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## Kuch Kuch Hota Hai 720p

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Kong led by Dr. Barbara Zuker described these associations in the journal *Alcohol Health & Behavior*. The overall risk of developing alcohol dependence in the study was estimated to be between three and four times higher for people with certain variants of the gene compared to those with the most common variant, the researchers said, noting that these variants were also associated with a heightened risk of alcoholism across different ethnic groups. One important implication of the study for health is that people who drink more than 100 grams of alcohol per day have an increased genetic risk for alcoholism.

"Although the number of genes involved in alcohol dependence is very large, very few of these genes have been identified, with only a few of these consistently showing an association with alcohol dependence," said Zuker. advertisement "These findings suggest that single nucleotide polymorphisms (SNPs) in alcohol-associated genes may be risk factors for alcohol dependence, which explains why those who drink more than 100 grams of alcohol daily are at higher risk of developing alcoholism," said Zuker. The researchers also found that alcohol-associated genes were not uniformly distributed in all parts of the human genome. They found that the distribution of these genes was 'clumpy,' and that the distribution of alcohol-associated genes showed significant variation within a population. "Our results provide an important clue about the role of genetic variation in the

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biology of alcoholism," said Zuker. "These novel findings demonstrate that more genetic variants are involved in alcohol dependence than previously suspected and that such a common and important

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XRCC1 polymorphism and radiation-induced lung fibrosis.  
Patients with asbestosis, radiation pneumonitis, and

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radiation-induced lung fibrosis (RILF) could develop pulmonary fibrosis after thoracic radiotherapy. DNA repair plays an important role in the development of RILF. The relationship between DNA repair genes and RILF has not been examined. In this study, we identified whether X-ray repair cross-complementing group 1 (XRCC1) polymorphism is associated with RILF. XRCC1 single-nucleotide polymorphisms (SNPs) were examined in 71 patients with RILF and 121 healthy controls by single-base extension method using the MassARRAY system. Genotype distribution of XRCC1 Arg399Gln (Arg399Gln) and Arg194Trp (Arg194Trp) polymorphisms did not differ significantly between RILF and control groups ( $P > 0$ )