Positive feedback regulation between Akt phosphorylation and fatty acid synthase expression in osteosarcoma

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Abstract. The activation of PI3K/Akt and the overexpression of fatty acid synthase (FASN) are frequently observed in human osteosarcoma (OS). In the present study, in order to investigate the possible association between the phosphorylation of Akt and FASN expression, immunohistochemical staining was conducted on 24 OS specimens from patients with pulmonary metastasis, which revealed a significant positive correlation between phosphorylated Akt (p-Akt) and the expression of FASN (R=0.469, P=0.04). To investigate the association between p-Akt and FASN in vitro, human U2-OS OS cells were treated with FASN-specific RNAi plasmid or LY294002 (an inhibitor of PI3K/Akt). The mRNA levels of Akt and FASN were measured by real-time PCR. Western blot analysis was also performed to detect the protein expression of PI3K, Akt, p-Akt and FASN. The results demonstrated that FASN may contribute to the metastasis of OS cells. In addition, the effects induced by the inhibition of the activity of p-Akt or FASN on the malignant phenotype of U2-OS cells were investigated, demonstrating that the malignant phenotype was inhibited by suppressing the activity of PI3K/Akt or FASN in the U2-OS cells. The findings from our study suggest the existence of a positive feedback regulation between Akt phosphorylation and FASN expression and that this loop may play an important role in the malignant phenotype of OS cells.

Introduction

Osteosarcoma (OS) is one of the most common primary malignant bone tumors in children and adolescents. Following the advent of effective chemotherapy, the five-year survival rate for patients with OS treated with intensive multidrug chemotherapy and aggressive local control has been reported to be 55-80% (1-3). However, numerous studies have reported that the five-year survival rate of patients with metastatic diseases is <20% (4-6). The development of lung metastasis is the main cause of mortality in patients with OS. The identification and understanding of the molecular mechanisms responsible for metastasis would pose a significant impact on the management of OS.

Fatty acid synthase (FASN) is an enzyme which is considered crucial for endogenous lipogenesis in mammals, and is responsible for catalyzing the synthesis of long-chain fatty acids. In the majority of normal cells, FASN expression is usually not observed, due to the presence of abundant amounts of dietary lipids (7). However, FASN is overexpressed in a variety of human tumors (8-12), and has been strongly linked to cancer cell proliferation and apoptosis (13-16). We previously demonstrated that FASN may contribute to the metastasis of OS cells (17,18). However, its potential molecular mechanisms of action remain unclear.

PI3K/Akt plays a crucial role in the cell-extracellular matrix (ECM) and cell-cell adhesion. Due to lack of correct adhesion, the adhesion-dependent signals are interrupted, resulting in adhesion-related apoptosis, namely anoikis. PI3K/Akt signaling has been implicated in the regulation of FASN expression in breast cancer and prostate cancer cells (19,20). Wang et al reported that there was a positive feedback regulation between Akt phosphorylation and FASN expression in ovarian carcinoma cells (21). However, to our knowledge, the association between the phosphorylation of Akt and FASN protein expression in OS has not yet been documented.

In this study, we found that the inhibition of Akt phosphorylation by LY294002 (an inhibitor of PI3K/Akt), resulted in the downregulation of FASN expression. In addition, the downregulation of FASN expression inhibited Akt phosphorylation. Based on these findings, we confirmed the existence of a positive feedback loop between Akt phosphorylation and...
FASN expression; this feedback loop may play an important role in the malignant phenotype of OS cells.

Materials and methods

Patient specimens. A total of 24 samples of OS tissues were obtained from patients with pulmonary metastatic disease who underwent surgery in our hospital (The First Hospital Affiliated to Nanchang University, Nanchang, China) from 2005 to 2012. The pulmonary metastasis survey was performed with plain films and chest CT scans at first diagnosis. All the patients had no history of prior treatment with anticancer drugs or radiotherapy. The samples were fixed with 10% formalin and embedded in paraffin and were then cut into 4-μm-thick sections. In all cases, informed consent was obtained from the relative departments and persons, and the study had the approval of the Ethics Committee of Nanchang University.

Immunohistochemistry. Immunohistochemical (S-P) staining with and hematoxylin and eosin (H&E) was performed on the paraffin-embedded tissue sections. Antigen retrieval was performed by heating the sections in 10 mmol/l citrate buffer (pH 6.0) for 20 min. FASN and phosphorylated Akt (p-Akt) antibodies (rabbit monoclonal antibody; antibody dilutions, 1:50; Epitomics, Inc., Burlingame, CA, USA) were used as the primary antibody at a final dilution as corresponding product specifications. The sections were then stained with diaminobenzidine (DAB) and counterstained using hematoxylin. The stained sections were evaluated and scored by two pathologists in a blinded manner without prior knowledge of the clinical pathological characteristics of the patients. According to the staining intensity by examining at least 500 cells in five representative areas, the expression level of p-Akt and FASN was measured and the intensity scores were recorded as follows: none, 0; weak, 1; moderate, 2; and intense, 3. According to the percentage of cancer cells with a positive expression of Akt and FASN, the percentage scores were recorded as follows: 0% (score 0); <10% (score 1); 10-49% (score 2); 50-79% (score 3); and 80-100% (score 4). The final score was averaged with the scores from the two pathologists; these scores were calculated by multiplying the intensity score by the percentage score. The sections with a final score of <4 were considered as negative (-), those with a score of 4-5 were considered as positive (+), those with a score of 6-8 as double positive (++), and those with a score of 9-12 were considered as triple positive (++++).

Cell culture and transfection. The human OS cell line, U2-OS, was purchased from the American Type Culture Collection (ATCC: Manassas, VA, USA), and the cells were routinely cultured in RPMI-1640 medium (HyClone, Logan, UT, USA) supplemented with 10% fetal bovine serum (FBS) (Sigma, Lenexa, KS, USA) in a humidified 37°C incubator containing 5% CO2. The U2-OS cells were seeded in 6-well plates till 40% confluence on the day prior to transfection. The U2-OS cells were transfected with FASN-specific RNAi plasmid (MR-FASN) and the negative control RNAi plasmid (MR-Neg) using Lipofectamine 2000 according to the instructions of the manufacturer (Invitrogen Life Technologies (Carlsbad, CA, USA).

Table I. Primer sequences of genes used in real-time PCR.

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<tr>
<th>Gene (size, bp)</th>
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<tr>
<td>FASN (171)</td>
<td>F AACTCCATGTTGTTGTTTG</td>
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<td></td>
<td>R CACATCGGTTTAATTGTTG</td>
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FASN, fatty acid synthase; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; F, forward, R, reverse.

Real-time PCR. Semi-quantitive (real-time) PCR was used to detect the FASN mRNA expression levels. Total RNA was extracted from the cells using TRIzol reagent (Invitrogen Life Technologies). The total RNA concentration was determined by spectrophotometry at 260 nm and the purity was determined by calculating the 260/280 ratio with a BioPhotometer (Eppendorf, Hamburg, Germany). The two-step kit (Promega Corp., Madison, WI, USA) was used to to obtain cDNA according to the manufacturer's instructions, which was then used as the template for amplification. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as an internal standard. The gene primer sequences are listed in Table I. The cycling conditions were as follows: initial denaturation at 94°C for 3 min, followed by 35 amplification cycles of 94°C for 30 sec, 55°C for 30 sec and 72°C for 12 sec. Each real-time PCR assay contained 2 µl cDNA template, 10 µl SuperMix and 0.5 µl of every forward and reverse primer in a 20 µl reaction mixture. All experiments were repeated six times over multiple days.

Western blot analysis of protein expression. Total protein from the cells was extracted using RIPA lysis buffer containing 60 µg/ml phenylmethylsulfonyl fluoride (PMSF). The protein concentrations were determined using the BCA protein assay kit (Pierce, Shanghai, China). The total RNA concentration was determined at 260 nm and the purity was determined by calculating the 260/280 ratio with a BioPhotometer (Eppendorf, Hamburg, Germany). The two-step kit (Promega Corp., Madison, WI, USA) was used to to obtain cDNA according to the manufacturer's instructions, which was then used as the template for amplification. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as an internal standard. The gene primer sequences are listed in Table I. The cycling conditions were as follows: initial denaturation at 94°C for 3 min, followed by 35 amplification cycles of 94°C for 30 sec, 55°C for 30 sec and 72°C for 12 sec. Each real-time PCR assay contained 2 µl cDNA template, 10 µl SuperMix and 0.5 µl of every forward and reverse primer in a 20 µl reaction mixture. All experiments were repeated six times over multiple days.

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FASN, fatty acid synthase; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; F, forward, R, reverse.
Cell proliferation assay. Cells (4x10^3/200 µl/well) were seeded in 96-well plates. Viable proliferating cells were detected by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazoliumbromide (MTT) assay at various time periods (24, 48 and 72 h), using five wells per time period. Cell viability was expressed as the optical density (OD), which was detected by an enzyme-linked immunoabsorbent assay reader (MK3; Thermo Scientific) at a 490-nm wavelength. All experiments were repeated six times over multiple days.

Analysis of cell apoptosis. Cells (5x10^5) were harvested, washed with PBS and resuspended in binding buffer, followed by mixing with Annexin V-FITC and propidium iodide (both from KeyGen Biotech. Co., Ltd., Nanjing, China). The cells were analyzed by a BD FACSCalibur flow cytometer (BD Biosciences, San Jose, CA, USA). All experiments were repeated six times over multiple days.

Transwell invasion assays. Cell invasion was measured in 24-well plates by Transwell assay using a chamber containing the polyethylene terephthalate filter membrane with 8-µm pores (BD Biosciences). The cells (6x10^4/200 µl/chamber) were seeded in the upper chamber with RPMI-1640 medium containing 10 g/l BSA, and the lower well was filled with 500 µl RPMI-1640 medium supplemented with 10% FBS as a chemoattractant. Following incubation for 24 h, the chambers were stained with crystal violet. The invaded cells were counted from ten randomly selected fields under an inverted microscope. All experiments were repeated six times over multiple days.

Wound healing assays. Cell migration was assessed by determining the ability of the cells to move into a cellular space in a two-dimensional in vitro ‘wound healing assay’. In brief, the cells were grown to confluence in 6-well tissue culture plastic dishes to a density of 5x10^5 cells/well. The cells were demuced by dragging a rubber policeman (Fisher Scientific, Hampton, NH, USA) through the center of the plate. The cultures were rinsed with PBS and replaced with fresh DMEM alone or containing 10% FBS, following which the cells were incubated at 37°C for 24 h. Images were captured at 0 and 24 h and the migrated distance was measured using ImageJ software (NIH, Bethesda, MD, USA). All experiments were repeated six times over multiple days.

Statistical analysis. Statistical comparisons were performed using SPSS software version 13.0 (SPSS Inc., Chicago, IL, USA). The correlation of FASN with p-Akt protein in the OS tissues was evaluated using the Wilcoxon rank sum test. All measurement data are presented as the means ± SD, and the one-way ANOVA with a post-hoc test (Student-Newman-Keuls test) was performed for statistical analysis. A P-value <0.05 was considered to indicate a statistically significant difference.

Results

Correlation between FASN and p-Akt protein expression in OS. To explore the possible association between FASN expression and the phosphorylation of Akt in OS, FASN and p-Akt (Ser473) protein expression in the OS tissues from patients with pulmonary metastatic disease was detected. The results revealed that FASN protein was expressed in the cytoplasm of the OS tissues, and the p-Akt protein was expressed in the nucleus and cytoplasm (Fig. 1A-C). There was a significant positive relationship between FASN and p-Akt expression (R=0.469, P=0.04). These data suggest that a possible connection between FASN expression and the phosphorylation of Akt exists in OS.

Inhibiting the phosphorylation of Akt suppresses the malignant phenotype of U2-OS cells. To determine the effects of the inhibition of Akt phosphorylation on the malignant phenotype of OS cells, the U2-OS cells were treated with LY294002 (an inhibitor of PI3k/Akt) at various concentrations (0, 5, 10, 20, 40, 80 and 160 µM) for 24, 48 and 72 h. Cell proliferation and apoptosis were assessed by MTT and FACS assays, respectively. The migration and invasion ability of the cells was investigated using wound healing and Transwell assays, respectively. In the MTT assays, the results revealed that LY294002 induced U2-OS cell apoptosis and inhibited cell growth in a dose- and time-dependent manner (Fig. 2A) and the IC50 was 47.96 µM for 24 h. The concentration of 40 µM was selected for further experiments. Furthermore, in FACS assays, the results revealed that LY294002 induced U2-OS...
Figure 2. Inhibition of Akt phosphorylation induces apoptosis and inhibits the growth of U2-OS cell. (A) The U2-OS cells were treated with LY294002 (an inhibitor of PI3K/Akt) at different concentrations for 24, 48 and 72 h. LY294002 inhibited U2-OS cell proliferation in a dose- and time-dependent manner. (B and C) FACS analysis of U2-OS cells treated with LY294002 at different concentrations for 24 h. LY294002 induced U2-OS cell apoptosis in a dose-dependent manner.

Inhibition FASN suppresses U2-OS cell migration and invasion. In a previous study, we demonstrated that the suppression of FASN expression induces U2-OS cell apoptosis and inhibits cell growth in vivo and in vitro (17). Therefore, in this study, we evaluated the effects of the downregulation FASN on U2-OS cell migration and invasion. The migration and invasion ability of the cells was significantly lower in the cells transfected with the FASN-specific RNAi plasmid than those transfected with the negative RNAi plasmid (P<0.05) (Figs. 3 and 4).

Downregulation of FASN expression inhibits the phosphorylation of Akt. To investigate the effects of silencing FASN on the phosphorylation of Akt in OS, the U2-OS OS cells were transfected with the FASN-specific RNAi plasmid to inhibit
The mRNA expression of FASN and Akt was detected by real-time PCR, and western blot analysis was used to measure the protein expression of FASN, PI3K, Akt and p-Akt. The downregulation of FASN inhibited the activation of the PI3K/Akt signaling pathway. However, the mRNA expression of Akt was not affected by the downregulation of FASN in the U2-OS cells (Fig. 5A and B).

**Figure 4.** (A) Transwell invasion assay to evaluate the effects of the downregulation of fatty acid synthase (FASN) or the inhibition of Akt phosphorylation on U2-OS cell invasion. (B) Bars represent the means ± SD (n=6); * P<0.05 vs. control group (MR-Neg).

**Figure 5.** (A) Western blot analysis for the detection of the expression of phosphorylated Akt (p-Akt), Akt and fatty acid synthase (FASN) protein. The downregulation of FASN inhibited Akt phosphorylation, and the inhibition of Akt phosphorylation downregulated FASN protein expression. (B) Real-time PCR was performed to measure the mRNA expression of Akt and FASN. The inhibition of Akt phosphorylation downregulated FASN mRNA expression; however, the downregulation of FASN had no significant inhibitory effect on Akt phosphorylation.

**Inhibition of Akt phosphorylation downregulates FASN in U2-OS cells.** In order to determine the effects of the inhibition of the phosphorylation of Akt on the mRNA and protein expression of FASN in OS, the U2-OS cells were treated with LY294002 (an inhibitor of PI3K/Akt). The mRNA and protein expression of FASN was measured by RT-PR and western blot analysis, respectively. The results revealed that both the
nRNA and protein expression of FASN was decreased by the inhibition of Akt phosphorylation (Fig. 5).

Discussion

Previous studies have demonstrated that cancers with a high expression of FASN always undergo a significant endogenous fatty acid biosynthesis and display a biologically aggressive subset (22,23). Moreover, the overexpression of FASN is an early event in tumor development and is more pronounced in tumors with a poor prognosis (24). Importantly, we previously demonstrated that the inhibition of FASN with pharmacological inhibitors is selectively cytotoxic to human OS cells and leads to a significant antitumor effect (17). Although very little is known about the mechanisms underlying the upregulation of the FAS protein in cancer cells, studies have revealed that FASN is also upregulated at the mRNA level (25,26), and increasing evidence indicates that Akt activity modulates FASN expression in tumor cells (20,27). Previous studies have revealed that activated PI3K stimulates the binding of sterol regulatory element-binding protein (SREBP)-1c, a SREBP family transcription factor, which controls genes involved in lipogenesis to a SREBP-binding site in the FASN promoter, thus inducing FASN transcription (28-30). Of note, the inhibition of FASN activity by either cerulenin or C75 has been shown to inhibit the production of p-Akt (19,31). These findings suggest the existence of a positive bidirectional association between p-Akt and FASN expression in cancer cells.

In the current study, we found that there was a positive correlation between p-Akt and FASN protein expression in OS tissues. This indicates that a possible connection between the phosphorylation of Akt and FASN expression may exist. In order to investigate whether the inhibition of the phosphorylation of Akt suppresses FASN expression in OS, LY294002, an inhibitor of PI3K/Akt, was used to downregulate the phosphorylation of Akt in U2-OS cells. The results revealed that the mRNA and protein expression of FASN was markedly inhibited by LY294002, which suggests that p-Akt regulates FASN by affecting the transcription and translation in U2-OS cells.

To determine the effects of the inhibition of FASN on the phosphorylation of Akt in OS cells, the U2-OS cells were transfected with the FASN-specific RNAi plasmid to inhibit FASN expression. We found that the protein expression of both total Akt and p-Akt was decreased in the FASN-silenced U2-OS cells. However, the mRNA expression of Akt did not differ between the cells treated with the FASN-specific RNAi plasmid and those treated with the negative RNAi plasmid. These data suggest that FASN regulates PI3K/Akt at the translational rather than the transcriptional level. However, the mechanism(s) responsible for the inhibition of Akt activity by the downregulation of FASN remain unclear. Currently, several mechanisms are likely to contribute to FASN-mediated p-Akt regulation: i) FASN-mediated lipogenesis produces phospholipids that are incorporated into cell membranes and partition into lipid rafts, which accommodate ErbBs and form signaling platforms. FASN blockade destabilizes these lipid rafts, which triggers the degradation of ErbBs and impedes the membrane recruitment of downstream mediators of Akt, thereby causing the downregulation of p-Akt (32,33); ii) the FASN promoter contains several sterol regulatory element-binding protein-1 (SREBP-1) sites enabling FASN to activate Akt (34,35).

Recently, a number of studies have demonstrated that FASN and PI3K/Akt play an important role in the proliferation, invasion and migration of cancer cells (36-39). In this study, we observed that the suppression of FASN expression by RNAi or the inhibition of the activity of PI3K/Akt blocked cell proliferation and migration and increased the apoptotic rate in the U2-OS cells. This suggests that FASN and PI3K/Akt play a key role in the maintenance of the malignant phenotype of OS cells.

In conclusion, the data presented in this study, confirm the existence of a positive feedback loop between Akt phosphorylation and FASN expression. Moreover, this feedback loop plays an important role in the malignant phenotype of OS cells. However, the detailed mechanisms of the bidirectional association between FASN expression and the phosphorylation of Akt in OS cells are currently unknown. Thus, further studies are required to provide further clarification.

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References


