ORIGINAL RESEARCH

Overexpression of IGFBP5 Enhances Radiosensitivity Through PI3K-AKT Pathway in Prostate Cancer

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Background: Radiotherapy is the main treatment for localized prostate cancer. The therapeutic effects of radiotherapy are highly dependent on radiosensitivity of target tumors. Here, we investigated the impact of insulin-like growth factor-binding protein 5 (IGFBP5) on irradiation therapy in prostate cancer.

Methods: *IGFBP5* gene was overexpressed in human prostate cancer cell lines, PC3 and DU145, with transfection of lentivirus expression vector. Radiosensitivity of the cell lines was assessed with colony formation, cell cycle and cell proliferation assays. The expression of proteins associated with the PI3K-AKT pathway was determined by Western blotting. The effect of IGFBP5 knockdown on PI3K-AKT pathway was tested using PI3K inhibitor.

Results: Higher expression of IGFBP5 improved the efficacy of radiotherapy for prostate cancer patients. The effects of IGFBP5 were linked to the PI3K-AKT signaling pathway. Overexpression of IGFBP5 enhanced radiosensitivity and induced G2/M phase arrest in prostate cancer cells. In contrast, it decreased PI3K, p-AKT expression and cell viability. These effects were reversed by IGFBP5 knockdown.

Conclusion: Our results reveal that IGFBP5 regulates radiosensitivity in prostate cancer via the PI3K-AKT pathway. It is, therefore, a potential biomarker of tumors that influences the therapeutic effect of radiotherapy.

Keywords: IGFBP5, irradiation, prostate cancer, radiosensitivity, PI3K-AKT pathway

Background

Prostate cancer is one of the most common malignancies and the leading cause of deaths in men worldwide.^{1,2} The incidence of prostate cancer has been on the rise and it is now the sixth leading cause of deaths in China.³ Radiotherapy is the main treatment for localized prostate cancer. External beam radiotherapies such as photon, proton and carbon-ion beam radiation, and brachytherapy have been successful in prostate cancer treatment.^{4–7} Radiotherapy exerts good local control on the growth of cancer cells and improves of prognosis of cancer patients who show high radiosensitivity. However, radioresistance increases the risk of local recurrence, distant metastasis, and poor prognosis.⁸ Thus, it is critical to understand the mechanism of radioresistance and radiosensitivity as this will reveal avenues for enhancing radiosensitivity.

Insulin-like growth factor-binding protein 5 (IGFBP5) regulates proliferation, invasion and metastasis of many cancers.^{9–11} IGFBP5 belongs to a protein family with six members and is the most conserved among them.¹² It has a high affinity for insulin-like growth factors (IGFs) which inhibits or promotes cancer development depending on the cellular context.¹³ For instance, it suppresses tumor growth or promotes cell proliferation in breast cancer, ovarian cancer, melanoma and osteosarcoma at different contexts.^{13,14}

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The PI3K is activated by different growth factors. Subsequent to its activation, it phosphorylates and activates AKT, which plays a central role in insulin receptor intracellular signaling.¹⁵ Both PI3K and AKT are important intracellular signals regulating various physiological and pathological processes. The PI3K-AKT pathway regulates cell growth, proliferation, survival, cell cycle, metabolism, and inflammation.¹⁵ Research indicates that activation of PI3K-AKT pathway contributes to radioresistance in many cancers including

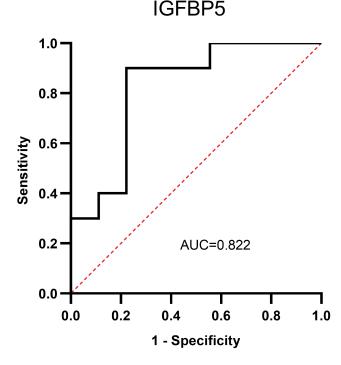


Figure I ROC curve of IGFBP5 expression in radiotherapy-patients of prostate cancer. ROC curve of IGFBP5 with irradiation in prostate cancer patients from previous proteomics analysis. The AUC was 0.822, indicating that IGFBP5 predicts immediate efficacy in these patients.

prostate cancer.^{16–18} Mechanistically, PI3K-AKT induces radioresistance by activating tumor-cell proliferation.^{17,19} Radioresistance is one of the causes of radiotherapy failure in prostate cancer.^{16,20} Thus, regulating PI3K-AKT may be an effective approach to prevent radioresistance.

In this study, we explored whether IGFBP5 influences radiosensitivity in prostate cancer cells. Specific tests were performed to determine whether IGFBP5 regulates the proliferation of PC3 and DU145 cells via PI3K-AKT pathway. The functions of IGFBP5 were explored using IGFBP5 knockdown or overexpression cell models.

Methods and Materials Cell Lines and Cell Culture

Prostate cancer PC3 cells and DU145 cells were bought from Cell Bank, Chinese Academy of Sciences. These cells were cultured in F12K medium and MEM medium (Gibco, MD, USA) supplemented with 10% fetal bovine serum (Gemini Foundation, USA), 100 U/mL penicillin and 100 µg/mL streptomycin. The cells were cultured at 37 °C in a humidified atmosphere with 5% CO₂.

Data Extraction

The expression data of IGFBP5 in cancer and normal tissues was downloaded from the Oncomine database (<u>https://www.oncomine.org/</u>), based on the selection filtering p<0.01, and fold change > 1.5.

Cell Transfection

Cells were plated in six-well plates and transfected with lentivirus to overexpress or knockdown IGFBP5 expression. The lentivirus expression vector (GV358) for overexpression IGFBP5 (LV-IGFBP5) was constructed by Genechem (Shanghai, China). To choose a suitable concentration of LV-IGFBP5, PC3 cells and DU145 cells were first seeded in 12well plates and transfected with three different concentrations of the vectors and the resultant fluorescence intensity was measured. Finally, the concentration of 1×10⁸ TU/mL was chosen. The cell lines were transfected in six-well plates for a defined time. During transfection, 2µg/mL puromycin was added to PC3 cells and 1.25 µg/mL puromycin to DU145 cells to stabilize the cells. For effective IGFBP5 knockdown with lentivirus expression vector, sh-RNA-IGFBP5 sequences and lentivirus expression sh-IGFBP5 (LV-IGFBP5-RNAi) were obtained from Genechem (Shanghai, China). The target sequence gcAAGTCAAGATCGAGAGAGA and the vector GV493 were used. The negative control group was a lentivirus vector lacking the gene of interest.

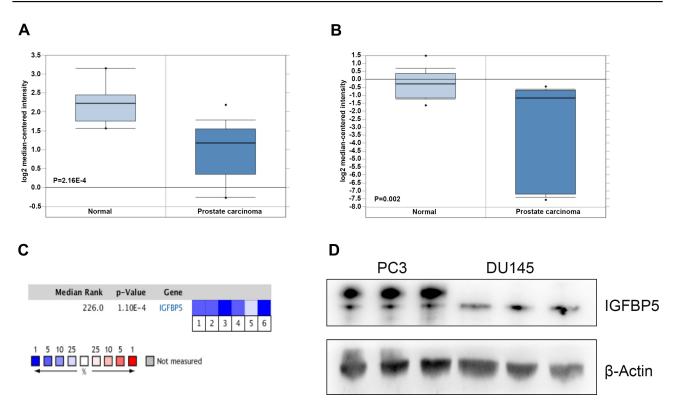


Figure 2 IGFBP5 expression in prostate cancer. (A-C) Data derived from Oncomine database. The *P*-value was set at 0.01 and fold change was defined as 1.5. (A) Comparison of IGFBP5 mRNA expression between prostate carcinoma and normal tissues in the Arredouani Prostate dataset; P=2.16E-4. (B) Comparison of IGFBP5 mRNA expression between prostate carcinoma and normal tissues in the Luo Prostate 2 dataset; P=0.002. (C) A meta-analysis of IGFBP5 gene expression in six Oncomine databases. Colored squares indicate the median rank of IGFBP5 across six databases, Arredouani Prostate (1), Luo Prostate (2), Luo Prostate 2 (3), Tomlins Prostate (4), Welsh Prostate (5), Wooster Cell Line (6). Blue color represents low expression and red color indicates high expression. (D) IGFBP5 protein expression in PC3 and DU145 cells determined by Western blotting, β -Actin serves as a loading control.

Colony Formation

Cells were plated in 60 mm dishes until they reached 70% confluence. They were irradiated with or without treatment. Each group was irradiated with X-ray dose of 2 Gy, 4 Gy, 6 Gy, and 8 Gy. Cells of control group were not irradiated. After irradiation, cells were dissociated by treatment with trypsin (Gibco, Grand Island, NY, USA). They were subsequently re-plated in appropriate dilutions to form colonies for 10–14 days. Colonies were fixed in 4% paraformaldehyde (Beyotime Biotechnology, Shanghai, China), stained with 0.1% crystal violet (Beijing Solarbio Science technology, Beijing, China) and counted using a stereomicroscope. More than 50 cells were plotted according to plating efficiency and results were normalized to the control group.

Western Blotting

Proteins were extracted from cells using the total protein extraction kit (Invent Biotechnologies, Minnesota, USA). The extracted proteins were mixed with loading buffer (TaKaRa, Tokyo, Japan) and heated at 100 °C for 8 min. Proteins were separated by TGX stain-free SDS-PAGE (Bio-Rad, California, USA) under 150 V for 50 min. Next, proteins were electrotransferred to a PVDF membrane at 250 mA for 60 min. The membrane was blocked in 5% non-fat milk for 1 h at room temperature. Next, it was incubated overnight at 4 °C with primary antibodies against IGFBP5 (1:500, Santa Cruz, Texas, USA), PI3K (1:1000, CST, MA, USA), AKT (1:1000, CST, MA, USA), p-AKT (1:1000, CST, MA, USA), beta-Actin and α -tubulin (1:5000, Proteintech, Wuhan, China). Next, the membrane was washed thrice by TBST for 10 min and incubated with secondary goat-anti-mouse antibody (1:5000, Proteintech, Wuhan, China) or goat-anti-rabbit (1:5000, Proteintech, Wuhan, China). Finally, the immunoblot bands were exposed to the enhanced chemiluminescence solution (Thermo Fisher Scientific, Massachusetts, USA) to detect protein bands.

Cell Cycle Assay

Cells were seeded in a six-well plate with or without irradiation at a density of 5×10^4 cells per well. The

irradiation group irradiated with 5 Gy. After 48 h of treatment, samples were collected and diluted to a concentration of 1×10^6 cells/mL. Cells were fixed in 70% cold ethanol at -20 °C overnight. Next, the samples were washed with PBS and centrifuged twice at 800 g, 5 min; followed by staining with Propidium Iodide (PI)/ RNase solution (Dojindo, Kumamoto, Japan) according to the manufacturer's protocol. Finally, flow cytometry (BD Biosciences, USA) was conducted to analyze the cell cycle. The data was analyzed by FlowJo software (Version X; TreeStar, Ashland, OR, USA).

Cell Proliferation Assay

Briefly, cells were plated in 96-well plates at a density of 4×10^3 cells per well, and treated with or without irradiation. Each group was prepared as three repeats. To investigate the proliferation of LV-IGFBP5 PC3 and DU145 cells, cells cultured 48 h after irradiation and then detected by microplate reader. The PI3K inhibitor, LY294002 (25 μ M, Selleck, USA) was added to LV-IGFBP5-RNAi cells and cultured for 24 h. After irradiation, all samples were further cultured for 24 h in the same conditions. Then, 10 μ L of the CCK-8 solution (Dojindo, Kumamoto, Japan) was added to each well. Three control wells were set up without any cells and treatment. The plates were incubated for 1–4 h in an incubator. Finally, absorbance of the cells was measured at 450 nm using a microplate reader. The control group.

Statistical Analysis

Data were analyzed with SPSS software version 23.0 (SPSS, Chicago, IL, USA) and GraphPad Prism Software version 8.01 (GraphPad Software, La Jolla, CA, USA) and are presented as mean \pm SD. Student's *t*-test and Mann–Whitney *U*-test were used to compare two group means. One-way ANOVA was performed for more than three groups. A *P* value less than 0.05 was considered statistically significant.

Results

Higher IGFBP5 Expression Correlates with Better Radiotherapy Efficacy

We previously applied proteomics to explore the efficacy of radiotherapy in prostate cancer patients. Results showed that higher expression of IGFBP5 predicts better efficacy (Figure 1, AUC=0.822, P=0.02). We re-analyzed these data with pathway enrichment analysis. Results showed that

IGFBP5 is highly correlated with PI3K-AKT signaling pathway. Details of these results were reported in the 58th conference of particle therapy co-operative group (PTC58-0345, unpublished data, <u>Supplementary 1</u>). This study was performed to investigate the function of IGFBP5 on irradiation efficacy in prostate cancer.

Expression of IGFBP5 in Prostate Cancer

The expression level of IGFBP5 in prostate cancer was determined by analyzing six independent microarray datasets from Oncomine database.^{21–25} We found that IGFBP5 is expressed in multiple cancers including prostate cancer (Table 1). Notably, the expression of IGFBP5 in prostate cancer tissues was significantly lower than in normal prostate gland tissues (Table 1). IGFBP5 mRNA expression in prostate carcinoma was lower compared to normal tissues (Figure 2A, P=2.16E-4; Figure 2B, P=0.002).^{21,23} The median rank of IGFBP5 among other downregulated genes in prostate cancer was 226 as revealed by a metaanalysis performed across the six datasets (Figure 2C, P=1.10E-4). Moreover, Western blotting assay revealed that IGFBP5 protein was expressed in PC3 cells and DU145 cells (Figure 2D).

Overexpression of IGFBP5 Enhances Radiosensitivity in Prostate Cancer

Previous studies found that IGFBP5 affects cancer development and progression. To date, it remains unknown whether IGFBP5 influences radiosensitivity of prostate cancer. We, therefore, explored the association of IGFBP5 expression level with radiosensitivity in prostate cancer cells using colony formation analysis. The effect of overexpression and knockdown IGFBP5 in PC3 cells and DU145 cells was showed in Figure 3. It was found that IGFBP5 overexpression in PC3 and DU145 cells decreased the survival fraction (Figure 4A and B). We preformed three identical experiments. After overexpression of IGFBP5, the mean survival fraction of PC3 cells was 0.6, 0.17, 0.06, 0.01 with defined irradiation, which significantly reduced survival fraction when compared to control group (0.73, 0.32, 0.32)0.08, 0.03, respectively). Similarly, the mean survival fraction of DU145 cells was 0.56, 0.29, 0.14, 0.07 with defined irradiation after overexpression IGFBP5, which was lower than control group (0.64, 0.45, 0.18, 0.08, respectively). These results suggest that overexpression of IGFBP5 enhances radiosensitivity.

Datasets (Sample Size)	Comparison Groups	Fold Change	P value	Under-Expression Gene Bank
Arredouani Prostate (21)	Prostate carcinoma vs Normal	2.356	2.16E-4	285 (in top 2%)
Luo Prostate (25)	Prostate carcinoma vs Benign Prostatic Hyperplasia	1.784	4.73E-6	167 (in top 4%)
Luo Prostate 2 (30)	Prostate carcinoma vs Normal	6.375	0.002	105 (in top 1%)
Tomlins Prostate (101)	Prostatic Intraepithelial Neoplasia vs Normal	5.486	2.93E-4	360 (in top 4%)
Welsh Prostate (34)	Prostate carcinoma vs Normal	1.946	0.001	996 (in top 12%)
Wooster Cell Line (318)	Prostate cancer	3.798	6.65E-16	61 (in top 1%)

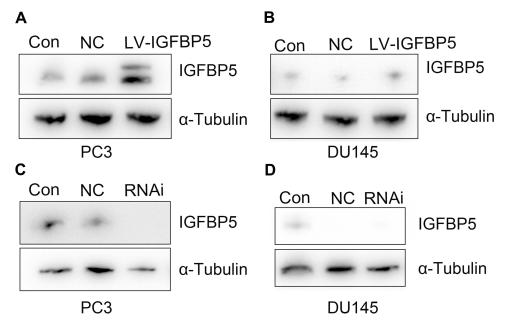
Table I IGFBP5 Expression in 6 Datasets of Oncomine Database About Prostate Cancer

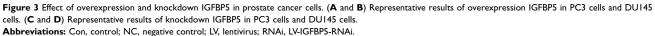
IGFBP5 Induced G2/M Phase Arrest in Prostate Cancer Cells

In further tests, cell cycle analysis was performed to investigate whether IGFBP5 overexpression enhanced radiosensitivity. Results showed that IGFBP5 overexpression in PC3 cells and DU145 cells induced G2/M phase arrest (P<0.05, Figure 4C and D). We have performed three times experiments to confirmed our results. The mean DNA content of G2/M phase in PC3 cells was 21% (LV-IGFBP5 group) versus 16.14% (control group, *P<0.05). The mean DNA content of G2/M phase in DU145 cells was 29.45% (LV-IGFBP5 group) versus 24.84% (control group, *P<0.05). G2/M phase arrest was reversed by IGFBP5 knockdown in PC3 cells and DU145 cells (*P<0.05, **P<0.01, Figure 4E and F). The mean DNA content of G2/M phase in PC3 cells was 18.62% (LV-IGFBP5-RNAi group) versus 28.35% (control group, **P<0.01). The mean DNA content of G2/M phase in DU145 cells was 24.66% (LV-IGFBP5-RNAi group) versus 26.63% (control group, *P<0.05).

IGFBP5 Regulates PI3K-AKT Pathway in Prostate Cancer Cells

Next, we elucidated the mechanism by which IGFBP5 regulated radiosensitivity. For this purpose, we established stable cell lines with IGFBP5 overexpression and IGFBP5 knockdown by lentivirus vector transfection. Results showed that IGFBP5 overexpression deceased the expression of PI3K and p-AKT in cells exposed to defined dose-irradiation (Figure 5A). By contrast, IGFBP5 knockdown enhanced the expression of p-AKT (Figure 5B), indicating PI3K-AKT pathway activation. The quantification of these results was also presented (Figure 5C–E). One-way ANOVA analysis indicated p<0.05.





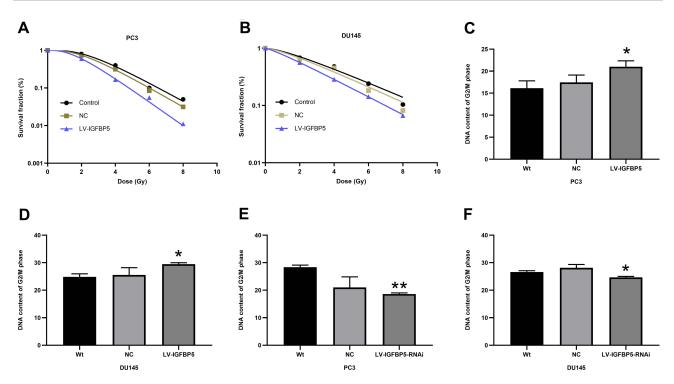


Figure 4 Overexpression of IGFBP5 enhanced radiosensitivity and induced G2/M phase arrest in prostate cancer cells. (A) Representative colony formation results of PC3 cells, LV-IGFBP5 group exhibited high sensitivity to irradiation. (B) Representative colony formation results of DU145 cells, LV-IGFBP5 group exhibited high sensitivity to irradiation. (Cand D) Overexpression of IGFBP5 in PC3 cells and DU145 cells induced G2/M phase arrests. (E and F) Knockdown of IGFBP5 expression reduced G2/M arrest in PC3 cells and DU145 cells. *P<0.05, **P<0.01.

Abbreviations: Wt, wild type; NC, negative control; LV, lentivirus.

IGFBP5 Modulates Cell Proliferation in Prostate Cancer Cells

The effect of IGFBP5 on the proliferation of prostate cancer cells lines was determined using the CCK8 assay. Each experiments performed triplicate times. The results demonstrated that the cell growth of cells overexpressing IGFBP5 was lower compared to that of cells from the control group (Figure 6A and B). The mean OD value of LV-IGFBP5 group in PC3 and DU145 cells was 1.06 and 1.56, while the control group was 1.2 (p<0.001) and 1.28 (p<0.0001). The effect of inhibition also observed by combining with irradiation, that is, irradiation inhibited cell proliferation compared to without irradiation in overexpression IGFBP5 cells (Figure 6C and D). The mean OD value of IR group in PC3 and DU145 cells was 0.73 and 0.80, while the control group was 1.07 and 1.28 (p<0.0001).

PI3K Inhibitor Reverses the Effect of IGFBP5 Knockdown on Cell Proliferation

We further tested whether inhibition of PI3K-AKT pathway with PI3K inhibitor, LY294002, could reverse the effect of IGFBP5 knockdown on cell proliferation. We found that LY294002 slowed the growth of PC3 cells and DU145 cells caused by IGFBP5 knockdown (Figure 6E and F). The mean OD value of LY294002 group versus control group was 0.46 versus 1.16 (p<0.0001) in PC3 knockdown cells and 0.58 versus 1.29 (p<0.0001) in DU145 knockdown cells. Interestingly, the inhibition of cell proliferation by LY294002 was more pronounced when combined with irradiation (****P<0.0001, Figure 6E and F) than LY294002 alone group (#P<0.05, ##P<0.01, Figure 6E and F). The mean OD value of IR+ LY294002 group versus LY294002 group was 0.31 versus 0.46 (p<0.05) in PC3 knockdown cells and 0.41 versus 0.58 (p<0.01) in DU145 knockdown cells.

Discussion

This pioneering study demonstrates that IGFBP5 influences radiosensitivity in prostate cancer via PI3K-AKT pathway. The human *IGFBP5* is a protein-coding gene, located on chromosome 2q33-q36.^{26,27} IGFBP5 comprises six family members, all of which bind insulin-like growth factors (IGFs) which modulate the bioavailability and function of IGFs.^{12,28} Recent studies showed that IGFBPs participate in cancer development and progression. Other studies have

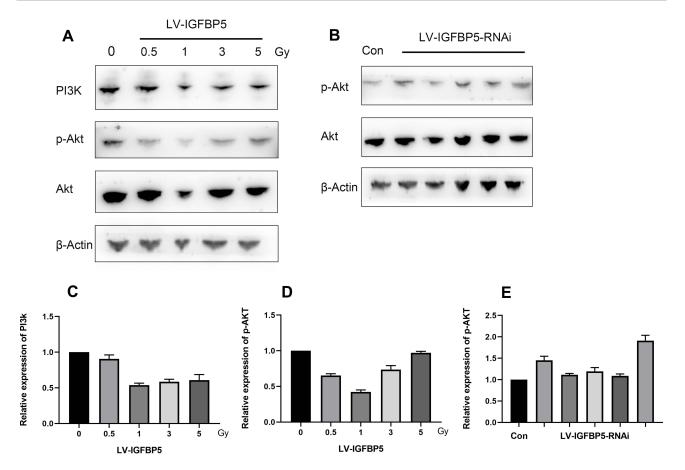


Figure 5 IGFBP5 regulated PI3K-AKT pathway in prostate cancer cells. (A) Overexpression of IGFBP5 decreased PI3K and p-AKT expression in response to irradiation. (B) Knockdown IGFBP5 increased p-AKT expression, indicating PI3K-AKT pathway activation. (C) Results of PI3K expression after overexpression of IGFBP5. To compare with control group, overexpression of IGFBP5 decreased PI3K expression (p<0.05). (D) Results of p-AKT expression after overexpression of IGFBP5. To compare with control group, overexpression of IGFBP5 inhibited p-AKT expression (p<0.05). (E) Results of p-AKT expression after knockdown IGFBP5. To compare with control group, knockdown IGFBP5 increased p-AKT expression (p<0.05). (E) Results of p-AKT expression after knockdown IGFBP5. To compare with control group, knockdown IGFBP5 increased p-AKT expression (p<0.05). (E) Results of p-AKT expression after knockdown IGFBP5. To compare with control group, knockdown IGFBP5 increased p-AKT expression (p<0.05).

reported their roles in predicting the prognosis of pancreatic cancer, ovarian cancer, and nasopharyngeal carcinoma.²⁹⁻³¹ IGFBP5 was first recognized to stimulate osteoblast mitogenesis dependent or independent of IGF.³² Several other studies have shown that IGFBP5 is a growth factor that increases bone formation.³³ Gyanendra et al reported that IGFBP5 inhibited cell proliferation independent of IGF signaling.³⁴ Accumulating evidence show that IGFBP5 inhibits or stimulates cell proliferation in several types of cancers leading to different outcomes. Some studies reported that IGFBP5 functioned as a growth inhibitor in breast cancer³⁵ and suppressed tumor metastasis in osteosarcoma.¹⁴ In contrast, Li et al found that IGFBP5 promoted cancer development and acted as a prognostic factor in breast cancer.³⁶ To date, few studies have explored the role of IGFBP5 in prostate cancer, and to our knowledge, none has reported the association of IGFBP5 with radiosensitivity in prostate cancer.

In the current study, we show that IGFBP5 overexpression combined with irradiation enhances radiosensitivity in prostate cancer cells. Notably, the survival fraction of cells in the LV-IGFBP5 group was significantly lower compared to that of control group. Overexpression of IGFBP5 induced G2/M cell cycle arrest. It has been shown that G2 and M phases are sensitive to radiations.³⁷ Evidence from our proteomic analysis indicated that higher expression of IGFBP5 predicted better radiotherapy efficacy in prostate cancer patients. Pathway analysis suggested that IGFBP5 was largely associated with the PI3K-AKT pathway. Consistently, our results show that overexpression of IGFBP5 decreased PI3K and p-AKT expression. These results demonstrate that IGFBP5 regulates radiosensitivity via the PI3K-AKT signaling pathway. Notably, inhibition of this pathway with PI3K inhibitor partially reversed the effects of IGFBP5 knockdown on cell viability.

Numerous studies indicate that PI3K-AKT pathway contributes to radioresistance in many cancers including

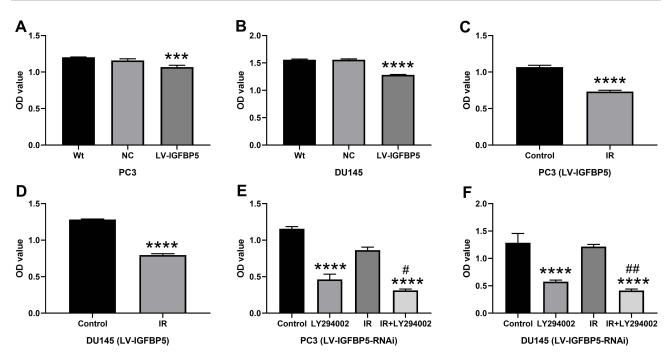


Figure 6 The effect of IGFBP5 on the proliferation of prostate cancer cells. (A) Overexpression of IGFBP5 reduced the growth of PC3 cells; (B) overexpression of IGFBP5 inhibited the proliferation of DU145 cells; (C) radiation inhibited cell proliferation in LV-IGFBP5 PC3 cells; (D) radiation inhibited cell proliferation in LV-IGFBP5 DU145 cells; (E and F) PI3K inhibitor, LY294002, administered with or without irradiation, rescued the effect of IGFBP5 knockdown in PC3 cells and DU145 cells. ***P<0.001, ****P<0.0001 compared to control group; #P<0.05, ##P<0.01 compared to LY294002 group. Abbreviations: Wt, wild type; NC, negative control; LV, lentivirus; OD, optical density.

prostate cancer.^{16,17} Radioresistance is the leading cause of radiotherapy-related failure. Thus, our findings imply that regulating PI3K-AKT pathway may be an effective strategy to prevent radioresistance and thus improve the therapeutic efficacy. These findings suggest that targeting IGFBP5 signaling could be a potential therapeutic strategy for enhancing radiosensitivity in prostate cancer. However, some limitations exist in this study. IGFBP5 overexpression induced G2/M phase arrest, which was associated with radiosensitivity. We did not investigate the detail of G2/M phase protein expression. G2/M phase arrest enhanced radiosensitivity in prostate cancer cells, which was consistent with Geldof et al reported.³⁸ Further studies to investigate the function of IGFBP5 will be meaningful. One the one hand, further studies in animal models are needed; on the other, clinical studies to investigate the status of IGFBP5 in a large number of patients with radiotherapy should performed in the future.

Conclusion

In conclusion, this study shows that IGFBP5 overexpression enhances radiosensitivity in prostate cancer via the PI3K-AKT pathway. This suggests that IGFBP5 can be leveraged to develop a novel strategy to improve prostate cancer radiotherapy.

Abbreviations

Con, control; IGFBP5, insulin-like growth factor-binding protein 5; NC, negative control; LV, lentivirus; Wt, wild type.

Ethics Approval and Consent to Participate

This study is an in vitro experiment, Cell lines bought from Cell Bank, Chinese Academy of Sciences.

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Disclosure

The authors declare no conflicts of interest.

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