

Restoration of *klotho* expression induces apoptosis and autophagy in hepatocellular carcinoma cells

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Accepted: 6 December 2012 / Published online: 18 December 2012
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Abstract

Purpose *Klotho* has been identified as a tumor suppressor in several human malignancies including hepatocellular carcinoma (HCC). However, the signaling pathways involved in the tumor suppressive role of *klotho* in HCC have not been reported. Here, we investigated the role of *klotho* in HCC cell proliferation, apoptosis, autophagy, and invasion, as well as its associated signal transduction pathways.

Methods Restoration of *klotho* gene expression was established by delivering a *klotho* gene expression vector into the human HCC cell lines HepG2 and MHCC-97-H. Cell viability was measured using a cell counting (CCK-8) assay and apoptosis was analyzed through flow cytometry. Autophagy was measured via LC3-I and LC3-II protein expression levels and tumor cell invasion was assessed using a Matrigel invasion chamber assay. Expression and phosphorylation of several apoptosis and survival related proteins were assessed using Western blot assays.

Results Exogenous *klotho* gene expression significantly inhibited HCC cell proliferation, induced HCC cell apoptosis, increased LC3-I and LC3-II protein expression in HCC cells, and decreased migration of HCC cells in a Matrigel invasion chamber assay. Exogenous *klotho* gene expression also down-regulated the phosphorylation levels of the IGF-1 receptor, and the downstream Akt, ERK, and p70S6K proteins. Both apoptosis and autophagy inhibitors decreased *klotho*-induced apoptosis and autophagy.

Conclusion *Klotho* is a tumor suppressor that, through the regulation of IGF-1R phosphorylation and subsequent activation of downstream Akt-p70S6K and ERK signaling, regulates HCC tumor cell proliferation, apoptosis, autophagy and invasion.

Keywords *Klotho* · Apoptosis · Autophagy · Invasion · Hepatocellular carcinoma

1 Introduction

Primary liver cancer is the fifth most common cancer and the third most frequent cause of cancer-related death worldwide [1, 2]. Hepatocellular carcinoma (HCC) accounts for 4 % of the currently diagnosed cancers, 42 % of which occur in China [3]. The incidence of HCC is increasing in the western world, particularly in the United States, Canada and southern Europe [4]. Despite overall improvements in cancer therapy, the 5 year survival rate of HCC is still approximately 5 %. This is partially due to the fact that HCC is highly aggressive and that the key molecular mechanisms underlying HCC initiation and progression have not been uncovered yet. However, a variety of HCC related signaling pathways has been identified.

Klotho has been shown to function as a tumor suppressor, and our recent work has shown that *klotho* also acts as a tumor suppressor in HCC and that decreased *klotho* expression is involved in HCC progression [5]. *Klotho* has been observed to inhibit insulin/IGF-1 signaling, to induce cell apoptosis and, consequently, to inhibit tumor growth [6, 7]. IGF-1 exerts its roles through the activation of Raf / Mek / Erk (extracellular signal-regulated kinases) and PI3K (phosphoinositide 3-kinase) / Akt / mTOR signaling. A number of studies have revealed that insulin/IGF-1, Raf / Mek / Erk, and PI3K / Akt / mTOR signaling pathways are also involved in hepatocarcinogenesis [8–10]. Therefore, we

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hypothesized that *klotho* may inhibit IGF-1 signaling in HCC and, subsequently, induce apoptosis through down-regulation of the Raf/Mek/Erk and PI3K/Akt/mTOR signaling pathways.

Autophagy is a physiological process that eliminates toxic wastes or damaged cellular components in response to stress. As a form of type II programmed cell death, overactive autophagy may cause cell death [11, 12]. A number of studies have reported that autophagy is activated in cancer cells in response to various anticancer therapies, and that autophagy can be considered as a way to kill apoptosis-resistant tumor cells [3]. Autophagy can be activated through multiple signaling pathways. There is increasing evidence supporting the idea that tumor suppressor genes promote autophagy, while oncogenes inhibit autophagy [13]. We, therefore, hypothesized that *klotho* might also be responsible for regulating autophagy in HCC.

Here, we investigated the involvement of *klotho* in HCC cell proliferation, apoptosis, autophagy and invasion, as well as its associated signaling pathways, by exogenously expressing the *klotho* gene in two human HCC cell lines. Our study provides evidence that *klotho* regulates signaling pathways involved in HCC cell proliferation, apoptosis, autophagy and invasion.

2 Materials and methods

2.1 Cell culture

HepG2 and MHCC-97-H are human hepatocellular cancer-derived cell lines, and L-02 is a normal human liver cell line. HepG2 and MHCC-97-H were chosen for this study because they express low levels of *klotho* [5]. All cell lines were obtained from the Shanghai Cell Bank, Chinese Academy of Sciences (Shanghai, China). Cells were cultured in either RPMI 1640 or DMEM with 10 % fetal bovine serum, 100 units/mL penicillin, and 100 µg/mL streptomycin at 37 °C, 5 % CO₂.

2.2 Construction of a *klotho* gene expression vector

The *klotho* gene was amplified from a cDNA library established from normal liver cells. The open read frame (ORF) *klotho* cDNA was amplified by a forward primer containing a Bgl II restriction site: ACTCAGATCTGAGCCGGGCGACGGCGCGCAGA and a reverse primer containing a Bam HI restriction site: CGGTGGATCCCCTATTTGTAACCTTCTTGCC. The amplified *klotho* ORF was cloned into a pZsGreen1-C1 vector using the Bgl II/Bam HI sites (Clontech, Mountain View, USA). The *klotho* ORF was fused to the C-terminal end of GFP. A pZsGreen1-C1 vector without insertion was used as a negative control.

2.3 *Klotho* gene expression analysis by RT-PCR

Cultured cells were homogenized in Trizol reagent (Invitrogen, Carlsbad, CA) using a PowerGen homogenizer (Fisher Scientific, Pittsburgh, PA). Total RNA was isolated following the manufacturer's instructions. Reverse transcription was performed using a RevertAid™ H Minus First Strand cDNA Synthesis Kit (Thermo Science, Waltham, MA, USA). The *klotho* cDNA was amplified using forward primer: 5'-CACGGCAAGGGTTCGTCAT-3' and reverse primer: 5'-TCGCGCCCACGAGATGGAGA-3'. The control *GAPDH* cDNA was amplified using forward primer: 5'-CTCATGACCACAGTCCATGC-3' and reverse primer: 5'-TTCAGCTCTGGGATGACCTT-3'. RT-PCR products were visualized on a 1.5 % agarose gel containing 0.5 µg/ml of ethidium bromide.

2.4 Cell viability assay

Cell Counting Kit-8 (CCK-8) (Beyotime, Shanghai, China) allows a sensitive colorimetric assessment of cell viability. Briefly, HepG2 and MHCC-97-H cells were seeded into 96-well plates at 1×10^4 cells per well 24 h before transfection. Cells were transfected with a *klotho* expression vector, an empty vector, or no vector (PBS) using lipofectamine 2,000 according to the user manual (Invitrogen, Grand Island, NY, USA). Cells were then cultured in Opti-MEM (Invitrogen) for 6 h and subsequently cultured in normal growth medium for 72 h. Finally, 10 µl of kit reagent was added to the cells followed by incubation for 1 h. Cell viability was assessed using a microplate reader at 450 nm. All results were normalized to OD values measured from identically conditioned wells with plain medium. Cell proliferation rate (%) = (OD value in treated cells—OD value in background control wells) / (OD value of control cells—OD value in background control wells) × 100 %.

2.5 Cell proliferation and apoptosis assay

HepG2 and MHCC-97-H cells were seeded in 10 cm culture dishes at a density of 2×10^6 cells per dish. After reaching 60 % confluency, cells were transfected with a *klotho* expression vector, an empty vector, or PBS as described above. Cells were then trypsinized and suspended in 500 µl binding buffer containing 5 µl of Annexin V-FITC and 5 µl Propidium Iodide (Abcam, Cambridge, MA, USA). After incubation in the dark for 1 h, cells were subjected to flow cytometry.

2.6 Treatment with autophagy and apoptosis inhibitors

HepG2 and MHCC-97-H cells at 70 % confluency were transfected with a *klotho* expression vector, an empty vector, or PBS as described above. Cells transfected with a *klotho*

expression vector or PBS were incubated with 10 mM 3-methyladenine (3-MA) or 20 μ M apoptosis inhibitor Z-VAD-FMK for 24 h. Cells were then harvested for Western blot and/or flow cytometry analysis.

2.7 Matrigel invasion assay

The invasive ability of tumor cells was assessed using a Matrigel invasion chamber (BD Biosciences, San Jose, CA) according to the instructions provided by the manufacturer. HepG2 and MHCC-97-H cells ($5 \times 10^4/750 \mu$ l) were loaded on the transwell chamber (8 μ m pore) covered with Matrigel. 10 ng/ml of EGF was placed in the lower chamber. The number of cells that migrated to the lower surface during a 24 h-incubation period was determined by counting the cells in 5 randomly chosen fields [14]. In this assay system, the pore filter was covered with a thin layer of Matrigel. Therefore, the cells have to first digest the Matrigel before they can reach the lower chamber.

2.8 Western blot analysis

Tumor cells were homogenized and Western blotting was performed as previously described [15]. The anti-*klotho*, anti-Akt, anti-phospho-Akt1, anti-phospho-p70S6K, anti-phospho-IGF-IR, anti-GAPDH, and HRP-conjugated secondary antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). The anti-LC3-I antibody was purchased from Epitomics (Burlingame, CA, USA). The anti-LC3-II, anti-ERK1/2, anti-phospho-ERK, and anti-p70S6K antibodies were purchased from Cell Signaling Technology (Danvers, MA, USA). Protein concentrations were measured using a BCA Protein Assay kit (Beyotime, Shanghai, China). Twenty μ g of total protein was loaded onto a 12 % SDS-PAGE gel and transferred to nitrocellulose membranes. After blocking with 5 % non-fat milk for 1 h, membranes were incubated with a primary antibody overnight at 4 °C and, subsequently, incubated with a HRP-labeled secondary antibody (1:2,000 dilution) for 2 h at room temperature. Reactive proteins were detected using chemiluminescent reagents (Pierce, Rockford, IL, USA). To control for loading efficiency, the blots were stripped and re-probed with an anti-GAPDH antibody. The images were scanned using Adobe photoshop (Adobe, San Jose, CA) and quantified using NIH Image J (see <http://rsb.info.nih.gov>). All protein expression levels were evaluated relative to α -tubulin.

2.9 Statistical analysis

Data were analyzed using SPSS 13.0 (statistical package for the Social Sciences Version 13.0). Two samples were compared using the χ^2 independence test. A $p < 0.05$ was considered statistically significant.

3 Results

3.1 *Klotho* expression in normal liver and HCC cells

Klotho expression in normal human liver cells and hepatocellular cancer cells was detected using Western blot analyses. As shown in Fig. 1a & b, *klotho* expression in HepG2 and MHCC-97-H tumor cells was clearly lower than that in L-02 normal cells ($p < 0.001$). The *klotho* expression vector (see materials and methods) was effectively transfected in about 70 % of HepG2 and MHCC-97-H cells (Fig. 1c, indicated by GFP expression). Subsequent Western blot analysis showed that the transfection resulted in over-expression of *klotho* in both HepG2 (Fig. 1d, e) and MHCC-97-H cells (Fig. 1f, g).

3.2 Restoration of *klotho* expression decreases tumor cell proliferation and increases tumor cell apoptosis

The CCK-8 cell count assay showed that exogenous over-expression of the *klotho* gene significantly decreased cell viability in both HepG2 (Fig. 2a) and MHCC-97-H (Fig. 2b) cells ($p < 0.001$). The flow cytometry assay revealed that exogenous over-expression of the *klotho* gene significantly induced cell apoptosis in both HepG2 (Fig. 2c) and MHCC-97-H (Fig. 2d) cells ($p < 0.001$). Also, over-expression of the *klotho* gene significantly increased the amount of G0 cells in both HepG2 (Fig. 2e) and MHCC-97-H cells (Fig. 2f) ($p < 0.001$), implying cell cycle arrest.

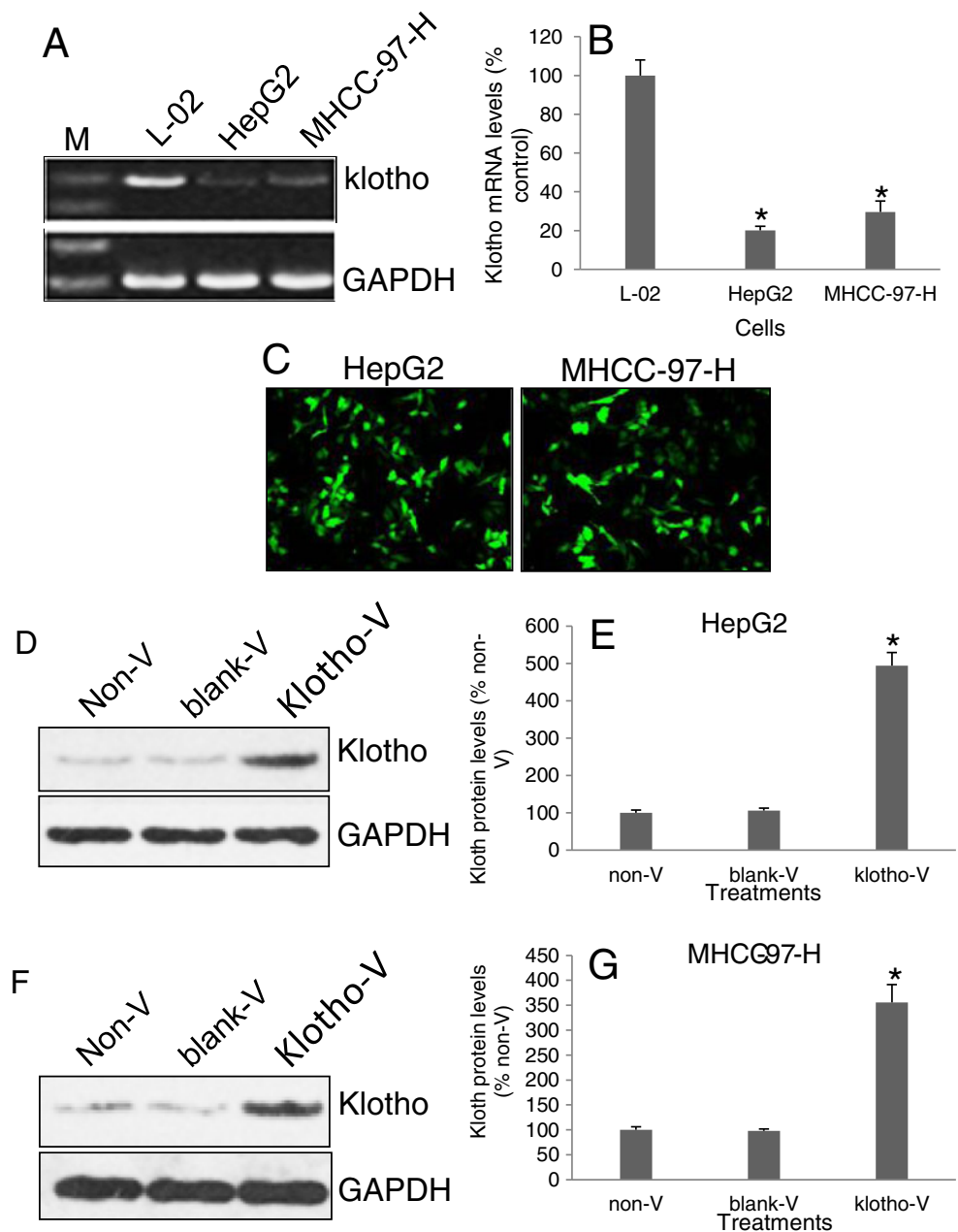
3.3 Restoration of *klotho* gene expression induces autophagy and apoptosis

LC3-I and LC3-II are markers for autophagy. Transfection of a *klotho* expression vector significantly increased LC3-I and LC3-II expression in both HepG2 (Fig. 3a, b, c, $p < 0.001$) and MHCC-97-H cells (Fig. 3d, e, f, $p < 0.001$). In HepG2 and MHCC-97-H cells transfected with a *klotho* expression vector, the autophagy inhibitor 3-MA significantly normalized LC3-I and LC3-II expression levels ($p < 0.01$). We further tested the effect of an apoptosis inhibitor and found that Z-VAD-FMK significantly decreased *klotho*-induced LC3-I and LC3-II expression in both HepG2 ($p < 0.001$) and MHCC-97-H cells ($p < 0.001$). Next we investigated the role of autophagy and apoptosis inhibitors on apoptosis in HepG2 (Fig. 3g) and MHCC-97-H (Fig. 3h) cells. The flow cytometry assay showed that both autophagy and apoptosis inhibitors significantly inhibited *klotho*-induced cell apoptosis ($p < 0.001$).

3.4 Restoration of *klotho* expression increases tumor cell invasion

The invasive capacity of tumor cells was assessed using a Matrigel invasion chamber. The number of cells that migrated

Fig. 1 *Klotho* expression in normal liver and tumor cells. **a** RT-PCR detection of *klotho* expression in L-02 normal liver cells, and HepG2 and MHCC-97-H HCC cells. **b** Quantification of *klotho* mRNA levels in (a). * $p < 0.001$ vs. L-02 cells. $N = 4$. **c** *Klotho* expression in HepG2 and MHCC-97-H cells transfected with a *klotho* expression vector. **d** Western blot analysis of *klotho* protein expression in HepG2 cells transfected with a *klotho* expression vector (*klotho*-V), an empty vector (*blank*-V), and PBS (*non*-V). **e** Quantification of *klotho* protein levels in (d). * $p < 0.001$ vs. *non*-V control. $N = 4$. **f** Western blot analysis of *klotho* protein expression in MHCC-97-H cells. **e** Quantification of *klotho* protein levels in (f). * $p < 0.001$ vs. *non*-V control. $N = 4$



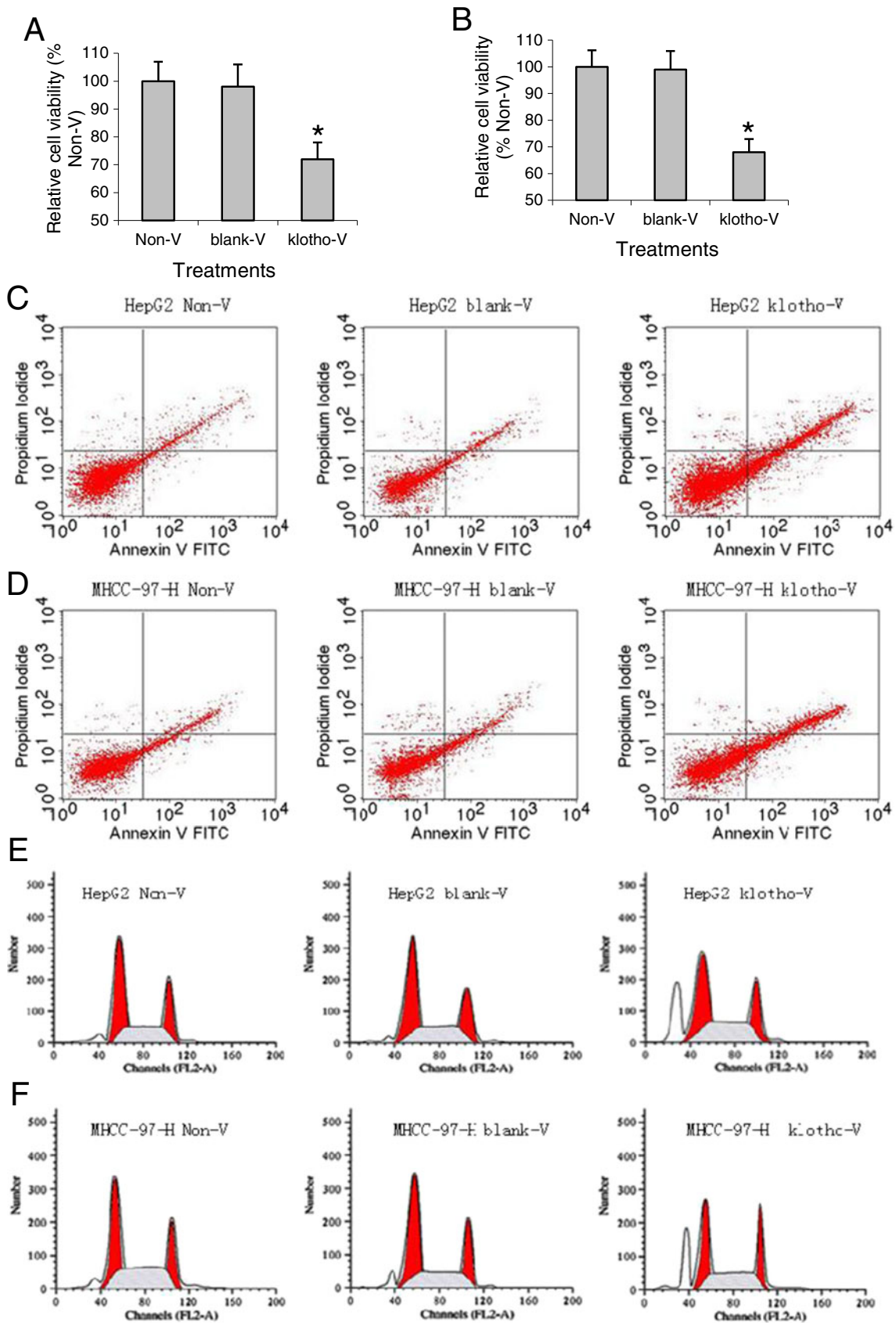
to the lower chamber was calculated. Transfection of a *klotho* expression vector significantly decreased the migration of HepG2 and MHCC-97-H cells to the lower chamber ($p < 0.001$), suggesting that the *klotho* gene can inhibit hepatocellular cancer cell invasion (Fig. 4).

3.5 Cellular signaling involved in the inhibitory role of *klotho*

Next we investigated the expression of signaling molecules involved in cell proliferation and survival. In both HepG2 and MHCC-97-H cells, exogenous over-expression of the

klotho gene significantly inhibited phosphorylation of several proliferation- and survival-related molecules. The levels of phospho-IGF-1R, phospho-Akt, phospho-ERK, and phospho-p70S6K were significantly lower in HepG2 (Fig. 5a, c) and MHCC-97-H (Fig. 5b, d) cells transfected

Fig. 2 *Klotho* inhibits cell proliferation and induces apoptosis. **a** Relative cell viability in HepG2 cells transfected with a *klotho* expression vector (*klotho*-V), an empty vector (*blank*-V), and PBS (*non*-V). * $p < 0.001$ vs. *non*-V. $N = 5$. **b** Relative cell viability of MHCC-97-H cells. * $p < 0.001$ vs. *non*-V. $N = 5$. **c** Flow cytometry of apoptotic HepG2 cells. **d** Flow cytometry of apoptotic MHCC-97-H cells. **e** Cell cycle assessment of HepG2 cells. **f** Cell cycle assessment of MHCC-97-H cells



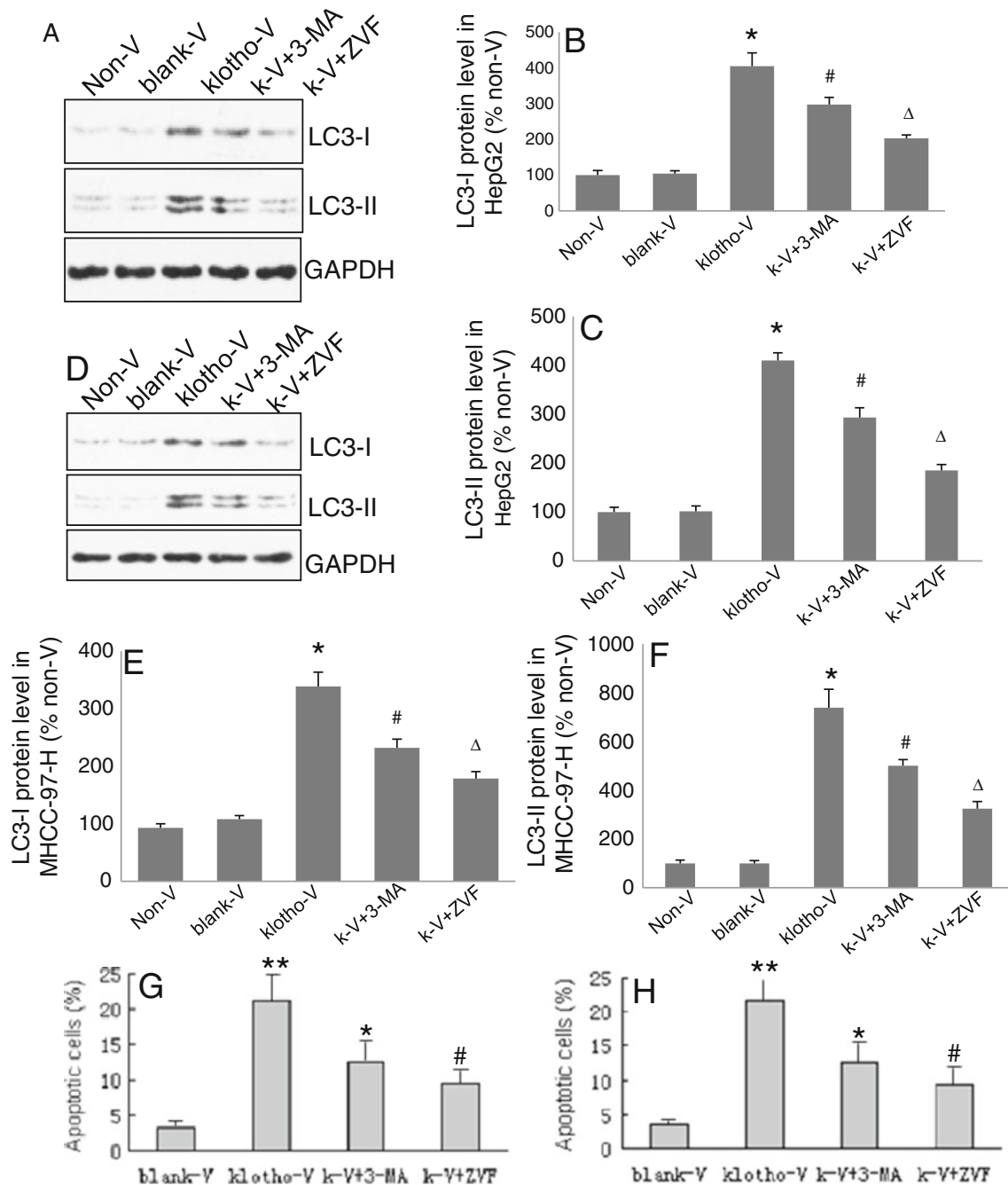


Fig. 3 The effects of apoptosis and autophagy inhibitors. **a** Western blot analysis of LC3-I and LC3-II expression in HepG2 cells. Cells were transfected with a *klotho* expression vector (klotho-V), an empty vector (blank-V), and PBS (non-V). LC3-I and LC3-II expression were also measured in HepG2 cells transfected with a *klotho* expression vector (K-V) and incubated with an autophagy inhibitor (k-V + 3-MA) or an apoptosis inhibitor (k-V + ZVF). ZVF: Z-VAD-FMK. **b** Quantification of LC3-I levels in HepG2 cells in (a). * $p < 0.001$ vs. non-V control. # $p < 0.01$ vs. klotho-V, $\Delta p < 0.001$ vs. klotho-V. $N = 4$. **c** Quantification of LC3-II levels in HepG2 cells in (a). * $p < 0.001$ vs. non-V

control. # $p < 0.01$ vs. klotho-V, $\Delta p < 0.001$ vs. klotho-V. $N = 4$. **d** Western blot analysis of LC3-I and LC3-II expression in MHCC-97-H cells. **e** Quantification of LC3-I levels in MHCC-97-H cells in (d). * $p < 0.001$ vs. non-V control. # $p < 0.01$ vs. klotho-V, $\Delta p < 0.001$ vs. klotho-V. $N = 4$. **f** Quantification of LC3-II levels in MHCC-97-H cells in (d). * $p < 0.001$ vs. non-V control. # $p < 0.01$ vs. klotho-V, $\Delta p < 0.001$ vs. klotho-V. $N = 4$. **g** Percentage of apoptotic cells in HepG2 cell with different treatments. ** $p < 0.001$ vs. blank-V; * $p < 0.01$ vs. klotho-V; # $p < 0.01$ vs. klotho-V. $N = 5$. **h** Percentage of apoptotic cells in MHCC-97-H cells. ** $p < 0.001$ vs. blank-V; * $p < 0.05$ vs. klotho-V; # $p < 0.01$ vs. klotho-V. $N = 5$

with a *klotho* expression vector than in cells transfected with an empty vector or PBS ($p < 0.001$). No differences were

observed in protein levels of total ERK, Akt, and p70S6K between the 3 treatment groups.

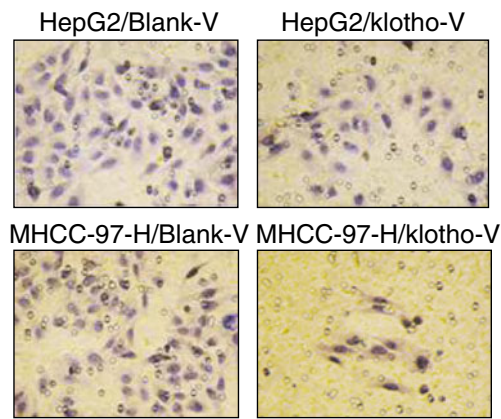


Fig. 4 *Klotho* increases tumor cell migration. Tumor cell migration was assessed using a Matrigel invasion chamber. The cells that migrated to the lower chamber are shown. Transfection of a *klotho* expression vector significantly decreased the migration of HepG2 and MHCC-97-H cells to the lower chamber ($p < 0.001$)

4 Discussion

Klotho has been shown to act as a tumor suppressor in several malignancies, including HCC. Our recent work has shown that absence of *klotho* gene expression is associated with progression of HCC. However, the signaling pathways associated with *klotho*'s mode of action in HCC survival and growth have thus far remained unknown. Here, the roles played by *klotho* in cell proliferation, apoptosis and autophagy were investigated through the delivery of a *klotho* gene expression vector into two HCC cell lines (HepG2 and MHCC-97-H) and one normal liver cell line (L-02). Our results show that restoration of *klotho* gene expression in HCC cells inhibits tumor cell proliferation, induces apoptosis and autophagy, and reduces tumor cell invasion. In addition, we found that *klotho* exerts a tumor suppressive role through down-regulation of IGF-1R, Akt, ERK and p70S6K phosphorylation.

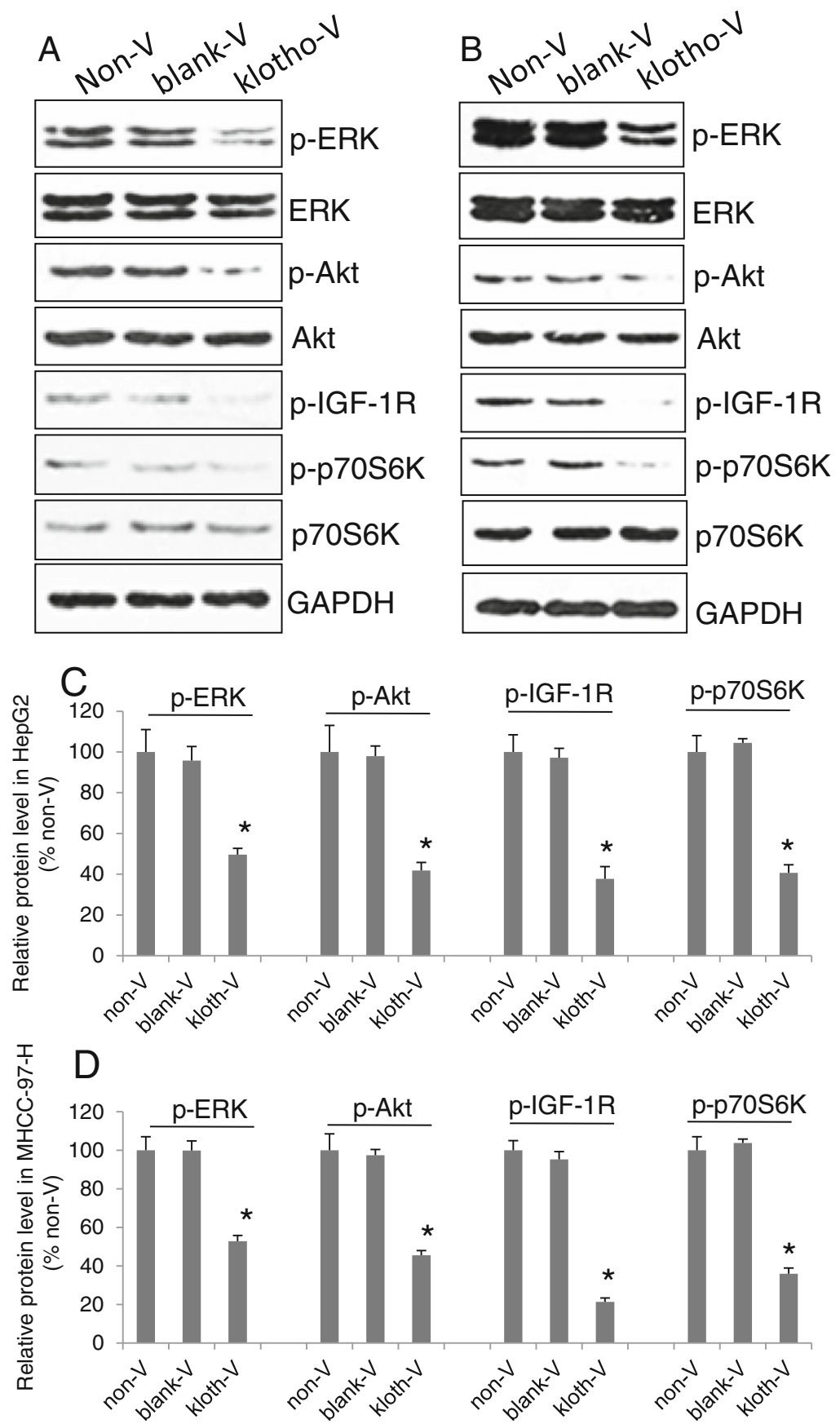
These results are similar to those previously reported in lung and colorectal cancers [13, 16]. The signaling pathways involved in the tumor suppressive role of *klotho* have, as yet, not been fully elucidated. *Klotho* has been found to inhibit insulin/IGF-1 signaling and, consequently, to induce cell apoptosis and to inhibit tumor growth [17–19]. *Klotho* regulates insulin/IGF-1 signaling by inhibiting tyrosine phosphorylation of the IGF-1 receptor [20]. Tyrosine phosphorylation of the IGF-1R is also involved in the activation of extracellular signal-regulated kinases (ERK) 1 and ERK2 and PI3K/Akt signaling [21]. In this study, we revealed that restoration of *klotho* gene expression significantly inhibits IGF-1R, Akt, and ERK phosphorylation in HCC cells. The PI3K-Akt pathway plays a pivotal role in the regulation of apoptosis in many cell types, and the ras-raf-MEK-ERK route has also been proposed to function in cell survival.

We, therefore, propose that through inhibiting IGF-1 receptor phosphorylation, *klotho* inhibits ERK and Akt activities and induces apoptosis in HCC cells. p70S6K is a downstream effector of the PI3K/Akt/mTOR signal transduction pathway. It has been found that inhibition of p70S6K phosphorylation induces cell apoptosis and cell cycle arrest in G1 and G2 in B-cell lymphoma cells [22].

Non-apoptotic cell death is mainly attributed to autophagy, which is considered a viable alternative to kill apoptosis-resistant tumor cells [3]. Autophagy is initiated by the formation of an autophagosome or autophagic vacuole, which then fuses with the lysosomal membrane to deliver its content into the autolysosome for degradation [23]. During autophagosome formation, the microtubule-associated protein light chain 3 I (LC3-I) is conjugated to phosphatidylamine to form LC3-phosphatidylamine, termed LC3-II. Translocation of LC3-II to the autophagosome membrane is an essential step in the formation of the autophagosome [23, 24]. Therefore, LC3-I and LC3-II levels can be used as markers to reflect the activation of autophagy. Here, we detected low LC3-I and LC3-II expression levels in HCC cells and that, after restoration of *klotho* expression, LC3-I and LC3-II levels significantly increased. Furthermore, the autophagy inhibitor 3-MA inhibited LC3-I and LC3-II expression in cells transfected with a *klotho* expression vector, suggesting that *klotho* is an inducer of autophagy in HCC. Thus, *klotho* is a tumor suppressor capable of inhibiting both apoptosis and autophagy in HCC. However, how *klotho* inhibits apoptosis and autophagy in tumor cells has so far not been elucidated.

Recent studies have demonstrated that autophagy can be induced by reduced growth factor signaling. Growth factor signaling involved in the insulin/IGF-1-PI3K-Akt-mTORC1 pathway has been shown to regulate cell autophagy through the insulin receptor [25]. In addition, activation of insulin/IGF signaling has been shown to suppress the autophagic-lysosomal pathway [26, 27]. Importantly, the *klotho* protein functions as a circulating hormone that represses intracellular signals of insulin and IGF-I [20, 28]. In this study, exogenous over-expression of the *klotho* gene significantly inhibited IGF-1R, Akt, and p70S6K phosphorylation. This suggests that the signaling pathway of *klotho*-IGF-1R/Akt may be responsible for both apoptosis and autophagy. Unexpectedly, autophagy inhibitors significantly reversed *klotho*-induced apoptosis, while apoptosis inhibitors significantly inhibited autophagy in HCC cells. This surprising observation suggests that apoptosis and autophagy act through similar pathways. However, the inhibition of apoptosis by an inhibitor to autophagy, and autophagy by an inhibitor to apoptosis, was not 100 % effective, suggesting that apoptosis and autophagy caused by over-expression of *klotho* may, at least partially, employ different cell death pathways.

Fig. 5 Klotho inhibits phosphorylation of IGF-1, Akt, p70S6K and ERK proteins. **a** Western blot analysis of protein and phosphorylation levels in HepG2 cells. **b** Western blot analysis of protein and phosphorylation levels in MHCC-97-H cells. **c** Quantification of protein levels in HepG2 cells in **a**). * $p < 0.001$ vs. non-V control. $N = 4$. **d** Quantification of protein levels in MHCC-97-H cells in **b**). * $p < 0.001$ vs. non-V control. $N = 4$. *P-IGF-1R* phospho-IGF-1 receptor; *p-Akt* phospho-Akt; *p-ERK* phospho-ERK; *p-p70S6K* phospho-p70S6K; *klotho-V* klotho expression vector; *blank-V* empty vector; *non-V*: PBS transfection



In this study, *klotho* was identified as a tumor suppressor that inhibits tumor cell proliferation, induces cell apoptosis and autophagy as well as inhibits invasion of HCC cells. These tumor suppressive roles of *klotho* may be associated with a down-regulation of IGF-1 receptor phosphorylation, and a subsequent decreases in Akt/p70S6K and ERK phosphorylation. Our study highlights a central role for *klotho* in HCC cell survival and suggests that *klotho* may serve as an ideal target for the development of novel therapeutic agents for HCC.

Conflict of interests All author declared no any conflict of interest.

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