# Nuclear export of phosphorylated C/EBP $\beta$ mediates the inhibition of albumin expression by TNF- $\alpha$

# Martina Buck<sup>1,2,3</sup>, Lian Zhang<sup>2,4</sup>, Nicholas A.Halasz<sup>4,5</sup>, Tony Hunter<sup>1</sup> and Mario Chojkier<sup>2,4,6</sup>

<sup>1</sup>Molecular and Cell Biology Laboratory, The Salk Institute for Biological Studies, La Jolla, CA 92037, Departments of <sup>2</sup>Medicine and <sup>5</sup>Surgery and <sup>6</sup>Center for Molecular Genetics, University of California, La Jolla, CA 92037 and <sup>4</sup>VA Medical Center, San Diego, CA 92161, USA

<sup>3</sup>Corresponding author e-mail: mbuck@vapop.ucsd.edu

Decreased albumin expression is a frequent feature of cachexia patients afflicted with chronic diseases, including cancer, and a major contributor to their morbidity. Here we show that tumor necrosis-a (TNF- $\alpha$ ) treatment of primary mouse hepatocytes or TNF- $\alpha$  overexpression in a mouse model of cachexia induces oxidative stress, nitric oxide synthase (NOS) expression and phosphorylation of C/EBPB on Ser239, within the nuclear localization signal, thus inducing its nuclear export, which inhibits transcription from the albumin gene. SIN-1, a NO donor, duplicated the TNF- $\alpha$  effects on hepatocytes. We found similar molecular abnormalities in the liver of patients with cancer-cachexia. The cytoplasmic localization and association of C/EBPB-PSer239 with CRM1 (exportin-1) in TNF- $\alpha$ -treated hepatocytes was inhibited by leptomycin B, a blocker of CRM1 activity. Hepatic cells expressing the non-phosphorylatable C/EBP $\beta$ alanine mutant were refractory to the inhibitory effects of TNF- $\alpha$  on albumin transcription since the mutant remained localized to the nucleus. Treatment of TNF-a mice with antioxidants or NOS inhibitors prevented phosphorylation of C/EBPB on Ser239 and its nuclear export, and rescued the abnormal albumin gene expression.

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# Introduction

Albumin is the most abundant protein in plasma, and the colloid pressure of plasma is maintained principally by the levels of circulating albumin (West, 1990). Albumin also performs important metabolic functions in the transport of free fatty acids, bilirubin and many drugs (West, 1990; Chojkier, 1995). In a normal individual, ~15 g of albumin are synthesized daily to maintain the albumin plasma steady-state concentration (~4 g/100 ml) (West, 1990). Therefore, decreased albumin synthesis results in hypo-albuminemia, which facilitates excessive transudation of fluids into extravascular spaces (edema and ascites) (Braunwald, 1994). Hypoalbuminemia is a frequent feature of cachectic patients afflicted with chronic diseases

(Tracey and Cerami, 1993), including cancer, AIDS and inflammatory disorders, and a major contributor to their morbidity (Voth et al., 1990; Beutler, 1992; Grunfeld and Feingold, 1992; Roubenoff et al., 1994). There is strong evidence to suggest that tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) is a critical mediator (Fong et al., 1989; Yoneda et al., 1991; Cheng et al., 1992), in concert with other cytokines (Grunfeld and Feingold, 1992; Strassman et al., 1992; Spiegelman and Hotamisligil, 1993; Todorov et al., 1996), of cachexia. Therefore, we evaluated the mechanisms leading to decreased albumin transcription (Brenner et al., 1990) in a murine model of cachexia (TNF- $\alpha$  mice), induced by chronically elevated serum TNF- $\alpha$  (Oliff *et al.*, 1987; Brenner et al., 1990; Buck and Chojkier, 1996) in hepatocytes and hepatoma cells treated with TNF- $\alpha$  and in patients with cancer-cachexia.

Here we show that TNF- $\alpha$  treatment of primary mouse hepatocytes or TNF-a overexpression in a mouse model of cachexia induces oxidative stress, nitric oxide synthase (NOS) expression and phosphorylation of C/EBP $\beta$  on Ser239, within the nuclear localization signal (NLS), thus inducing its nuclear export, which inhibits transcription from the albumin gene. We found similar molecular abnormalities in the liver of patients with cancer-cachexia. The cytoplasmic localization and association of C/EBPβ-PSer239 with CRM1 (exportin-1) in TNF-α-treated hepatocytes was inhibited by leptomycin B, a blocker of CRM1 activity. Hepatic cells expressing the non-phosphorylatable C/EBPB alanine mutant were refractory to the inhibitory effects of TNF- $\alpha$  on albumin transcription since the mutant remained localized to the nucleus. Treatment of TNF- $\alpha$  mice with antioxidants or NOS inhibitors prevented phosphorylation of C/EBP $\beta$  on Ser239 and its nuclear export, and rescued the abnormal albumin gene expression.

## Results

# Oxidative stress inhibits albumin expression in the liver of TNF- $\!\alpha$ mice

Chinese hamster ovary (CHO) cells stably transfected with either a cytomegalovirus (CMV) vector expressing human TNF- $\alpha/neo$  (TNF- $\alpha$  cells) or *neo* alone (CHO cells, control) (Oliff *et al.*, 1987; Brenner *et al.*, 1990; Buck and Chojkier, 1996) were used in our experiments. The TNF- $\alpha$ cells, but not the CHO cells, secreted human TNF- $\alpha$ . Athymic nude mice were injected intramuscularly with either CHO cells or TNF- $\alpha$  cells. The TNF- $\alpha$  and CHO mice maintained essentially the same weight for the first 2–3 weeks after inoculation. We have reported that albumin mRNA and albumin transcription are markedly decreased in TNF- $\alpha$  mice before the onset of weight loss (Brenner *et al.*, 1990). After this period, TNF- $\alpha$  animals began to develop symptoms of cachexia, including



**Fig. 1.** Increased oxidative stress and NOS2 in the liver of cachectic mice. The experimental groups are as described in Materials and methods. Representative examples (n = 8 in each group) of the immunohistochemistry for malondialdehyde (MDA)–protein adducts and NOS2. PV indicates portal venules. Negligible staining was observed in all immunohistochemistries when the first antibody was omitted.

decreased weight, muscle wasting, anemia and abnormal wound healing (Buck and Chojkier, 1996; Buck *et al.*, 1996).

Because TNF- $\alpha$  may initiate a cascade leading to oxidative stress (Wong *et al.*, 1989; Henkel *et al.*, 1993; Schulze-Osthoff *et al.*, 1993), we assessed whether aldehyde products of lipid peroxidation, such as malondialdehyde (MDA) (Chaudhary *et al.*, 1994), were present in the liver of TNF- $\alpha$  mice. Using specific antibodies against MDA–lysine adducts for immunostaining (Houglum *et al.*, 1990; Buck and Chojkier, 1996), we found a high level of MDA–protein adducts in the liver of TNF- $\alpha$  mice, while the liver of control mice had negligible levels (Figure 1).

Because of this evidence, supporting an oxidative stress pathway in the liver of TNF- $\alpha$  mice, and because aldehydes may form adducts with proteins critical for differentiated function, we treated these animals with antioxidants in an attempt to elucidate the mechanisms responsible for the decrease in albumin gene expression of TNF- $\alpha$  mice (Brenner *et al.*, 1990). When TNF- $\alpha$  mice received a diet supplemented with D- $\alpha$ -tocopherol (Buck and Chojkier, 1996), a lipophilic antioxidant, the increased oxidative stress in the liver was blocked, as determined by the absence of MDA-protein adducts (Figure 1), a sensitive indicator of lipid peroxidation (Houglum *et al.*, 1990). More importantly, the characteristic decrease in albumin mRNA of TNF- $\alpha$  mice was prevented, to a significant extent, by a 30-day treatment with D-α-tocopherol (Brenner et al., 1990) (Figure 2A). We also assessed whether the inhibition of albumin mRNA induced by TNF- $\alpha$  could be rescued in short-term experiments with antioxidants. In normal mice that had received eight highdose TNF- $\alpha$  injections (Buck *et al.*, 1996) to induce a rapid decrease in liver albumin mRNA (Figure 2B), three high-dose D-\alpha-tocopherol injections (Chojkier et al., 1998) were sufficient to rescue the impaired albumin gene expression observed in this animal model (Figure 2B). As expected, the non-lipophilic antioxidant, 3-amino-1-(3-trifluoromethylphenyl)-2-pyrazoline hydrochloride (BW755c) (Buck and Chojkier, 1996), did not affect the generation of MDA-protein adducts (not shown) since it does not alter the peroxidation of polyunsaturated fatty acids in membranes. However, BW755c treatment normalized the albumin mRNA in TNF- $\alpha$  mice (not shown), indicating that the oxidative stress cascade initiated by TNF- $\alpha$  can be blocked effectively at different levels.

The D-site, which is present in both the enhancer and promoter of the albumin gene (Zaret, 1994; Chojkier, 1995), is critical for the expression of liver-specific genes (Maire et al., 1989; Descombes et al., 1990; Zaret, 1994; Chojkier, 1995). Therefore, its binding and transcriptional activities, which are contributed mainly by C/EBPB and C/EBP $\alpha$ , have been used as indicators of liver-specific differentiation (Descombes et al., 1990). We analyzed the D-site binding activities of liver nuclear proteins from the various experimental groups. There was substantially less D-site binding activity in liver nuclear extracts from TNF- $\alpha$  mice than in those from control animals (Figure 2C), and this deficiency was reversed, albeit incompletely, in TNF- $\alpha$  mice treated with either D- $\alpha$ tocopherol or BW775c (Figure 2C). The D-site nuclear protein complex was supershifted by specific anti-C/EBPB antibodies as described previously (Descombes et al., 1990) (data not shown). In contrast, the level of protein binding to the B-site, another important *cis*-element for the efficient transcription of the albumin gene (Maire et al., 1989; Zaret, 1994; Chojkier, 1995), was only slightly decreased in liver nuclear extracts from TNF- $\alpha$  mice (Figure 2C). Liver nuclear extracts from TNF- $\alpha$  mice also showed normal levels of protein binding to the Sp1 site, a site commonly used in housekeeping genes (Figure 2C), indicating a selective defect in the binding of C/EBP $\beta$  to the albumin D-site. The binding of liver nuclear extracts to these labeled cognate DNAs was suppressed by competition with excess unlabeled homologous, but not heterologous, oligonucleotide (data not shown).

Next, we assessed the effects of TNF- $\alpha$  on albumin transcription in HepG2 human hepatoma cells. Because HepG2 cells express negligible amounts of C/EBP $\beta$  (Descombes *et al.*, 1990; Trautwein *et al.*, 1993; Buck *et al.*, 1994), in these experiments, cells were transfected with a CMV vector expressing C/EBP $\beta$  (Buck *et al.*, 1994, 1999) and with an albumin–enhancer (–9 to –12 kb)/ promoter (–282/+28bp) chimeric reporter gene (ALB-CAT) (Kioussis *et al.*, 1979). C/EBP $\beta$  binds to two critical *cis*-elements within the promoter (D-site) and the enhancer of the albumin gene (Descombes *et al.*, 1990; Zaret, 1994;

Chojkier, 1995). As expected, C/EBP $\beta$  stimulated transcription from the ALB-CAT reporter gene (Figure 2D). TNF- $\alpha$  treatment of HepG2 cells markedly inhibited ALB-CAT expression induced by C/EBP $\beta$ , and this effect was blocked with the antioxidants D- $\alpha$ -tocopherol (Figure 2D) or BW755c (data not shown), suggesting that an oxidative stress pathway mediates the inhibitory effects of TNF $\alpha$  on albumin transcription.

#### NO inhibits albumin gene expression

Because oxidative stress pathways interact with NO to modulate cytoprotective or cytotoxic effects (Lipton et al., 1993; Stamler, 1994; Buck and Chojkier, 1996), and NO itself plays a role in liver function (Geller, 1993), we analyzed whether this pathway affects albumin transcription. N<sup>5</sup>-[nitroamidino]-L-2,5-diaminopentanoic acid (nitro-L-arginine), a potent inhibitor of NOS (Kobzik et al., 1994; Buck and Chojkier, 1996), blocked the inhibition of ALB-CAT transcription initiated by TNF- $\alpha$  (Figure 2D). Additional support for a role for NO was obtained by treating the hepatoma cells with 3-morpholinosydnonomine (SIN-1), an NO donor (Lipton et al., 1993; Buck and Choikier, 1996). SIN-1 treatment (Lipton et al., 1993; Buck and Chojkier, 1996) markedly decreased ALB-CAT transcription in HepG2 cells in the absence of TNF- $\alpha$ (Figure 2D).

Because the data derived from HepG2 cells suggested that the NO pathway may mediate (or act synergistically with) the effects of oxidative stress on liver cells, we studied these interactions in TNF- $\alpha$  mice. As shown in Figure 1, the expression of NOS was increased substantially in the livers of TNF- $\alpha$  mice compared with control mice. The increased expression of NOS in TNF- $\alpha$  mice was prevented by treatment with the antioxidants D- $\alpha$ tocopherol (Figure 1) or BW755c. These results support

Fig. 2. Oxidative stress and NO inhibit albumin expression in the liver of cachectic mice. (A) The experimental groups are as described in Materials and methods. Representative examples (n = 6 in each group)of albumin mRNA detected by an RNase protection assay with a specific riboprobe, using equal amounts of total RNA. The 18S RNA was utilized as a correction factor for loading. Albumin mRNA in livers from CHO (lane 1), TNF-α (lane 2), TNF-α/D-α-tocopherol (lane 3) and TNF- $\alpha$ /nitro-L-arginine (lane 4). *P* <0.05 for TNF- $\alpha$ . (B) Mice received TNF- $\alpha$  alone from 8 to 92 h (closed circles) or with D- $\alpha$ -tocopherol from 24 to 72 h (open circles) as described in Materials and methods. Animals were sacrificed as indicated and liver albumin mRNA was determined as in (A). The results are averages of at least triplicate samples. P < 0.05 for D- $\alpha$ -tocopherol at 72 and 96 h. (C) Mobility shift analysis of liver nuclear extracts was performed using equal amounts of nuclear protein (5  $\mu$ g) and following incubation with a <sup>32</sup>P-labeled oligonucleotide (1 ng) spanning the D-site or the B-site of the albumin enhancer/promoter or the control Sp1 site. The position of the bound DNA is indicated by arrows. Representative samples (n = 5 in each group) are shown: CHO (2); TNF- $\alpha$  (3); TNF-α/D-α-tocopherol (4); TNF-α/BW755c (5); and TNF-α/nitro-Larginine (6). On lane 1, the probe was processed without nuclear extracts. The D-site binding activities were CHO (100%), TNF- $\alpha$ (22%), TNF-a/D-a-tocopherol (84%), TNF-a/BW755c (67%) and TNF- $\alpha$ /nitro-L-arginine (76%). (D) HepG2 human hepatoma cells were transfected with ALB-CAT (1  $\mu$ g) alone or with CMV-C/EBP $\beta$  (1  $\mu$ g) and treated every 24 h with TNF- $\alpha$  (10 ng/ml); TNF- $\alpha$  + D- $\alpha$ tocopherol (50  $\mu$ M); TNF- $\alpha$  + nitro-L-arginine (500  $\mu$ M); and SIN-1 (0.6 µM) as indicated. After 48 h, the cells were collected and CAT expression was determined. The results are averages ( $\pm$  SEM) of triplicate samples, and representative of four independent experiments. P < 0.05 for C/EBP $\beta$  + TNF- $\alpha$  and C/EBP $\beta$  + SIN-1 compared with C/EBPβ.



the hypothesis that oxidative stress induces NOS expression in the liver of TNF- $\alpha$  mice.

Treatment of TNF-α mice with the NOS inhibitor, nitro-L-arginine (Buck and Chojkier, 1996), ameliorated the decrease in both albumin mRNA (Figure 2A) and binding of liver nuclear extracts to the albumin D-site (Figure 2C). Nitro-L-arginine did not affect the induction of either MDA-protein adducts or NOS (Figure 1), upstream components of the cascade leading to the synthesis of NO, in the liver of TNF-α mice. Treatment of TNF-α mice with either antioxidants or nitro-L-arginine did not affect the production of TNF-α, or some of the other end-organ biological effects of TNF-α, such as anemia or inhibition of collagen α1(I) expression (Buck and Chojkier, 1996; Buck *et al.*, 1996), indicating a selective effect on specific TNF-α pathways.

# Phosphorylation of C/EBP $\beta$ on its DNA-binding domain mediates the inhibition of albumin transcription by TNF- $\alpha$

C/EBPβ contributes, as homo- and heterodimers, ~70% of the binding activity of liver nuclear proteins to the D-site (Descombes et al., 1990). However, the expression of C/EBPB (LAP, NF-IL6, IL-6DBP) protein (Akira et al., 1990a; Descombes et al., 1990; Poli et al., 1990) was not decreased in whole liver lysates from TNF- $\alpha$  mice (Figure 3A). This observation could be reconciled with the decreased D-site binding activities of liver nuclear extracts from TNF- $\alpha$  mice, if TNF- $\alpha$  stimuli were to induce a modification of C/EBPB, such as phosphorylation, that impairs its function. Therefore, we investigated the role of C/EBP $\beta$  phosphorylation in the inhibition of albumin transcription induced by TNF- $\alpha$  in primary mouse hepatocytes. As in normal adult liver, these quiescent hepatocytes expressed nuclear C/EBPB (Buck et al., 1999).

To test whether TNF- $\alpha$  induces C/EBP $\beta$  phosphorylation, highly differentiated, quiescent mouse hepatocytes cultured on a collagen type I matrix (Buck et al., 1999) were incubated with 30 mCi of [32P]orthophosphate before treatment for 30 min with TNF- $\alpha$ . Phosphopeptide mapping of immunoprecipitated endogenous C/EBPB showed that TNF- $\alpha$  induced a major site-specific phosphorylation of C/EBP $\beta$  on Ser239, as well as other, as yet unidentified, sites (Figure 3B). Phosphorylation of Ser240 in rat C/EBPB (homologous to mouse PSer239), which was identified previously using recombinant wild-type and mutant C/EBPβ-Ala240 and activated protein kinase C (Trautwein et al., 1994), was shown to inhibit in vitro binding of C/EBPB to cognate DNA (Trautwein et al., 1994). This serine phosphoacceptor, within the DNAbinding domain and NLS of C/EBPB (Williams et al., 1997), is identical in rat, mouse, chicken, bovine and human, as well as in C/EBPa (Akira et al., 1990a; Descombes et al., 1990; Cao et al., 1991; Katz et al., 1993; Yamaoka et al., 1997) (Figure 3C). Endogenous C/EBPβ was not phosphorylated on Ser239 in control hepatocytes (Figure 3B). Furthermore, nuclear extracts from these mouse hepatocytes treated for 30 min with TNF- $\alpha$ displayed decreased binding activity for the albumin D-site (Figure 4A), which was restored (data not shown) by phosphatase treatment of the nuclear extracts as described previously (Trautwein et al., 1994).



Fig. 3. TNF- $\alpha$  stimulates phosphorylation of C/EBP $\beta$  in its DNAbinding domain. (A) Representative examples (n = 5 in each group) of C/EBPß protein immunoblots in liver lysates from CHO (lane 1) and TNF- $\alpha$  (lane 2) groups. The antigen-antibody complexes were visualized using the Renaissance detection system (DuPont). (B) Tryptic phosphopeptide maps of C/EBPß immunopurified from day-5 primary mouse hepatocytes cultured on a collagen type I matrix. Cells were labeled with [32P]orthophosphate (2 mCi/ml; total 30 mCi) for 18 h and either treated with TNF- $\alpha$  (10 ng/ml) for 30 min before harvest or not treated (control). C/EBPB tryptic peptides were separated by highvoltage electrophoresis (horizontal dimension) followed by ascending thin-layer chromatography (vertical dimension). The level of the phosphopeptide containing Ser239 (arrowhead), identified as described previously (Trautwein et al., 1994), was increased after treatment with TNF- $\alpha$ . (C) The basic domain serine phosphoacceptor is conserved through evolution. Rat (S240), mouse (S239), chicken (S271), bovine (S291) and human (S288) phosphoacceptors, within the basic domain of C/EBPB, are shown. An identical phosphoacceptor is present in human (h) C/EBPa (S299).

Because phosphorylation of C/EBP $\beta$  within its basic domain impaired binding of this transcription factor to cognate DNA (Figure 4A), we analyzed the role of this phosphoacceptor in albumin gene expression. In HepG2 cells, the rat C/EBP $\beta$ -Ala240 mutant protein, which lacks the Ser240 phosphoacceptor (Trautwein *et al.*, 1994), stimulated albumin transcription as much as wild-type C/EBP $\beta$  but was refractory to the inhibitory effects of TNF- $\alpha$  (Figure 4B). These findings suggest that phosphorylation of rat C/EBP $\beta$  on Ser240 is critical for the TNF- $\alpha$  signal transduction pathway to cause inhibition of albumin transcription. The effects of Ser240 phosphorylation of rat C/EBP $\beta$  on its DNA binding affinity can be mimicked by introduction of a negatively charged residue (Trautwein *et al.*, 1994). As expected, cells expressing the mutant C/EBP $\beta$ -Asp240 protein exhibited minimal transcription from the ALB-CAT reporter gene (Figure 4B).

Next, we developed antibodies against a peptide containing the phosphorylated Ser239 domain, that recognized C/EBP<sub>β</sub>-PSer239 but not unphosphorylated C/EBP<sub>β</sub>. These antibodies were purified by affinity chromatography on the PSer239 phosphoacceptor peptide (CIAVRK-PSer239-RDKAK). Immunohistochemical staining with purified anti-C/EBPB-PSer239 antibodies was positive for this epitope in liver sections from TNF- $\alpha$  mice, but negative in liver sections from control and D-α-tocopherolor nitro-L-arginine-treated TNF- $\alpha$  mice (Figure 4C). Unexpectedly, the localization of C/EBPB was almost exclusively cytoplasmic, correlating with Ser239 phosphorylation in TNF- $\alpha$  mice, whereas C/EBP $\beta$  was mainly nuclear in control and D-\alpha-tocopherol- or nitro-L-argininetreated TNF- $\alpha$  mice, correlating with lack of Ser239 phosphorylation. In contrast to the findings in TNF- $\alpha$  mice, following the administration of lipopolysaccharide (LPS), another inducer of inflammation (Akira et al., 1990a; Alonzi et al., 2001), to normal mice, C/EBPB expression, although increased, was predominantly nuclear and not phosphorylated on Ser239 (Figure 4C).

To delineate the mechanisms responsible for the nuclear exclusion of C/EBP $\beta$ -PSer239, we transfected primary mouse hepatocytes isolated from C/EBP $\beta$ -/- mice (Screpanti *et al.*, 1995; Buck *et al.*, 1999) with CMV vectors expressing mouse C/EBP $\beta$ -hemagglutinin (HA), C/EBP $\beta$ -Ala239-HA, or C/EBP $\beta$ -Asp239-HA. Using confocal microscopy, we detected C/EBP $\beta$  mainly in the nucleus, and CRM1 (exportin-1), a nuclear export receptor (Ossareh-Nazari *et al.*, 1997; Kudo *et al.*, 1999; Nachury and Weis, 1999), in a nuclear and perinuclear localization in hepatocytes (Figure 5A). Following treatment for 30 min with TNF- $\alpha$ , C/EBP $\beta$  was mainly cytoplasmic and colocalized in the perinuclear region with CRM1. Treatment of hepatocytes with leptomycin B, a blocker of protein

Fig. 4. Phosphorylation of C/EBPB in its basic domain induced by TNF- $\alpha$  inhibits albumin expression. (A) Mobility shift analysis of nuclear extracts from day 5 primary mouse hepatocytes was performed using equal amounts of nuclear protein  $(5 \ \mu g)$  and following incubation with a <sup>32</sup>P-labeled oligonucleotide (1 ng) spanning the D-site of the albumin enhancer/promoter. The position of the bound DNA is indicated by the arrow and the supershifted DNA is indicated by the arrowhead. Representative samples are shown: control (2); TNF- $\alpha$ (10 ng/ml for 30 min) (3); and control + C/EBPB antibody (4). On lane 1, the probe was processed without nuclear extracts. (B) HepG2 cells were transfected with ALB-CAT (1 µg) alone or with 1 µg of CMV-C/EBPβ, CMV-C/EBPβ-Ala240 or CMV-C/EBPβ-Asp240, and treated with TNF- $\alpha$  (10 ng/ml) every 24 h as indicated. After 48 h, the cells were collected and CAT expression was determined. The results are averages (± SEM) of triplicate samples, and representative of five independent experiments. P < 0.05 for C/EBP $\beta$  + TNF- $\alpha$  and C/EBP $\beta$ -Asp240 compared with C/EBP $\beta$ . (C) Representative examples (n = 8 in each group) of the triple channel immunofluorescence microscopy using antibodies for C/EBPB and C/EBPB-PSer239 simultaneously. Hepatocytes from control, TNFα/D-α-tocopherol-, TNF-α/nitro-Larginine- and LPS-treated mice displayed nuclear C/EBPB (in red) but not C/EBP $\beta$ -PSer-239 (in green). In hepatocytes from TNF- $\alpha$  livers, the cytoplasmic co-localization of antibodies against C/EBP $\beta$  (red) and C/EBPβ-PSer-239 (green) is shown in yellow. Nuclei are stained with DAPI (blue); co-localization of C/EBPB (red) and DAPI (blue) is shown in white.

nuclear export through its interactions with CRM1 (Kudo *et al.*, 1999), prevented the effects of TNF- $\alpha$  on C/EBP $\beta$  nuclear export (Figure 5A), but nuclear C/EBP $\beta$  was still phosphorylated on Ser239 judging by the positive staining with anti-PSer239 antibodies (data not shown; see below). Hepatocytes treated only with leptomycin B displayed unphosphorylated, nuclear C/EBP $\beta$  (data not shown). These experiments indicate that the cytoplasmic localization of C/EBP $\beta$ -PSer239 induced by TNF- $\alpha$  is mediated



by CRM1. Like wild-type C/EBPβ, the non-phosphorylatable C/EBPβ-Ala239 mutant displayed a predominantly nuclear localization but it was refractory to the induction of nuclear export by TNF- $\alpha$  (Figure 5B). Furthermore, the phosphorylation-mimic C/EBPβ-Asp239 mutant was excluded from the nucleus and co-localized in the perinuclear region with CRM1 (Figure 5B), closely resembling the pattern of C/EBPβ-PSer239 induced by TNF- $\alpha$  treatment in both mice and cells. These data strongly suggest that phosphorylation of Ser239 is required for the nuclear export of C/EBPβ following TNF- $\alpha$  treatment.

Because mouse C/EBP $\alpha$  also has the same phosphoacceptor (Ser300) (Christy *et al.*, 1991) and because C/EBP $\alpha$  also plays an important role in liver-specific gene transcription (Landschulz *et al.*, 1988), we analyzed the effects of TNF- $\alpha$  on C/EBP $\alpha$ , in primary hepatocytes isolated from C/EBP $\beta$ -/- mice as described previously (Buck *et al.*, 1999). C/EBP $\alpha$  was found mainly in the nucleus of control C/EBP $\beta$ -/- hepatocytes (Figure 5C). However, following TNF- $\alpha$  treatment, C/EBP $\alpha$  was excluded from the nucleus and co-localized predominantly in the perinuclear region with CRM1 (Figure 5C). These results indicate that C/EBP $\alpha$  is also susceptible to the TNF- $\alpha$  induction of both phosphorylation within its NLS (Williams *et al.*, 1997) and nuclear export.

The mechanisms affecting C/EBP $\beta$  following TNF- $\alpha$  treatment of hepatocytes appear to involve CRM1. Therefore, we analyzed whether CRM1 was associated with C/EBP $\beta$  in primary mouse C/EBP $\beta$ -/- hepatocytes by immunoprecipitation of C/EBP $\beta$  followed by immunoblotting with anti-CRM1 antibodies. Consistent with the confocal microscopy studies, immunoprecipitation of C/EBP $\beta$ -HA from TNF- $\alpha$ -treated hepatocytes immunoprecipitated with antibodies against HA was associated with CRM1 (Figure 6). This association was negligible in control and in TNF- $\alpha$ /leptomycin B-treated hepatocytes (Figure 6). Moreover, phosphorylation of C/EBP $\beta$  on

Fig. 5. Nuclear export of C/EBPβ-PSer-239 is mediated by CRM1 in hepatocytes treated with TNF $\alpha$ . Day 5 primary hepatocytes were isolated form C/EBPβ-/- mice. Representative confocal microscopy (n = 5 in each group). (A) Mouse hepatocytes were transfected with C/EBPB, treated as described in Materials and methods and stained with antibodies against HA and CRM1. In C/EBP $\beta$  and TNF- $\alpha$  + leptomycin B cells, C/EBPB (red) is nuclear and CRM1 (green) is nuclear and perinuclear. In TNF-a cells, C/EBPB and CRM1 are colocalized in the cytoplasm (in yellow). Nuclei are stained with TOTO-3 (blue). (B) Mouse hepatocytes were transfected with CMV vectors expressing C/EBPβ-Ala239 or C/EBPβ-Asp239 for 24 h, treated as described in Materials and methods and stained with antibodies against HA and CRM1. C/EBPβ-Ala239 localized to the nucleus (in red) in untreated and TNF-α-treated hepatocytes. C/EBPβ-Asp239 co-localized with CRM1 in the perinuclear region (in yellow). Nuclei are stained with TOTO-3 (blue). (C) Cells were stained with antibodies against C/EBPa and CRM1. Control hepatocytes expressed nuclear C/EBPa. TNF-α treatment induced the nuclear export of C/EBPα, which colocalized with CRM1 in the perinuclear region (in yellow). Nuclei are stained with TOTO-3 (blue).  $(\mathbf{D})$  Mouse hepatocytes were treated with TNF-a or SIN-1 as described in Materials and methods, and stained with antibodies against HA and PSer239. TNFa and SIN-1 treatments induced the phosphorylation of C/EBPB on Ser239 and its nuclear export. The cytoplasmic co-localization of C/EBPβ-PSer239 (green) and C/EBPB (red) antibodies is shown in yellow. Control hepatocytes displayed mainly nuclear C/EBPB (red) but not C/EBPB-PSer239. Nuclei are stained with TOTO-3 (blue).

Ser239 was induced by TNF- $\alpha$  (Figure 6), as we determined by phosphopeptide mapping (Figure 3B) and by immunohistochemistry (Figure 5D). Also, TNF- $\alpha$  treatment of C/EBP $\beta$ -/- hepatocytes induced the phosphorylation of endogenous C/EBP $\alpha$  on Ser300 (Christy *et al.*, 1991) (homologous to human C/EBP $\alpha$  Ser299), as detected with anti-PSer239 antibodies, as well as the association of C/EBP $\alpha$  with CRM1 (Figure 6). In addition,





Fig. 6. C/EBP $\beta$ -PSer239 is associated with CRM1 in hepatocytes treated with TNF- $\alpha$ . Day 5 primary mouse C/EBP $\beta$ -/– hepatocytes were transfected with C/EBP $\beta$ -HA or not. After 24 h, cells were treated for 30 min as indicated. Cell lysates (500 µg) were precipitated with either HA or C/EBP $\alpha$  antibodies. Protein blotting of immuno-precipitates was performed as described in Materials and methods, using specific antibodies against CRM1, C/EBP $\beta$ -PSer239, C/EBP $\beta$  or C/EBP $\alpha$ .

leptomycin B decreased the phosphorylation of C/EBP $\beta$  on Ser239 induced by TNF- $\alpha$  (Figure 6).

Because inducible NOS (NOS2) expression is increased in the livers of TNF- $\alpha$  mice, and nitro-L-arginine, a blocker of NOS, prevents the phosphorylation of C/EBPβ and its nuclear export (Figure 4C), as well as the inhibition of albumin gene expression (Figure 2) induced by TNF- $\alpha$ , we treated primary mouse hepatocytes with SIN-1, a NO donor, and assessed the cellular localization of C/EBPB using confocal microscopy. SIN-1, like TNF- $\alpha$ , was sufficient to induce the phosphorylation of C/EBPB on Ser239 and its cytoplasmic localization. The cytoplasmic co-localization of C/EBPB-PSer239 and C/EBPB in TNF- $\alpha$ - and SIN-1-treated cells is depicted in yellow by an overlay of both channels (Figure 5D). These results suggest that NO, following its induction by TNF- $\alpha$ , is sufficient to stimulate the phosphorylation of C/EBPB on Ser239 and its cytoplasmic localization.

Given that NO may activate guanylyl cyclase, which in turn increases cGMP levels and the cGMP-dependent protein kinase II (PKG) (El-Husseini *et al.*, 1998; Idriss *et al.*, 1999), we analyzed whether this signal transduction pathway mediates phosphorylation of C/EBP $\beta$  on Ser239. We found no association of C/EBP $\beta$  with PKG in livers from TNF- $\alpha$  mice or in TNF- $\alpha$ -treated hepatocytes (data not shown), suggesting that PKG is not the protein kinase that phosphorylates C/EBP $\beta$  following TNF- $\alpha$  signaling.

# Enhanced oxidative stress and NOS expression in the liver of patients with cancer-cachexia

To assess the relevance to human cachexia of the cellular and animal models of cachexia, we analyzed, in preliminary studies, the mechanisms leading to decreased albumin gene expression in six patients with cancercachexia (see Materials and methods). These patients ( $62 \pm 5$  years) had low serum albumin ( $2.5 \pm 0.4$  g/dl; P < 0.05). Liver biopsies from cachectic patients afflicted with cancer also displayed a high level of both MDA-protein adducts and NOS2 expression (Figure 7A). Although whole lysates of liver biopsies from cachectic patients had normal amounts of C/EBP $\beta$  protein



Fig. 7. Nuclear export of C/EBPβ-Ser288 in the liver of patients with cancer-cachexia. (A) Representative immunohistochemistry for MDA-protein adducts, NOS2, C/EBPB and C/EBPB-PSer288 of control individuals (control) and cancer-cachexia patients (cachexia) was performed as described in Figures 1 and 4.  $(\hat{\mathbf{B}})$  A representative C/EBPß protein immunoblot in C/EBPß immunoprecipitates from liver protein lysates (250 µg) of control (lane 1) and cancer-cachexia (lane 2) subjects was performed as described in Materials and methods. (C) Mobility shift analysis of liver nuclear extracts (5 µg of protein) and the <sup>32</sup>P-labeled D-site of the albumin enhancer/promoter (1 ng) was performed as described in Figure 2. The position of the bound DNA is indicated by an arrow. Samples shown are: control (lanes 2 and 4); cancer-cachexia (lanes 3 and 5). On lane 1, the probe was processed without nuclear extracts. (D) Representative immunohistochemistry using antibodies for C/EBPB (in red) and C/EBPB-PSer288 (in green). simultaneously. In control liver, C/EBPB was localized in the nucleus and C/EBPB-PSer288 was undetectable. In cancer-cachexia liver, C/EBPB-PSer288 was detected in the cytoplasm in yellow due to the superimposition of C/EBPB (red) and C/EBPB-PSer288 (green).

(Figure 7B), there was a substantial decrease in the DNAbinding activity of nuclear extracts to the D-site (Figure 7C), congruent with the abnormal cytoplasmic localization of C/EBPB (Figure 7D) compared with samples from four control patients (67  $\pm$  4 years; serum albumin  $3.7 \pm 0.2$  g/dl). We detected cytoplasmic C/EBPβ-PSer288 (homologous to mouse C/EBPβ-PSer239) in the cytoplasm of hepatocytes from patients with cancer-cachexia, while in control subjects C/EBPB was mainly nuclear and not phosphorylated on Ser288 (Figure 7D). The impaired D-site binding activities of nuclear extracts from cachectic patients were normalized by the addition of recombinant C/EBPB or nuclear extracts from control individuals (data not shown). The binding of liver nuclear extracts to the cognate DNA was suppressed by competition with excess unlabeled homologous, but not heterologous, oligonucleotide.

### Discussion

In this study, we have shown that in normal hepatocytes, TNF- $\alpha$  induces both a site-specific phosphorylation of C/EBPB within the NLS and its cytoplasmic localization. This phosphorylation is required for C/EBPB's cytoplasmic localization, which is mediated by CRM1 and results in decreased transcription from the albumin enhancer/ promoter. Our results provide a relevant physiological example of mammalian transcriptional regulation exerted through phosphorvlation-dependent nucleocytoplasmic protein transport (Hogan and Rao, 1999). Inhibition of albumin synthesis is a common finding in patients with cachexia of cancer, AIDS and diseases characterized by chronic inflammation, and a major contributor to the morbidity of these diseases (Tracey and Cerami, 1993). Because the TNF- $\alpha$  mouse model of cachexia closely resembles human cachexia (Oliff et al., 1987: Brenner et al., 1990; Tracey et al., 1990), it provides a valuable system to analyze the molecular mechanisms responsible for inhibition of albumin synthesis. The TNF- $\alpha$  serum levels in TNF- $\alpha$  mice are only moderately increased (100-300 pg/ml) at the onset of weight loss (Buck and Chojkier, 1996) but, at the time of sacrifice, values are similar to those found in patients with trauma or infectious, parasitic and neoplastic diseases (Scuderi et al., 1986; Waage et al., 1987; Grau et al., 1989; Goodman et al., 1990).

We demonstrated the presence of MDA-protein adducts in the livers of TNF- $\alpha$  mice, indicating activation of an oxidative pathway (Houglum et al., 1990; Chaudhary et al., 1994). These findings are in agreement with evidence that TNF- $\alpha$  can stimulate oxidative stress in many cells and tissues (Wong et al., 1989; Schulze-Osthoff et al., 1993). In addition, NOS2 expression was induced markedly in the livers of TNF- $\alpha$  mice. This effect was rescued by treating these animals with the antioxidants D-a-tocopherol or BW755c, indicating that the liver induction of NOS2 in TNF- $\alpha$  mice is mediated by an enhanced oxidative stress. In addition, in primary mouse hepatocytes, SIN-1, a NO donor, was sufficient to induce the phosphorylation of C/EBPβ on Ser239 within the NLS, and its nuclear export. NO plays an important role in redox signaling by interacting with superoxide to generate peroxynitrite (Lipton et al., 1993; Stamler, 1994) and by nitrosylation

of mitochondrial complex I (Clementi *et al.*, 1998). NOS expression is increased markedly in both the liver of patients with chronic viral hepatitis and hepatoma cells transfected with the hepatitis B virus cDNA (Majano *et al.*, 1998). Therefore, NO may also mediate the inhibition of albumin expression in chronic viral hepatitis. In addition, wild-type p53 and tumor-derived p53 mutants can repress C/EBP $\beta$ -mediated transactivation of the albumin promoter (Kubicka *et al.*, 1999).

The phosphorylation of C/EBPB on Ser239 and its cytoplasmic localization, as well as the decreased albumin gene expression in TNF- $\alpha$  mice, were reversed by treating these animals with antioxidants (D- $\alpha$ -tocopherol or BW755c) or a NOS inhibitor (nitro-L-arginine), indicating that oxidative pathways and activation of NOS are critical for the phosphorylation of C/EBPB on Ser239 and the inhibition of albumin transcription in TNF- $\alpha$  mice (Brenner et al., 1990). Neither the supplemental dose of D- $\alpha$ -tocopherol nor the nitro-L-arginine treatment utilized in our study were toxic to control or TNF- $\alpha$  mice (Buck and Chojkier, 1996; Chojkier et al., 1998) for up to 8 weeks. The effects of antioxidants and nitro-L-arginine were not the spurious result of decreased synthesis of TNF- $\alpha$ , since they affected neither the secretion of biologically active TNF- $\alpha$  by these cells, nor the serum levels of TNF- $\alpha$  in TNF- $\alpha$  animals (Buck and Chojkier, 1996). This pathway may include activation of other cytokines such as interleukin (IL)-1ß and IL-6 (Dinarello et al., 1986; Akira et al., 1990b; Schulze-Osthoff et al., 1993), which in turn could contribute to the inhibition of albumin gene expression in cachexia (Flores et al., 1989; Fong et al., 1989; Strassman et al., 1992; Spiegelman and Hotamisligil, 1993). However, LPS administration to mice, another inducer of inflammation, resulted in the increased nuclear expression of C/EBPB, which remained unphosphorylated on Ser239. The induction of C/EBPB mRNA, C/EBPß protein in the nucleus and C/EBPß binding activities by LPS has been reported previously (Akira et al., 1990a; Alonzi et al., 2001).

Although binding of the albumin enhancer/promoter D-site (Zaret, 1994; Chojkier, 1995) by liver nuclear proteins from TNF- $\alpha$  mice and cachectic patients was substantially decreased, the total hepatocyte expression of C/EBPB, a major D-site activator protein (Akira et al., 1990a; Poli et al., 1990; Trautwein et al., 1993; Chojkier, 1995), remained unchanged. We demonstrated that in highly differentiated, quiescent primary mouse hepatocytes, TNF- $\alpha$  was able to stimulate phosphorylation of endogenous mouse C/EBPB on Ser239 within its NLS (Williams et al., 1997), which inhibits C/EBPB's characteristic nuclear localization in normal hepatocytes (Descombes et al., 1990; Buck et al., 1994), and its binding to cognate DNA sequences necessary for high level transcription from the albumin gene (Maire et al., 1989; Descombes et al., 1990; Trautwein et al., 1993; Chojkier, 1995). Serine phosphorylation of the Saccharomyces cerevisiae transcription factor SWI5 within the NLS prevents its nuclear import (Moll et al., 1991). In addition to the neutralization by phosphorylation on Ser239 of the mouse C/EBPB NLS, C/EBPB also has a sequence within the leucine zipper domain (L<sup>271</sup>-SRE-L-ST-L-RN-L) that closely conforms to the consensus leucine-rich nuclear export signal (NES) (Mattaj and Englmeir, 1998). Our results suggest that this putative NES becomes available for interaction with CREM1 following the phosphorylation of C/EBPB's NLS induced by TNF- $\alpha$  or NO. Cells expressing the non-phosphorylatable rat C/EBPβ-Ala240 mutant (homologous to mouse C/EBPß Ala239) were refractory to the inhibitory effects of TNF- $\alpha$  on albumin transcription. As expected, TNF- $\alpha$  did not induce the cytoplasmic localization of the non-phosphorylatable mutants of rat C/EBPB-Ala240 or mouse C/EBPB-Ala239. Nuclear proteins from cells expressing the phosphorylation-mimic rat C/EBPβ-Asp240 mutant, which, like phosphorylated C/EBPB did not localize to the nucleus, displayed negligible binding to the D-site and these cells did not transcribe albumin reporter chimeric genes. There are other examples of how modification of nucleocytoplasmic transport can regulate gene expression (for reviews see Mattaj and Englmeir, 1998; Hogan and Rao, 1999).

In our experiments, C/EBPB's normal nuclear localization in differentiated hepatocytes was disrupted by TNF- $\alpha$  in both mice and primary cell cultures. When C/EBP $\beta$  is phosphorylated on Ser239 by TNF- $\alpha$  (or by its mediator, NO), it is found in the cytoplasm. This could be the result of either impaired nuclear import or enhanced nuclear export of C/EBPB. However, our studies suggest that the latter is a probable explanation for the following reasons: (i) in hepatocytes, a 30 min TNF- $\alpha$  or SIN-1 treatment was sufficient to redistribute C/EBPB from the nucleus to the cytoplasm, before protein synthesis and impaired nuclear import could contribute significantly to the cytoplasmic localization; (ii) in TNF- $\alpha$  mice, although C/EBPB was localized predominantly in the cytoplasm of hepatocytes, the total hepatic C/EBPB was unchanged, ruling out increased nuclear degradation of C/EBPB as an explanation; (iii) C/EBPB's nuclear localization is conserved in TNF- $\alpha$ -treated hepatocytes, by blocking with leptomycin B the activity of CRM1, which is critical for nuclear protein export (Ossareh-Nazari et al., 1997; Kudo et al., 1999); and (iv) in TNF- $\alpha$ -treated hepatocytes, C/EBPB was co-localized in the cytoplasm and associated with CRM1. Nonetheless, the inhibition of CRM1 by leptomycin B was associated with a more moderate phosphorylation of C/EBPB on Ser239 compared with TNF- $\alpha$ -treated cells, suggesting that the impaired nuclear export of C/EBPB-PSer239 may allow dephosphorylation of C/EBP $\beta$ -PSer239 by a protein phosphatase within the nucleus. Alternatively, the appropriate localization and/or activity of the as yet unidentified TNF- $\alpha$ -induced protein kinase for C/EBPB-Ser239 could be decreased by leptomycin B.

It is of interest that, in transformed rat hepatocyte and human colorectal cancer cell lines, which exhibit the abnormal absence of nuclear C/EBP $\beta$ , TNF- $\alpha$  and antioxidants, respectively, promote the nuclear import of C/EBP $\beta$  (Yin *et al.*, 1996; Chinery *et al.*, 1997), indicating a reversal of the normal nucleocytoplasmic transport in transformed and cancer cell lines through as yet unidentified mechanisms. Furthermore, in the human colorectal cancer cell line, activation of PKA leads to phosphorylation of C/EBP $\beta$  on Ser299, which is required for C/EBP $\beta$ 's nuclear translocation (Chinery *et al.*, 1997). Nonetheless, following the administration of LPS or partial hepatectomy, there is an increase in the nuclear expression of C/EBP $\beta$  and of the transcriptional repressor C/EBP $\beta$  21 kDa (LIP) (Akira *et al.*, 1990a; Diehl and Yang, 1994; Alonzi *et al.*, 2001), which may be physiologically relevant for liver regeneration induced by hepatectomy and acute phase reaction induced by LPS. In contrast to the partial hepatectomy model, CCl<sub>4</sub>induced hepatic regeneration following hepatocellular necrosis and inflammation (a common feature of human liver disease) (Houglum *et al.*, 1995; Buck *et al.*, 1999; Rudolph *et al.*, 2000) is associated with a marked decrease in D-site binding affinity of liver nuclear extracts (Mueller *et al.*, 1990).

Why do TNF- $\alpha$  mice, but not C/EBP $\beta$ -/- mice (Screpanti et al., 1995; M.Buck, unpublished data), have a decreased expression of albumin? Based on our results, there are two mechanisms that can explain the apparent discrepancy. One would expect other C/EBPs expressed in hepatocytes to substitute, but because C/EBPB-PSer239 has a normal leucine zipper domain, it could act in hepatocytes of TNF- $\alpha$  mice as a dominant-negative by forming homo- and heterodimers with leucine zipper proteins, including other C/EBPs (Descombes et al., 1990; Cao et al., 1991), and inducing their nuclear export. Moreover, C/EBP $\alpha$ , which is expressed (Greenbaum *et al.*, 1998) and functions as a transcription factor (Wang et al., 1995; Greenbaum et al., 1998) normally, under basal conditions, in C/EBPB-/- mice (Screpanti et al., 1995) is also a target of the TNF- $\alpha$  signal transduction pathway, given the presence of the conserved serine phosphoacceptor in its DNA-binding domain (Cao et al., 1991). We found that TNF- $\alpha$  also stimulates the phosphorylation on Ser300 and the nuclear export of C/EBPa in C/EBPB-/hepatocytes.

Our preliminary findings in patients with cancercachexia indicate that the cascade leading to decreased albumin gene expression also involves oxidative stress, NOS2 expression, phosphorylation of C/EBP $\beta$  on Ser288 (the human homolog to mouse Ser239), impaired nuclear localization of C/EBP $\beta$  and albumin-binding activities. Therefore, this study provides insights into the mechanisms responsible for the decreased albumin expression in cachexia, which may in turn lead to novel therapeutic approaches for patients with cancer, AIDS and chronic inflammatory diseases.

## Materials and methods

### Mouse model of cachexia and LPS injection

CHO cells stably transfected with either the human TNF- $\alpha$  gene cloned into a CMV mammalian expression vector (TNF-α cells) or with the mammalian expression vector alone (CHO cells, control) were grown in Dulbecco's modified Eagle's essential medium (DMEM) supplemented with 10% fetal calf serum. The TNF- $\alpha$  cells, but not the CHO cells, produced TNF-a. Four-week-old male homozygous nude mice were injected intramuscularly with either  $3.5 \times 10^6$  CHO cells or  $3.5 \times 10^6$ TNF-a cells as described (Brenner et al., 1990; Buck and Chojkier, 1996). Animals in the treatment groups received D-α-tocopherol (8 IU/g of foodstuff), BW755c (10 mg/kg in 100 µl of 1% sucrose, twice a day orally) or nitro-L-arginine (50 µg/ml of drinking water) (Buck and Chojkier, 1996). Animals had free access to food and water and they were sacrificed at 30 days post-inoculation. Also, 4-week-old male heterozygous nude mice received hTNF-a (R&D) (500 ng, intramuscularly every 12 h from 8 to 92 h) alone or with D- $\alpha$ -tocopherol acetate (150 IU, intraperitoneally every 24 h from 24 to 72 h). Animals were sacrificed throughout a 96 h period. TNF-α levels in cell culture media and mouse sera were measured by a biological cytolytic assay and by an enzymelinked immunosorbent assay (ELISA), using monoclonal antibodies against human TNF- $\alpha$  (Brenner *et al.*, 1990; Buck and Chojkier, 1996). LPS (*Escherichia coli* serotype 026:B6; Sigma, St Louis, MO) (1 mg/kg body weight) was administered intraperitoneally and mice were sacrificed after 9 h.

#### Cell culture and transfection

Primary hepatocytes isolated from C/EBP $\beta$ +/+ and C/EBP $\beta$ -/- mice and cultured in a serum-free medium on a collagen type I matrix remained quiescent and highly differentiated until day 12 (Buck *et al.*, 1999). Hepatocytes were either treated with TNF- $\alpha$  for 30 min (10 ng/ml) or not treated (control). In some experiments, hepatocytes were treated for 30 min with SIN-1 (0.6  $\mu$ M) or for 45 min with leptomycin B (10 ng/ml). HepG2 cells (ATTC) were cultured in DMEM supplemented with 10% fetal bovine serum (Gibco). Treatments were started after transfection and were provided every 24 h. DNA transfection was carried out with Lipofectin as recommended by the manufacturer (BRL). The ALB-CAT plasmid (1  $\mu$ g) (Kioussis *et al.*, 1979) was transfected with CMV-C/EBP $\beta$ , CMV-C/EBP $\beta$ -Ala240, CMV-C/EBP $\beta$ -Asp240 and/or control pcDNA (1  $\mu$ g each). Cells were harvested 48 h after transfection and CAT content determined by ELISA following the manufacturer's (5 Prime $\rightarrow$ 3 Prime) recommendations (Buck and Chojkier, 1996).

#### Nuclear extracts preparation and gel retardation assay

Tissue or cells were homogenized in the presence of protease and phosphatase inhibitors as described (Descombes *et al.*, 1990; Buck *et al.*, 1994). The nuclei were sedimented in a 30% sucrose cushion by a 4000 g centrifugation at 4°C for 20 min, and lysed (Descombes *et al.*, 1990; Buck *et al.*, 1994; Buck and Chojkier, 1996). Gel retardation analyses of protein–DNA complexes were performed with equal amounts of nuclear extracts (5 µg protein) as described (Descombes *et al.*, 1990; Trautwein *et al.*, 1993; Buck *et al.*, 1994). The sense oligonucleotides were D-site (5'-TGGTATGATTTTGTAATGGGG-3'), B-site (5'-AGTATGGTT-AATGATCTA-3') and Sp1 (5'-GGGGCGGGCC-3'). Expression of recombinant C/EBP $\beta$  protein was induced with isopropyl- $\beta$ -D-thiogalactopyranoside (IPTG), from the coding region of C/EBP $\beta$  cDNA (Descombes *et al.*, 1990) cloned into a pRSET vector, and purified with Probond Resin (Invitrogen) (Buck and Chojkier, 1996).

#### Phosphorylation of C/EBP $\beta$ in vivo

Day 5 hepatocytes  $(60 \times 10^6 \text{ in } 30 \text{ ml of serum-free}, \text{phosphate-free}$  hepatocyte medium) were labeled with 2 mCi/ml (total 30 mCi) of [<sup>32</sup>P]orthophosphate (ICN, Irvine, CA) for 18 h (Buck *et al.*, 1999). Cells were either treated with TNF- $\alpha$  (10 ng/ml) for 30 min before harvest or not treated (control). Phosphopeptide mapping of endogenous C/EBP $\beta$  was performed as described (Boyle *et al.*, 1991).

# Immunohistochemistry, RNA determination and immunoblotting

Immunohistochemical detection of MDA-protein adducts and NOS2 was performed in liver sections (Houglum et al., 1990; Buck et al., 1994), using the avidin-biotin-alkaline phosphatase system (Vector Laboratories, Inc.) and antibodies against MDA-lysine epitopes (Houglum et al., 1990) and NOS2 (Transduction Laboratories). Immunofluorescence staining was observed in livers using a triple-channel fluorescence microscope. Fluorochromes utilized were fluorescein isothiocyanate (FITC), Texas red, 4',6-diamidino-2-phenylindole (DAPI) and TOTO-3 (Molecular Probes). Fluorescent labels were also analyzed by confocal microscopy in hepatocytes using purified antibodies reactive to both human and mouse C/EBPB, C/EBPa, CRM1 (Santa Cruz Biotechnologies) and C/EBPB-PSer239 (induced in rabbits with the epitope CIAVRK-PSer239-RDKAK linked to keyhole limpet hemocyanin). RNA was purified from liver with Trizol following the manufacturer's protocol (BRL). RNase protection assays were performed with specific riboprobes for albumin or 18S RNA (Ambion) as described previously (Buck and Chojkier, 1996).

#### Immunoprecipitation and immunoblotting

C/EBPβ, C/EBPβ-Ala239, C/EBPβ-Asp239, C/EBPα and CRM1 were detected by immunoblotting immunoprecipitates from hepatocyte lysates (Buck *et al.*, 1994) following the chemiluminescence protocol (DuPont) and using purified IgG antibodies as described (Trautwein *et al.*, 1993).

#### Human subjects

Preliminary studies were conducted in six male patients ( $62 \pm 5$  years; serum albumin  $2.5 \pm 0.4$  g/dl) with cachexia (30% decrease in ideal body weight over the preceding 6 months) due to cancer (anal, esophageal, lung

and kidney carcinomas, multiple myeloma and malignant mesothelioma) but without liver metastasis, and in four male control subjects ( $67 \pm 4$  years; serum albumin  $3.7 \pm 0.2$  g/dl) with normal body weight and without cancer (University of California, San Diego, CA IRB #991275). Liver biopsies were obtained during elective intra-abdominal surgery and processed as described above.

#### Statistical analysis

Results are expressed as the mean ( $\pm$  SEM) of at least triplicate experiments unless stated otherwise. Either Student's *t*- or Fisher's exact test was used to evaluate the differences of the means between groups, with a *P*-value of <0.05 as significant.

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