used for the liver lysate proteins. Anti-SOD2 (Abcam), anti-Grp75 (mitochondrial heat shock protein70; Abcam), or anti-NDUFB8 (mitochondrial complex I) antibody (Abcam) were used for the mitochondrial lysates. The proteins were blocked for 1 h at room temperature and then incubated overnight at 4°C with a Phospho-stat3 (pSTAT3) antibody (Cell Signaling Technology Inc., Danvers, MA, USA) and a Phospho-Smad1/Smad5/Smad8 (pSMAD1/5/8) antibody (Cell Signaling Technology Inc.). The anti-acetyl-

histoneH3K9 and anti-histoneH3 (Cell Signaling Technology Inc.) were used for the nuclear lysates.

#### Human BCAA supplementation study design

We screened 68 HCV RNA-positive patients who were aged >65 years (Fig. S1). We enrolled 25 patients with HCV-related advanced fibrosis who satisfied the following criteria: serum albumin = 3.5–4.2 g/dl; platelet counts  $<15 \times 10^4/\mu\text{l}$ ; amino acid imbalance [based on the ratio of BCAA relative to tyrosine (BTR)  $<4.40$ , which was lower than the normal limits]; and no HCC or symptoms of chronic liver failure such as ascites, varices or hepatic encephalopathy. Advanced fibrosis defined liver specimens (METAVIR fibrosis staging:  $>F3,4$ ) or Fib-4 index ( $>3.25$ ). The patients were assigned randomly to receive BCAA supplementation (BCAA group;  $n = 12$ ) or follow-up without treatment (non-BCAA group;  $n = 13$ ). BCAA group were given a 4 g BCAA preparation (LIVACT Granules; Ajinomoto, Tokyo, Japan) administered orally three times daily after meals. We measured the plasma oxidized/reduced albumin and serum dROM and BAP as oxidative stress markers at 12, 24 and 48 weeks after starting the treatment. We also measured the levels of serum iron, ferritin, transferrin saturation (TSAT) and hepcidin-25 to evaluate the oxidative stress-associated iron metabolism. Moreover, type IV collagen 7s, type III procollagen peptide (PIIP) and Fib-4 index were measured to confirm the degree of hepatic fibrosis.

Written informed consent was obtained from each study participant. This study was conducted in accordance with the provisions of the 1975 Declaration of Helsinki and it was approved by the Institutional Ethics Committee of Kawasaki Medical School.

#### Statistical analysis

The results were expressed as mean  $\pm$  SD. The group results were compared using Levene's or Welch's tests. The changes in the levels of the iron metabolism and oxidative stress markers between the BCAA and the non-BCAA groups were analysed using Wilcoxon rank-sum tests. Pearson's product moment correlation coefficient was used to assess associations between the dihydroethidium-positive areas and the BAP and dROM ratios. Differences were considered statistically significant at  $P < 0.05$ . The statistical analyses were performed using SPSS software (IBM SPSS Statistics 20.0 for Windows).

#### Results

##### AST, fasting blood sugar, plasma BCAA and tyrosine levels in HCVtgM

The dietary intake and body weight did not differ significantly between the three groups of mice. BCAA administration for 6 months significantly reduced the serum

AST ( $P < 0.05$ ) and fasting blood sugar (FBS) levels ( $P < 0.05$ ) compared with HCV TgM fed the excess-iron diet with casein (casein/iron group) (Table 1). However, the FBS levels remained higher in the HCV TgM fed the excess-iron diet with BCAA (BCAA/iron group) compared with the HCV TgM fed a normal rodent diet (control group) ( $P < 0.05$ ). The casein/iron group had significantly lower plasma BCAA and the ratio of BCAA relative to tyrosine (BTR) levels ( $P < 0.05$ ) compared with the BCAA/iron and control groups (Table 1). The tyrosine levels were significantly higher in the casein/iron group than the control group ( $P < 0.05$ ).

#### Hepatic iron contents and hepcidin-25 levels in HCV TgM

The hepatic iron contents of HCV TgM fed the excess-iron diet with casein were significantly higher than those of HCV TgM fed an excess-iron diet with BCAA or a control diet at 6 months after the treatment commenced (Fig. 1A). The hepcidin levels of HCV TgM fed the excess-iron diet with BCAA were significantly higher than those of HCV TgM fed the excess-iron diet with casein or the control diet (Fig. 1A). The serum hepcidin to ferritin ratio was lower in patients with HCV (25). The serum hepcidin-25 to hepatic iron ratio was significantly higher in HCV TgM fed the excess-iron diet with BCAA compared with those fed the excess-iron diet with casein or the control diet.

#### ROS generation

BCAA administration resulted in significantly lower dROM levels and an increased BAP to dROM ratio (BAP/dROM) compared with casein administration ( $P < 0.05$ ; Fig. 1B). Hepatic ROS production, which

was determined by dihydroethidium staining, was significantly higher in HCV TgM fed the excess-iron diet with casein compared with those fed the excess-iron diet with BCAA or the control diet (Fig. 1C). The BAP/dROM ratios were negatively correlated with hepatic ROS production in all mice ( $r = 0.8985$ ;  $n = 15$ ;  $P < 0.01$ ; Fig. 1D).

#### Factors that affected hepcidin upregulation

HCV-induced ROS production downregulates hepcidin transcription by inhibiting the C/EBP $\alpha$  DNA-binding activity of CHOP (14). Thus, we examined CHOP expression and the hepcidin mRNA levels. Hepatic CHOP expression was significantly lower and hepatic hepcidin expression was significantly higher in HCV TgM fed the excess-iron diet with BCAA compared with the levels in HCV TgM fed the excess-iron diet plus casein (Fig. 2A,B). The IL-6-gp130/signal transducer and activator of transcription are involved in the regulation of hepcidin transcription (26). Another pathway that regulates hepcidin expression involves the TGF- $\beta$ /bone morphogenetic protein superfamily (27, 28). Thus, we examined the STAT-IL6 and SMAD-BMP signalling pathways. There were no differences in the phosphate STAT3, IL6, phosphor-SMAD1/5/8 and BMP6 expression levels between the BCAA and casein groups (Fig. 2C). In addition, HCV-induced oxidative stress inhibited hepcidin expression through increased histone deacetylase (HDAC) activity in cell culture system (29). HDAC activity of HCV TgM fed the excess-iron diet with BCAA was significantly lower than those of HCV TgM fed the excess-iron diet with casein or the control diet (Fig. S2). These results suggested that BCAA induced the upregulation of hepatic hepcidin by enhancing the antioxidant potential.

**Table 1.** Effects of casein/iron and branched-chain amino acids (BCAA)/iron diets on the liver to body weight ratios and blood chemistry results in hepatitis C virus transgenic mice

	Control	Casein/iron	BCAA/iron
Mice (N)	6	7	5
Liver weight/ Body weight (%)	3.32 $\pm$ 0.30	3.50 $\pm$ 0.60	2.97 $\pm$ 0.28
AST (IU/L)	61 $\pm$ 23	92 $\pm$ 47	39 $\pm$ 7‡
ALT (IU/L)	14 $\pm$ 4	61 $\pm$ 60	16 $\pm$ 4
FBS (mg/dl)	115 $\pm$ 11	299 $\pm$ 49†	184 $\pm$ 47†‡
Insulin (ng/ml)	0.89 $\pm$ 0.36	1.19 $\pm$ 0.20	0.93 $\pm$ 0.39
Albumin (g/dl)	2.82 $\pm$ 0.04	2.77 $\pm$ 0.15	2.96 $\pm$ 0.15
BCAA (nmol/ml)	313 $\pm$ 22	275 $\pm$ 31†	318 $\pm$ 35‡
Tyrosine (nmol/ml)	63 $\pm$ 5	82 $\pm$ 11†	69 $\pm$ 11
BTR	5.01 $\pm$ 0.20	3.41 $\pm$ 0.40†	4.67 $\pm$ 0.40‡

\*Results are mean  $\pm$  SD.

† $P < 0.05$  vs. transgenic mice expressing hepatitis C virus polyprotein (HCV TgM) on control diet for 6 months.

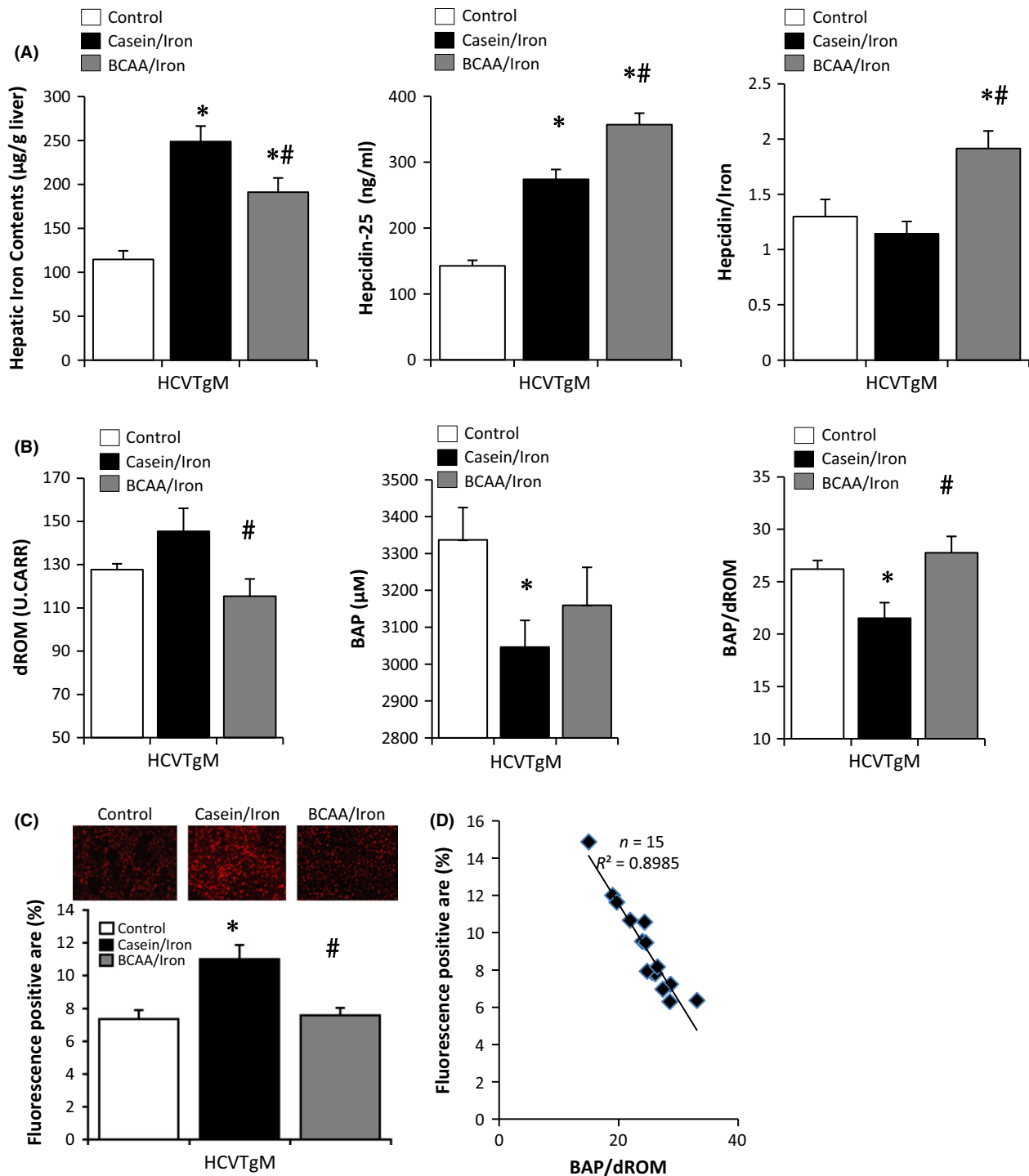
‡ $P < 0.05$  vs. HCV TgM on excess-iron diet with casein for 6 months.

ALT, alanine aminotransferase; AST, aspartate aminotransferase; FBS, fasting blood sugar.

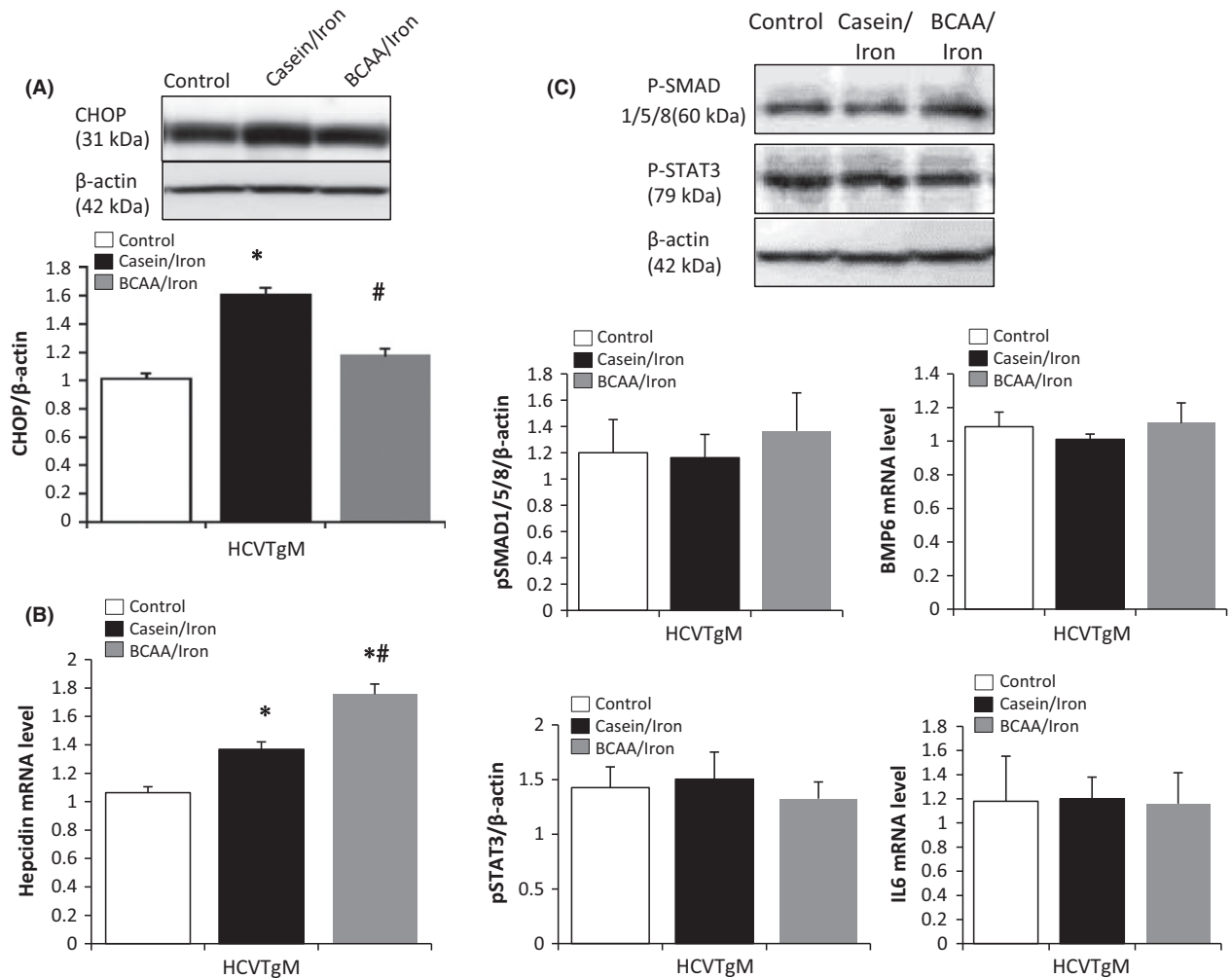
#### Hepatic steatosis and CPT1 expression

HCV TgM fed the excess-iron diet developed severe steatosis, including the centrilobular microvesicular type (15, 17). Previous studies showed that the antioxidant drugs N-acetylcysteine (NAC) and Stronger Neo-Minophagen C (SNMC) reduce the hepatic triglyceride levels in a dose-dependent manner (30, 31). In the present study, BCAA administration tended to reduce the hepatic triglyceride levels ( $P = 0.055$ ; Fig. 3A).

Thus, we examined the effects of BCAA on CPT1 and CPT2, which are proteins that regulate long-chain fatty acid oxidation in mitochondria, and SREBP1 expression, which is a transcription factor that activates genes required for lipogenesis. Our previous study indicated that decreased CPT1 and increased SREBP1 expression contribute to the development of hepatic steatosis in HCV TgM fed an excess-iron diet (30). In the present study, CPT1 expression increased significantly in HCV TgM fed the excess-iron diet with BCAA after 6 months ( $P < 0.05$ , Fig. 3C), whereas CPT2 expression



**Fig. 1.** (A) Hepatic iron contents, hepcidin-25 levels, and hepcidin-25 to iron content ratios (hepcidin/iron). (Left) Hepatic iron contents in mice at 6 months after starting treatment for the control ( $n = 6$ ), casein/iron ( $n = 7$ ) and BCAA/iron groups ( $n = 5$ ). (Centre) Serum hepcidin-25 levels. (Right) The hepcidin/iron ratios were used as an index of the sensitivity of hepcidin upregulation against iron overload. (B) Oxidative stress markers in serum. (Left) dROM and (centre) BAP were measured at 6 months after starting treatment. (Right) The antioxidant status was determined as the BAP to dROM ratio. (C) Dihydroethidium fluorescence intensity was quantified for three randomly selected areas in digital images for the control ( $n = 3$ ), casein/iron ( $n = 7$ ), and BCAA/iron groups ( $n = 5$ ) at 6 months after starting treatment. (D) Correlations between the BAP/dROM ratios and fluorescence-positive areas in liver. \* $P < 0.05$  vs control group; # $P < 0.05$  vs casein/iron group.



**Fig. 2.** (A) Immunoblots for CHOP after 6 months of treatment. (B) Liver hepcidin expression was determined for four mice in each group (C) Immunoblots for p-SMAD1/5/8 and (D) p-STAT3 after 6 months of treatment. (E) BMP6 and (F) IL6 mRNA expression was determined for four mice in each group. The protein expression levels were normalized against that of  $\beta$ -actin. \* $P < 0.05$  vs control group; # $P < 0.05$  vs casein/iron group.

did not increase significantly. However, SREBP1 expression did not decrease in HCVtgM fed the excess-iron diet with BCAA ( $P = 0.082$ ; Fig. 3B). These results suggest that the administration of BCAA was insufficient to prevent iron-induced steatosis in HCVtgM because BCAA failed to reduce SREBP1 expression.

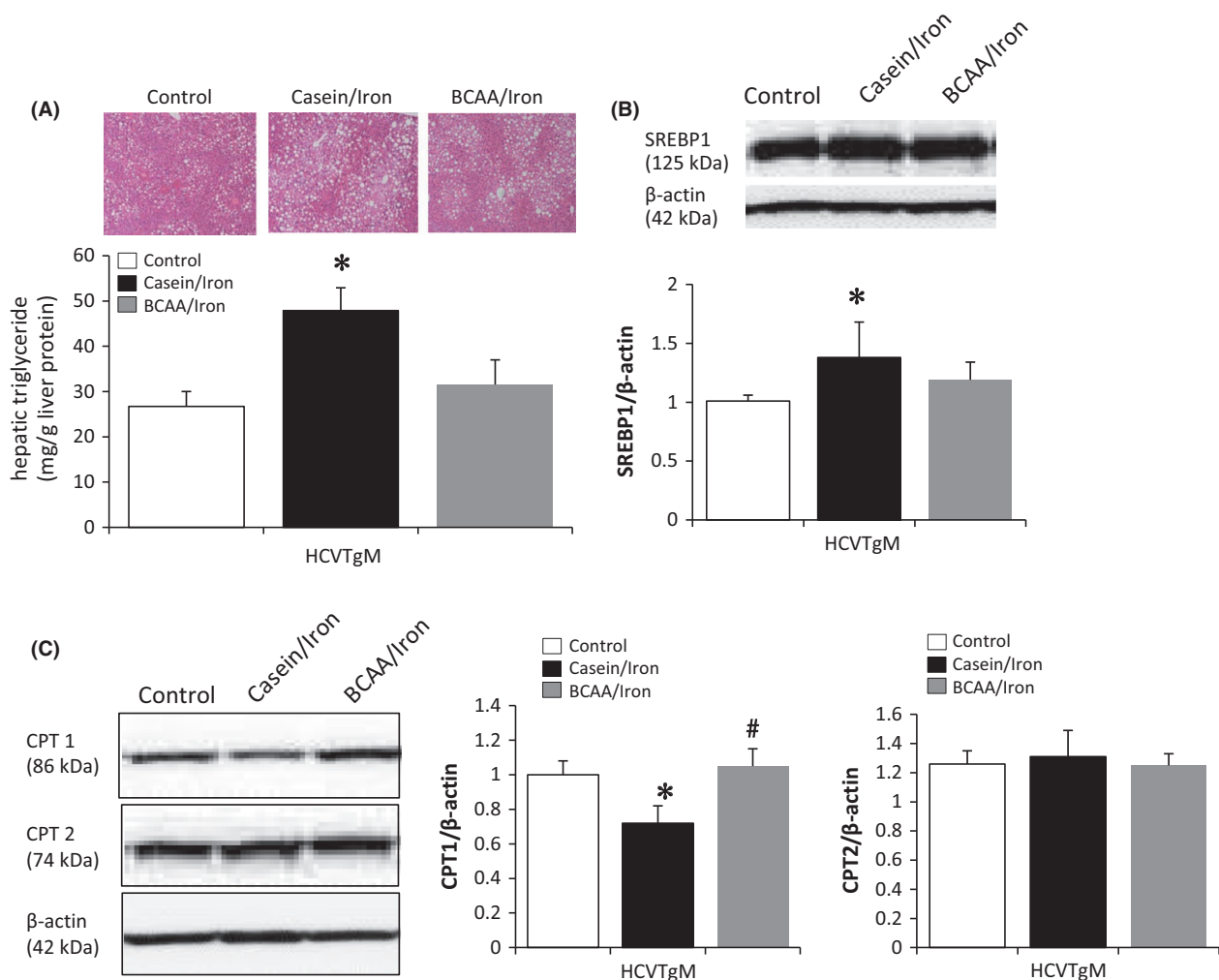
#### SOD2 expression and mitochondrial complex I activity

CPT1 is localized to the mitochondrial outer membrane. Decreased CPT1 expression may be related to the HCV core protein's association with the mitochondrial outer membrane. The HCV core protein interacts with mitochondria complex I, which generates ROS (13). Alterations in the mitochondrial ultrastructure were observed in HCVtgM fed the excess-iron diet after 6 months, as described previously (15, 30). We exam-

ined whether BCAA supplementation reduced iron- and HCV-induced mitochondrial injury.

The mitochondrial SOD2 mRNA levels were significantly higher in HCVtgM fed the excess-iron diet with BCAA compared with those fed the excess-iron diet with casein or the control diet. The SOD2 expression levels in mice fed the excess-iron diet with casein were significantly lower than those fed the control diet. However, the SOD2 expression levels were restored by BCAA supplementation (Fig. 4A). After 6 months, the mitochondrial complex I expression levels were significantly lower in mice fed the excess-iron diet with casein compared with those fed the control diet. Similar to SOD2, the mitochondrial complex I expression levels were restored by BCAA supplementation (Fig. 4B).

The enzymatic activity of mitochondrial complex I was significantly lower in mice fed the excess-iron diet



**Fig. 3.** (A) Hepatic steatosis in HCVtgM fed the excess-iron diet with BCAA and HCVtgM fed the excess-iron diet with casein after treatment for 6 months (haematoxylin and eosin, original magnification  $\times 100$ ). The hepatic triglyceride levels were determined. (B) Immunoblots of SREBP1, (C) CPT1, and (C) CPT2 in the livers of three mice from each group. \* $P < 0.05$  vs control group; # $P < 0.05$  vs casein/iron group.

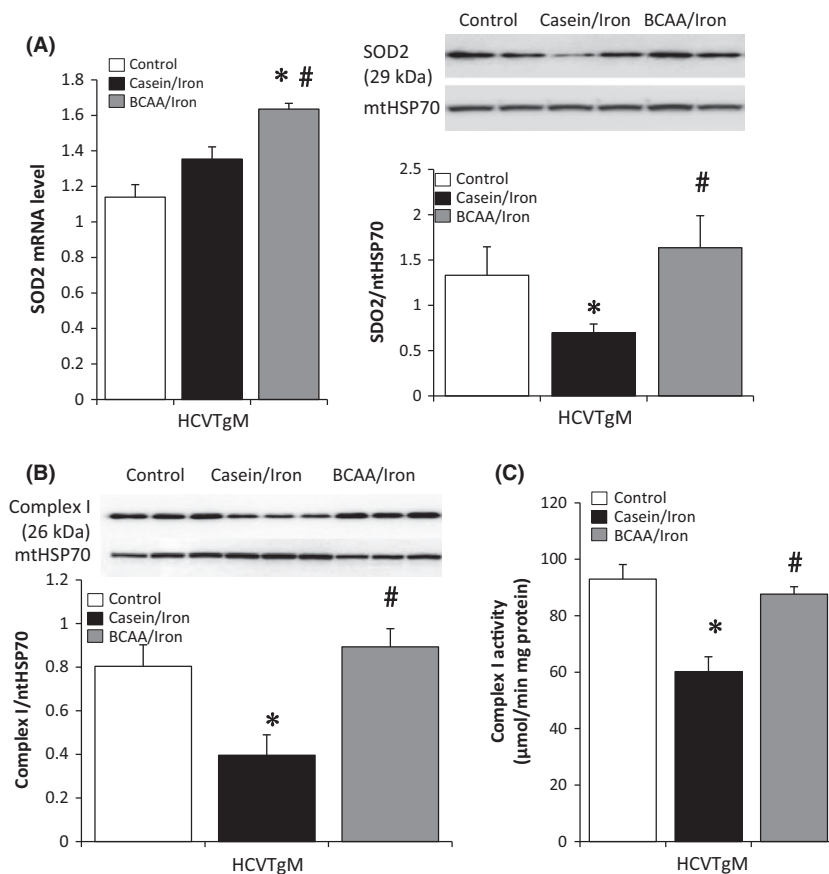
with casein compared with those fed the control diet. The activity was restored by BCAA supplementation (Fig. 4C). Thus, these improvements in the mitochondrial complex I activity and CPT1, SOD2, and mitochondrial complex I expression indicate that BCAA may protect against the mitochondrial injury induced by HCV proteins and iron overload.

#### Antioxidant effects of BCAA supplementation in patients with HCV-related severe fibrosis

Next, we determined whether oral BCAA supplementation reduced oxidative stress and affected iron metabolism in patients with HCV-related advanced liver fibrosis. We assigned 25 patients to receive either BCAA supplementation (BCAA group;  $n = 12$ ) or follow-up without treatment (non-BCAA group;  $n = 13$ ). There were no differences in the clinical characteristics, oxidative stress

markers, or iron metabolic markers at baseline between these groups (Table 2). Serum albumin and AST levels in BCAA group tended to be lower than those in non-BCAA group, although these differences were not statistically significant ( $P = 0.071$  and  $P = 0.074$  respectively).

The dROM levels increased significantly at weeks 24 and 48 in the non-BCAA group, whereas they did not in the BCAA group. The BAP levels also increased at weeks 12 and 24 in the non-BCAA group, and at weeks 12, 24 and 48 in the BCAA group (Table 3). The BAP/dROM ratio, an indicator of antioxidant potential, decreased significantly at week 48 in the non-BCAA group, but increased at weeks 24 and 48 in the BCAA group. This suggests that the BAP levels of the non-BCAA group increased in response to oxidative stress, while the increased BAP levels in the BCAA group indicated enhanced antioxidant potential.



**Fig. 4.** (A) SOD2 mRNA expression in the livers of four mice from each group. (B) Immunoblots of SOD2 and mitochondrial complex I in the mitochondria of four mice from each group after treatment for 6 months. The protein expression levels were normalized against that of mitochondrial heat shock protein 70. \* $P < 0.05$  vs control group; # $P < 0.05$  vs casein/iron group.

In agreement with the antioxidant status, the serum ferritin levels were significantly lower after week 48 of BCAA supplementation ( $137 \pm 109$  mg/dl;  $P < 0.05$ ) compared with those before treatment (Table 3). BCAA supplementation significantly increased the serum hepcidin-25 levels at week 48 ( $20.2 \pm 14.5$  mg/dl;  $P < 0.05$ ). In addition, we determined the level of albumin synthesis after BCAA supplementation, because the oxidized albumin to total albumin ratio increases with cirrhosis progression and it is related to oxidative stress (32,33). In the present study, there were no differences in the total albumin changes in the non-BCAA or BCAA groups (Table 4). However, the amount of albumin present in the reduced form increased significantly in the BCAA group at week 48 compared with that before the study. By contrast, the level of reduced albumin decreased significantly at week 48 in the non-BCAA group. This suggests that long-term BCAA supplementation reduced iron overload by upregulating antioxidant potential and this improved the albumin status in patients without hypoalbuminaemia and chronic liver failure.

## Discussion

Hepatic iron overload and ROS production are both pathophysiological features of HCV-associated chronic liver disease (34) and risk factors for HCC development (35). The reduced hepatic oxidative stress observed after oral BCAA supplementation may be related to changes in the albumin redox state (32, 36). However, previous studies did not determine how BCAA affects iron metabolism and ROS generation.

The mouse model used in the present study shared similarities with the patients who had HCV-associated chronic liver disease in terms of hepatic ROS production and steatosis (14) at 6 months after treatment, followed by hepatocarcinogenesis (15). Furthermore, the hepatic iron concentrations in HCVtgM fed the excess-iron diet were comparable to those of a large number of patients with chronic hepatitis C (30, 37, 38). Thus, HCVtgM fed the excess-iron diet is a suitable model for assessing the effects of long-term supplementation with BCAA on disordered iron metabolism and ROS production in HCV infection.



**Table 2.** Patient baseline characteristics

	non-BCAA	BCAA	P-value
Patients (N)	13	12	N.S.
Age (years)	73.5 (65–87)	74.9 (65–83)	N.S.
Sex (male/female)	6/7	6/6	N.S.
White blood cell count ( $\times 10^2/\text{mm}^3$ )	46.2 (30.1–63.4)	45.1 (27.1–84.5)	N.S.
Haemoglobin concentration (g/dl)	13.3 (10.6–16.3)	13.2 (11.4–15.7)	N.S.
Platelet counts ( $\times 10^4/\text{mm}^3$ )	12.2 (4.9–15)	10.5 (3.7–15)	N.S.
Total bilirubin (mg/dl)	0.7 (0.5–1.1)	1.0 (0.3–1.7)	N.S.
Albumin (g/dl)	4.0 (3.5–4.2)	3.9 (3.5–4.2)	N.S.
ALT (IU/L)	33 (23–47)	41 (21–55)	N.S.
AST (IU/L)	43 (32–54)	48 (32–54)	N.S.
ALP (IU/L)	286 (158–435)	302 (145–491)	N.S.
GTP (IU/L)	46 (15–177)	53 (16–137)	N.S.
FBS (mg/dl)	94 (70–130)	107 (72–158)	N.S.
Insulin ( $\mu\text{U}/\text{ml}$ )	14 (5.3–39)	14 (6.3–28)	N.S.
Tyrosine (nmol/ml)	104 (76–123)	103 (63–149)	N.S.
BCAA (nmol/ml)	424 (319–606)	401 (269–617)	N.S.
BTR	4.1 (2.6–4.9)	4.0 (2.7–4.4)	N.S.
AFP (ng/dl)	11 (2–61)	18 (2–95)	N.S.
Serum iron ( $\mu\text{g}/\text{ml}$ )	134 (50–255)	136 (37–256)	N.S.
TSAT (%)	38 (11–70)	44 (12–88)	N.S.
Ferritin (ng/ml)	120 (30–429)	190 (30–346)	N.S.

Results are mean (range). Comparisons between branched-chain amino acids (BCAA) and non-BCAA groups were made using Levene's or Welch's tests. AFP,  $\alpha$ -foetoprotein; ALP, alkaline phosphatase; ALT, alanine aminotransferase; AST, aspartate aminotransferase; GGT, gamma glutamyltransferase; BTR, the ratio of BCAA relative to tyrosine; FBS, fasting blood sugar; N.S., Not significant; TSAT, transferrin saturation.

BCAA supplementation improves the nutritional status, prognosis and quality of life for patients with cirrhosis (39, 40). A randomized, controlled trial demonstrated that BCAA supplementation reduced the frequency of HCC in obese male patients with cirrhosis and HCV infection (18). BCAA treatment also reduced the hepatocarcinogenic activity in obese diabetic animals with insulin resistance (9, 10). Insulin resistance promotes hepatocarcinogenesis by activating the mitogen-activated protein kinase (MAPK) pathway and insulin-like growth factor 1 (IGF-1) receptors, which further activates the Raf/MAPK kinase/MAPK cascade (41, 42). BCAA suppress the IGF/IGF-1R axis by down-regulating IGF-1, IGF-2 and IGF-1R mRNA expression, thereby leading to the inhibition of mitosis and cell growth (9). BCAA reduce HCC development by inhibiting insulin resistance (43).

In the present study, the FBS levels of HCVTgM fed the excess-iron diet with casein increased after 6 months. BCAA supplementation reduced the iron overload-induced elevation of the FBS. There was no intrahepatic inflammation or fibrosis in the HCVTgM fed the excess-iron diet, but those fed the excess-iron diet with casein had significantly lower plasma BCAA levels and a lower BTR compared with those fed excess-iron with BCAA and the control diet. An amino acid imbalance, which is indicated by a lower BTR, has been observed in patients with compensated cirrhosis or chronic hepatitis (44, 45). This suggests that BCAA might potentially reduce hepatic iron accumulation and ROS in patients with HCV-related advanced fibrosis.

**Table 3.** Changes in oxidative stress and iron metabolism markers during branched-chain amino acids (BCAA) administration\*

	Week 0	Week 12	Week 24	Week 48
Hepcidin (ng/ml)				
non-BCAA	11.6 $\pm$ 7.9	10.4 $\pm$ 9.8	11.8 $\pm$ 9.5	10.5 $\pm$ 8.8
BCAA	9.5 $\pm$ 8.7	9.2 $\pm$ 9.5	11.0 $\pm$ 9.1	20.2 $\pm$ 14.5 <sup>†</sup>
Ferritin (ng/ml)				
non-BCAA	120 $\pm$ 121	112 $\pm$ 105	100 $\pm$ 108	118 $\pm$ 120
BCAA	190 $\pm$ 135	164 $\pm$ 129	163 $\pm$ 130	137 $\pm$ 109 <sup>†</sup>
Serum iron ( $\mu\text{g}/\text{ml}$ )				
non-BCAA	134 $\pm$ 52	143 $\pm$ 60	133 $\pm$ 55	142 $\pm$ 38
BCAA	136 $\pm$ 64	131 $\pm$ 63	134 $\pm$ 68	117 $\pm$ 57
TSAT (%)				
non-BCAA	38 $\pm$ 14	42 $\pm$ 19	39 $\pm$ 15	42 $\pm$ 19
BCAA dROM (U.CARR)	45 $\pm$ 29	42 $\pm$ 24	35 $\pm$ 19 <sup>†</sup>	33 $\pm$ 16 <sup>†</sup>
non-BCAA	342 $\pm$ 64	405 $\pm$ 84	431 $\pm$ 76 <sup>†</sup>	455 $\pm$ 96 <sup>†</sup>
BCAA	360 $\pm$ 113	372 $\pm$ 80	361 $\pm$ 118	359 $\pm$ 65
BAP ( $\mu\text{M}$ )				
non-BCAA	2369 $\pm$ 386	2772 $\pm$ 487 <sup>†</sup>	2798 $\pm$ 337 <sup>†</sup>	2630 $\pm$ 64
BCAA	2139 $\pm$ 587	2516 $\pm$ 678 <sup>†</sup>	2601 $\pm$ 647 <sup>†</sup>	2758 $\pm$ 413 <sup>†</sup>
BAP/dROM				
non-BCAA	7.0 $\pm$ 1.0	7.1 $\pm$ 1.7	6.6 $\pm$ 0.7	6.0 $\pm$ 1.0 <sup>†</sup>
BCAA	6.1 $\pm$ 1.3	6.8 $\pm$ 1.5	7.5 $\pm$ 1.6 <sup>†</sup>	7.8 $\pm$ 1.5 <sup>†</sup>

\*Results are mean  $\pm$  SD.

<sup>†</sup>P < 0.05 vs. before BCAA treatment, Wilcoxon rank-sum test; U.CARR, Cartelli Units (1 U.CARR = 0.8 mg/L of H<sub>2</sub>O<sub>2</sub>), TSAT, transferrin saturation.

**Table 4.** Changes in the serum albumin characteristics during branched-chain amino acid (BCAA) administration\*

	Week 0	Week 12	Week 24	Week 48
Albumin (g/dl)				
non-BCAA (13)	4.0 ± 0.2	4.0 ± 0.2	4.0 ± 0.2	4.0 ± 0.2
BCAA (12)	3.9 ± 0.3	3.8 ± 0.4	3.9 ± 0.3	4.0 ± 0.3
Reduced albumin (%)				
non-BCAA (10)	66 ± 4.5	66 ± 5.3	66 ± 3.9	63 ± 4.9†
BCAA (10)	66 ± 3.9	68 ± 2.8	68 ± 5.2	70 ± 3.2†
Type IV collagen 7s (U/ml)				
non-BCAA (13)	5.8 ± 1.7	6.1 ± 2.5	6.0 ± 2.2	6.1 ± 2.1
BCAA (12)	6.8 ± 2.1	7.1 ± 1.8	7.0 ± 2.1	6.7 ± 2.0
P-III-P (U/ml)				
non-BCAA (13)	0.89 ± 0.19	0.83 ± 0.19	0.91 ± 0.24	0.83 ± 0.15
BCAA (12)	0.88 ± 0.23	0.90 ± 0.16	0.88 ± 0.18	0.86 ± 0.22
Fib4-index				
non-BCAA (13)	5.0 ± 2.7	5.0 ± 2.5	5.4 ± 2.9	5.5 ± 3.7
BCAA (12)	6.8 ± 4.2	7.2 ± 3.7	6.4 ± 4.0	6.3 ± 4.0

\*Results are mean ± SD.

† $P < 0.05$  vs. before treatment, Wilcoxon rank-sum test, P-III-P: Type III procollagen peptide.

Our previous study indicated that the antioxidant N-acetylcysteine (NAC) almost completely blocked ROS production and abrogated the hepatic steatosis induced by HCV proteins and iron (30). In the present study, the hepatic triglyceride levels tended to be lower in mice fed the excess-iron diet with BCAA compared with those fed the excess-iron diet with casein, although these differences were not statistically significant ( $P = 0.055$ ). This may have been because BCAA reduce ROS production to a lesser degree than NAC, or because BCAA supplementation did not completely inhibit the ROS-associated unfolded protein response or improve glucose intolerance compared with the control diet. SREBP1 expression is positively regulated by insulin signalling pathways (46). Therefore, further studies are needed to determine whether BCAA reduce hepatic iron accumulation without affecting hepatic steatosis.

CPT1, a transmembrane enzyme in the mitochondrial outer membrane, is negatively regulated at the transcriptional level by malonyl-CoA, which is an intermediate product of fatty acid synthesis (47). Decreased CPT1 expression may be related to the HCV core protein, which is also located in the mitochondrial outer membrane and it generates mitochondrial ROS production indirectly (13). BCAA enhanced protection against mitochondrial injury by restoring the mitochondrial antioxidant potential and mitochondrial complex I activity. Thus, how does BCAA protect from HCV-induced ROS production and mitochondrial injury? The hepatic ROS production increased more in the HCVtgM fed the control diet compared with non-transgenic mice (14), but we did not test whether BCAA supplements reduced ROS production in HCVtgM without the excess-iron diet. However, there were no differences in the liver enzyme, glucose, insulin, BCAA and tyrosine levels of HCVtgM and non-transgenic mice. Furthermore, HCVtgM without the excess-iron

diet did not develop severe steatosis and HCC. This indicates that HCVtgM without the excess-iron diet are not a suitable model for long-term treatments with BCAA.

BCAA supplementation increases the reduced form of albumin, which is a predictor of the cirrhosis prognosis (32, 33), while it also improves oxidative stress and iron metabolism in patients with decompensated cirrhosis (36), and in rats exposed to a fibrogenic agent (48). This suggests that the antioxidant effects of BCAA may be related to qualitative changes in serum albumin or the upregulation of albumin synthesis (4, 5, 9, 49). BCAA itself activates the mammalian target of rapamycin, which subsequently upregulates the downstream molecules, eukaryotic initiation factor 4E-binding protein-1 and 70-kDa ribosomal protein S6 kinase, thereby regulating mRNA translation and synthesis (50). In the present clinical study, we confirmed that long-term BCAA supplementation increased the BAP/dROM ratios and serum hepcidin-25 levels, whereas it decreased the serum ferritin levels in patients with HCV-related advanced fibrosis. Moreover, we found that BAP continued to increase from week 12 to week 48, and the level of the reduced albumin form increased at week 48, but without changes in the serum albumin levels, in our BCAA group. The mechanism that allows BCAA to protect against HCV and iron-induced oxidative stress remains uncertain, but BCAA may improve iron metabolism by upregulating the antioxidant potential in patients without decompensated cirrhosis.

In addition, amino acid imbalance is a risk factor for HCC development in patients without hypoalbuminemia (40), which suggests that BCAA supplementation should be recommended to patients with amino acid imbalances with advanced fibrosis, who may have a decreased antioxidant potential and reduced albumin level. In this study, we could not show any effect by

which BCAA supplementation prevented fibrotic progression (Table 4). However, Fib-4 index in BCAA group at 48 weeks tended to be decreased compared with those at initial point, although these differences were not statistically significant ( $P = 0.061$ ). Long-term BCAAs treatment might inhibit hepatic fibrosis in HCV patients with advanced fibrosis.

Our clinical study had some limitations, including a higher number of older patients who had higher serum albumin and ferritin levels than those in the cohorts reported in other studies, although they used small sample sizes and were not randomized. Further studies should use large cohorts to clarify these effects.

In conclusion, we demonstrated that BCAA administration reduced the hepatic iron contents and ROS levels, which were induced by HCV proteins and iron overloading in mice, probably by protecting the function of mitochondrial complex I. Furthermore, we confirmed that BCAA supplementation improved disordered iron metabolism and the antioxidant status in patients with HCV-related advanced fibrosis. These effects of BCAA may partially account for their inhibitory effects on HCC development in patients with cirrhosis.

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## Supporting information

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

**Fig. S1.** Twenty five patients with HCV-related advanced fibrosis enrolled Human BCAA supplementation study. Advanced fibrosis defined liver specimens (METAVIR fibrosis staging: >F3,4) or Fib-4 index (>3.25).

**Fig. S2.** HDAC activity of HCV TgM fed the excess-iron diet with BCAA was significantly lower than those of HCV TgM fed the excess-iron diet with casein or the control diet. All samples were nuclear which was extracted from liver tissue.