

**IKBKE GENE-A NEWLY POTENTIAL THERAPEUTIC TARGET FOR TREATING NON-SMALL CELL LUNG CANCER**

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**ABSTRACT**

The IKBKE gene is part of a family of enzyme complexes involved in increasing cellular inflammation. IKBKE over expression has been associated with breast and prostate cancers. However, it had not been linked to environmental carcinogen, such as tobacco smoke, until now.

Tobacco smoke is the strongest documented initiator and promoter of lung cancer. The traditional model holds that tobacco components promote carcinogenesis through a process that leads to DNA damage.

**KEY-WORDS:** IKBKE, Lung cancer, Gene, Tobacco

**INTRODUCTION**

Recent studies have shown that tobacco smoke can also promote lung cancer through changes in the pathways that regulate cell proliferation and survival. This study explored identifying and understanding one of the signaling pathways in order to find potential drug targets to treat non-small cell lung cancer. In this study, IKBKE was

found to be induced by two tobacco carcinogens: nicotine and a nicotine-derived nitrosamine ketone found in tobacco smoke. Their findings suggest that IKBKE is a key molecule related to tobacco-induced lung cancer. Since IKBKE kinase is induced by tobacco, small molecular inhibitors of IKBKE could have a therapeutic drug potential for lung cancer<sup>1,2</sup>.

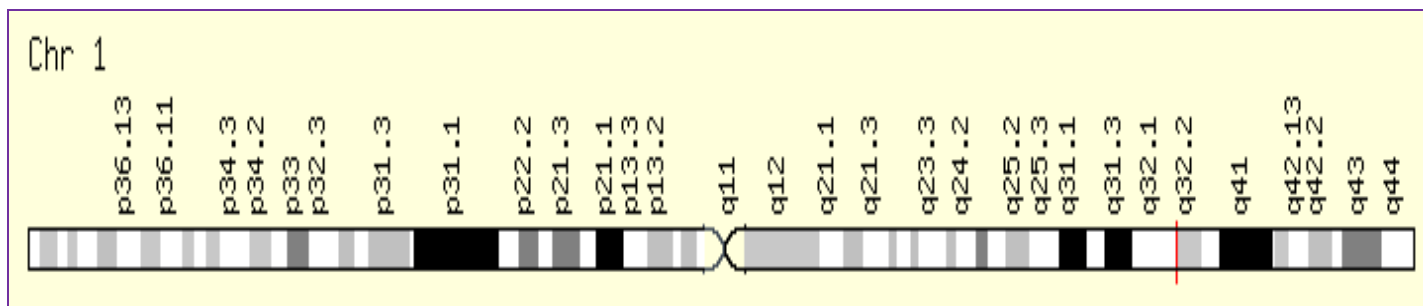


Figure 1: IKBKE Gene in genomic location<sup>3</sup>

**OTHER NAMES<sup>4,5</sup>:**

1. IKK-E
2. IKKI
3. IKK-i
4. IKKE

**ENTREZ GENE SUMMARY FOR IKBKE:**

IKBKE is a noncanonical I-kappa-B kinase (IKK) that is essential for regulating antiviral signaling pathways. IKBKE has also been identified as a breast cancer oncogene and is amplified and over expressed in over 30% of breast carcinomas and breast cancer cell lines<sup>5</sup>.

**ASSOCIATED RESEARCH ABOUT IKBKE GENE:**

Researchers at Moffitt Cancer Center have found a potential targeted therapy for patients with tobacco-associated non-small cell lung cancer. It is based on the newly identified oncogene IKBKE, which helps regulate immune response<sup>2</sup>.

The study appeared in the Feb. 13 online issue of Oncogene. The IKBKE gene is part of a family of enzyme complexes involved in increasing cellular inflammation. IKBKE overexpression has been associated with breast and prostate cancers. However, it had not been linked to environmental carcinogen, such as tobacco smoke, until now. Tobacco smoke is the strongest documented initiator and promoter of lung cancer. The traditional model holds that tobacco components promote carcinogenesis through a process that leads to DNA damage. Recent studies have shown that tobacco smoke can also promote lung cancer through changes in the pathways that regulate cell proliferation and survival. This study explored identifying and understanding one of the signaling pathways in order to find potential drug targets to treat non-small cell lung cancer. In this study, IKBKE was found to be induced by two tobacco carcinogens: nicotine and a nicotine-derived nitrosamine ketone found in tobacco smoke. Their findings suggest that IKBKE is a key molecule related to tobacco-induced lung cancer.

"Since IKBKE kinase is induced by tobacco, small molecular inhibitors of IKBKE could have a therapeutic drug potential for lung cancer," explained lead author Jin Q. Cheng, senior member of the Molecular Oncology Department at Moffitt. Current treatments for non-small cell lung cancer include surgery, radiotherapy and chemotherapy. However, patients eventually develop resistance to treatment. There is a great need to better understand the molecular mechanism of resistance and develop new gene-targeted therapies that can circumvent resistance, said the authors. In this study, the researchers

also reported for the first time that IKBKE is a target of STAT3, a transcription factor that plays a key role in many cellular processes, such as cell growth and programmed cell death.

According to the researchers, STAT3 is frequently activated in various types of human cancers and, when activated, STAT3 increases IKBKE overexpression and protein levels. In non-small cell lung cancer, nicotine-induced IKBKE depends on STAT3. The authors noted that the activation stage of STAT3 represents an attractive therapeutic potential because IKBKE is a STAT3 target. While IKBKE induces chemotherapy resistance, knocking down IKBKE sensitizes cancer cells to chemotherapy and reduces cancer cell survival.

"Since the IKBKE kinase overexpression is induced by tobacco smoke and IKBKE levels increase in response to nicotine and nicotine-derived nitrosamine ketone, this evidence can be potentially used to develop a non-small cell lung cancer intervention strategy that targets IKBKE," concluded Cheng. The work was supported by National Cancer Institute Grants CA137041 and P50 CA119997 and James & Esther King Biomedical Research Program 1KG02, 1KD04, and 1KN08<sup>6,7</sup>.

**TREATMENTS FOR NON-SMALL CELL LUNG CANCER:**

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The activation stage of STAT3 represents an attractive therapeutic potential because IKBKE is a STAT3 target. While IKBKE induces chemotherapy resistance, knocking down IKBKE sensitizes cancer cells to chemotherapy and reduces cancer cell survival<sup>1,2,6</sup>.

**CONCLUSION:**

Tobacco smoke is the strongest documented initiator and promoter of lung cancer. Since the IKBKE kinase over-expression is induced by tobacco smoke and IKBKE levels increase in response to nicotine and nicotine-derived nitrosamine ketone, this evidence can be

potentially used to develop a non-small cell lung cancer intervention strategy that targets IKBKE.

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