

**RISK FACTORS AND GENETIC POLYMORPHISMS OF GSTM1,
GSTT1 AND CYP1A1 IN DEVELOPMENT OF ORAL CANCER IN
AN INDONESIAN POPULATION**

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ABSTRACT

Introduction

Cancer is one of the foremost causes of death after infections and heart diseases in all societies and is the fatal leading disease in the world. Oral cancer is the sixth most common cancer worldwide and the most common in South East Asia. The geographic pattern and trends in incidence of oral cancer vary widely between countries and geographical areas of the world. This difference in distribution of oral cancer worldwide is influenced by their risk factors. The major risk factors of oral cancer which have already been established are tobacco use, alcohol drinking and betel quid chewing. Furthermore, epidemiology studies have also implicated other factors such as genetic susceptibility and the role of diet. Jakarta, the capital city of Indonesia is the most populous place in Indonesia. It is inhabited with more than 50% of Indonesia's population. Jakarta city comprises of inhabitants of various ethnicity and practicing different habits. To date no studies have been undertaken to determine the risk factors including genetic polymorphism of metabolizing gene (GSTM1, GSTT1 and CYP1A1) in the development of oral cancer in the Jakarta population.

Objective/Aim

This study aims to identify and determine the risk factors including genetic polymorphisms in GSTM1, GSTT1 and CYP1A1 in the development of oral cancer in a Jakarta population, Indonesia.

Material and Method

This is a cross sectional study with case-control design involving hospital-based subjects matched for age and sex. Two hundred forty three subjects participated in this study from 5 selected hospitals which represent each district of Jakarta, with a 1:2 ratio of cases and

controls. This study consists of two parts. The first part consists of data collection on the subjects' socio demographic characteristics, risk habits (smoking, alcohol consumption and betel quid chewing) and dietary pattern using a structured questionnaire. The second part consists of laboratory work on blood samples of cases and controls to assess the polymorphism of GSTM1, GSTT1 and CYP1A1 by means of PCR and Restriction Fragment Length Polymorphism (RFLP). The data was analysed using conditional logistic regression (STATA 8) and factor analysis (SPSS12) for dietary pattern.

Results

In terms of socio demographic characteristic, 42.3% of oral cancer cases were more than 49 years of age (mean 47.4 years \pm 12.4), males being more frequently affected than females with a ratio of 6:4 and the number increases with increasing age-group. Ninety percent of cases were married, and the most affected ethnic group was Deutro Melayu (87.7 %).

The most common habit practiced among the cases (54.2 %) and controls (49.3%) was smoking, either as a single habit or in combination with alcohol and betel quid chewing. Smokers accounted for 55.6% of the cancer cases whilst there was an almost similar proportion of smoker and non smoker among the controls (48.8% cases vs 51.2% controls). Among cases and controls, the majority had smoked for more than 10 years. *Kretek* was the most preferred type of cigarettes for both cases and controls. Almost all smoking factors (number of sticks per day, duration of smoking, and type of tobacco and pack-years of exposure of tobacco) displayed higher risks of oral cancer by two to three times after allowing for confounding factors.

Alcohol drinking habit was practiced by less than 10% of case and control groups (8.6% and 4.3%, respectively). The crude OR showed that alcohol drinking habit did not

contribute to the risk of oral cancer ($p>0.250$). The exception was that those who consumed wine had 11 times higher risk of oral cancer.

Similarly, betel quid chewing habit was also practiced by a very small number of case (7.4%) and control subjects (1.8%). The findings showed that duration of chewing and number of quid per day did not significantly contribute to the risk of oral cancer. However, current betel quid chewer and quid combination of betel leaf, tobacco, areca nut, and lime were significantly associated ($p<0.05$) with increasing risk of oral cancer (OR 5.32, 95% CI 1.03-27.52; OR 4.19, 95% CI 1.05-16.82 respectively).

The genetic polymorphism of GSTM1, GSTT1 and CYP1A1 assay showed that there was no statistically significant difference observed between cases and controls. The polymorphism of GSTM1, GSTT1 and CYP1A1 either singly or in combination did not have association with risk of oral cancer even in the smoker, non smoker as well as among betel quid chewer and non chewer in this study.

Analysis of dietary pattern found four factors retained from factor analysis, referred to as preferred food, combination, chemical related and traditional that were associated with oral cancer after adjusting for smoking, alcohol and betel quid chewing. The preferred food consisted of fast food, fermented food, canned food, snacks high in fat and sugar, cooked and raw vegetables, and seafood showed increased risk of oral cancer by two times (OR 2.17, 95% CI 1.05-4.50, χ^2 trend 5.446, $p<0.05$). The “chemical related” pattern (which is consisted of MSG and processed food) also showed an increase risk (OR 2.85, 95% CI 1.34-6.05). The “combination” pattern (consisted of meat, dairy product and fruit) displayed protective effects in relation to oral cancer (adjusted OR 0.46, 95% CI 0.23-0.91, χ^2 trend 7.335, $p<0.01$). Finally, “traditional” pattern (consist of drinks and carbohydrate)

showed an increased of risk by two-fold (OR 2.04, 95%CI 1.01-4.41, χ^2 trend 5.649, $p < 0.05$). The fast food was the most common food type consumed in this study (Communality : 0.818)

Conclusion

Smoking and betel quid chewing were found to be risk factors for oral cancer in the Jakarta population. In contrast genetic polymorphism of GSTM1, GSTT1 and CYP1A1 did not contribute to the development of oral cancer in this study. The dietary pattern referred to as “preferred food”, “chemical related” and “traditional” were found to be risk factors of oral cancer whereas the “combination” pattern had protective effect of oral cancer in this study.

Recommendations

Future research is recommended to include a bigger sample size and greater cross section that constitute the Indonesian population and using community controls. Investigation involving nutritional content of food consumed is also advocated.

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ABBREVIATION

bp	: base pair
BMDP	: BioMeDical Package (statistic software)
CYP	: Cythochrome P450
DNA	: Deoxyribonucleic acid
dNTPs	: deoxyribonucleotide triphosphate
EDTA	: Ethylene Diamine Tetraacetic Acid
FFQ	: Food Frequency Questionnaire
GSTM	: Gluthathione S transferase mu
GSTT	: Glutathione S transeferase tetha
KMO	: Kaiser-Meyer-Olkin (Measure of Sampling Adequacy)
MgCL	: Magnesium Chloride
mL	: milliliter
mM	: milliMolar
OR	: Odds Ratio
PCR	: Polymerase Chain Reaction
RFLP	: Restriction Fragment Lenght Polymoprhism
SCC	: Squamous Cell Carcinoma
95% CI	: 95 % Confidence Interval