

学校编码：10384

学号：24520141153540

廈門大學

硕士学位论文

胰岛素样生长因子2 mRNA结合蛋白
2 (IGF2BP2) 在肺癌发生发展中的作用研究

The role of insulin-like growth factor 2
mRNA binding protein 2 (IGF2BP2) in the
development of lung cancer

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专业名称：外科学

答辩日期：2017年5月

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摘要

背景：当今世界，癌症成为威胁人类健康的头号杀手，其中肺癌更是最常见的恶性肿瘤之一，由于其早期症状不明显，恶性程度高，发现时往往已是中晚期，其每年发病率和死亡率均位于所有癌症的前列。尽管手术方式和化学药物不断更新，但均不能达到较好的治疗效果，肺癌的发病率和死亡率仍没有降低的趋势。因此，早期发现、早期治疗成为治愈肺癌的关键。随着分子生物学的进展，肿瘤相关抗原（TAA）逐渐成为了现代医学的关注热点，因为它不仅能使肿瘤患者早期得到诊断，而且也为分子靶向治疗提供了分子基础。类胰岛素生长因子2 mRNA结合蛋白（IGF2BP2）是一种细胞质蛋白，被证明和多种肿瘤的发生发展有关，如肝癌、乳腺癌、神经胶质瘤、食管癌等，但与肺癌的关系尚不清楚。

目的：本研究旨在通过对IGF2BP2在肺癌中的表达量及其对肺癌细胞生物学行为的影响进行研究，以便初步认识IGF2BP2与肺癌的关系，并初步阐明IGF2BP2在肺癌发生发展的作用。

方法：本研究分别从组织水平和细胞水平来研究分析IGF2BP2在肺癌中的表达及其临床意义。首先在组织水平上，通过收集临床肺癌组织标本，提取组织蛋白和RNA，利用Western Blot、RT-PCR技术统计分析IGF2BP2的表达特点；在细胞水平，通过Western Blot、RT-PCR、细胞转染等技术研究肺正常细胞与肺癌细胞IGF2BP2的表达差异以及对细胞生物学特征的影响。

结果：结果发现，肺癌组织中的IGF2BP2表达量明显多于癌旁正常肺组织（ $P < 0.01$ ），IGF2BP2在肺癌中的表达与患者性别和肿瘤大小有明显统计学关系（ $P < 0.01$ ），与年龄和临床分期没有明显的统计学关系（ $P > 0.05$ ）；肺癌细胞中IGF2BP2的表达量比肺正常细胞的表达量多（ $P < 0.01$ ）；沉默肺癌细胞中IGF2BP2的表达降低了细胞的增殖和迁移侵袭能力（ $P < 0.01$ ）。

结论：IGF2BP2促进了肺癌的增殖和侵袭，在肺癌的发生发展中起到促进作用，这为临床肺癌的早期诊断和预后提供一定的科学实验依据。

关键词：肺癌；IGF2BP2；TAA

Abstract

Background: At present, cancer has become the leading threat to the health of mankind. The lung cancer is one of the most common malignant tumors worldwide. For the hidden symptoms and high malignant of initial lung cancer, it is always late when the disease is diagnosed and the morbidity and mortality rates are always at the top. Despite of continuous development of surgery and chemical drugs, the treatment outcome is not satisfied. Early diagnosis and timely treatment become the key to prevent the human's health from lung cancer. With the development of molecular biology, tumor-associated antigen (TAA) has been gradually becoming the hot spot of modern medical attention. Not only could patients be diagnosed in the early stage, but also the TAA can provide the basis of tumor targeted therapy. Insulin-like growth factor 2 mRNA binding protein 2 (IGF2BP2) is a kind of cytoplasmic protein, which has been proved having a tight connection with many tumors, such as hepatocellular carcinoma, breast cancer, glioma, esophageal carcinoma, etc. But the relationship of IGF2BP2 and lung cancer is still not clear.

Objective: This experiment aims to test the expression level of IGF2BP2 in the lung cancer tissues and study the effects of IGF2BP2 on the biological behavior on lung cancer cells. So that we can explore the association of IGF2BP2 and lung cancer, and to clarify the IGF2BP2's effects on the lung cancer.

Methods: We explored and analyzed the IGF2BP2 expression and its clinical value through the organization and cell level. Firstly, we collect the lung cancer tissue samples, from which we extracted the total proteins and RNAs. Then we made an analysis through the Western Blot, RT-PCR technology. On the other hand, we explored the influence of expression difference on the biological characteristics of IGF2BP2, using the Western Blot, RT-PCR and cell transfection technique on the cell level.

Results: The result shows that the IGF2BP2 in cancer tissues expresses significantly more than that in para-carcinoma tissues($P < 0.05$). The IGF2BP2 in cancer cells expresses significantly more than that in normal lung cells ($P < 0.01$). Knocking out the IGF2BP2 gene in lung cancer cells could reduce the ability of cell proliferation and migration ($P < 0.01$)

Conclusion: IGF2BP2 contributes to proliferation and invasion of the lung cancer cells, and plays an important role in the development of lung cancer, which establishes the scientific basic for the clinical early diagnosis and prognosis of lung cancer.

Keywords: Lung cancer; IGF2BP2; TAA

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