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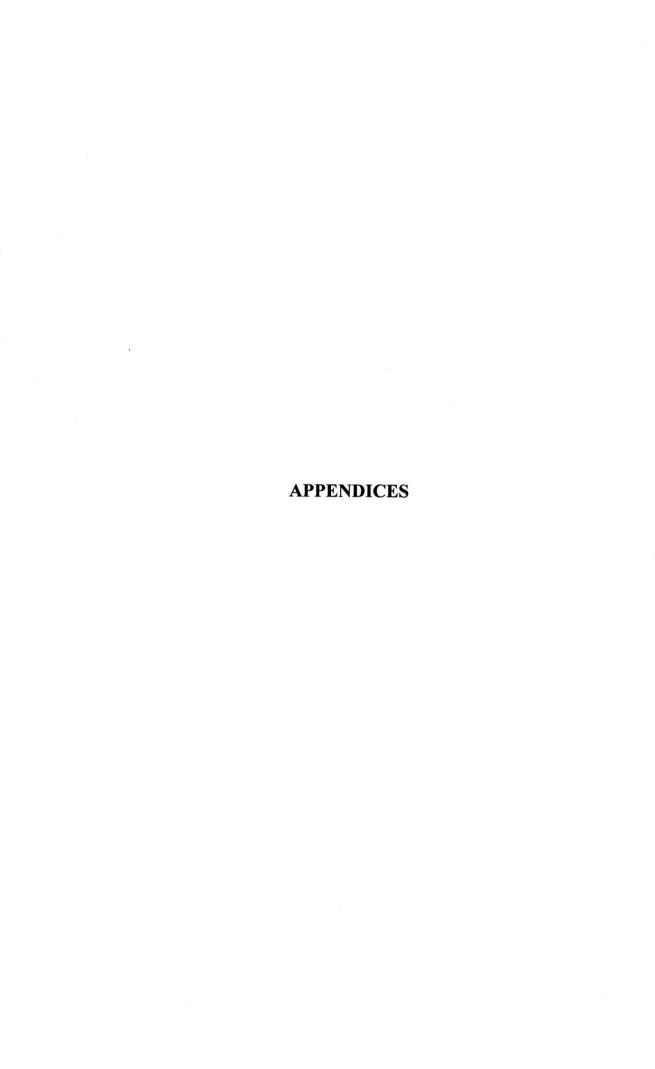
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## APPENDIX A REAGENTS

#### 1. Reagents for cell membrane lysis buffer

0.32 M Sucrose

5% V/V Triton X-100

5 mM MgCl2-6H2O

12 mM Tris-HCl, pH 7.6

Prepare cell membrane lysis buffer in distilled water, autoclave and store at 4°C.

#### 2. Reagents for nuclear membrane lysis buffer

4 M Guanidine HCL

12 mM Tris-HCl, pH 7.6

375 mM NaCl

12 M EDTA, pH 8.0

0.5% Sodium N-Lauroyl Sarcosinate

0.1 M β-Mercaptoethanol

Prepare nuclear membrane lysis buffer in distilled water, autoclave and store at 4°C.

#### 3. Reagents for 10X TBE buffer

0.9 M Tris-Base

0.9 M Boric acid

5 M EDTA, pH 8.0

Prepare 10X TBE buffer in distilled water, autoclave and store at 4°C.

#### 4. Reagents for 6x loading dye buffer

40% Sucrose

0.25% Bromophenol blue

Prepare loading dye buffer in distilled water and store at -20°C.



### มหาวิทยากัยขอนแก้น หนังสือฉบับนี้ให้ไว้เพื่อแสดงว่า

โครงการวิจัยเรื่อง

: ความผิดแผกทางทับธุกรรมของ NAD(P)H-quinone oxidoreductase-i (NQOI) และ glutathione S-transferases (GSTM1, GSTT1 และ GSTP1) กับความเพี่ยงต่อมะเร็ง ท่อน้ำดีในประชากรไทย กาลตะวันขอกเดืองเหนือ

(Polymorphisms of NAD(P)H-quinone oxidoreductase-1 (NQO1) and glutathione S-transferases (GSTM1, GSTT1 and GSTP1) and susceptibility to cholangiocarcinoma in Northcastern Thais)

พัวหนัาโครงการวิจัย : อาจารย์เชื้อมเดือน ประวาพ และคณะ

หม่วยงานที่ขังกัด

: ภาควิชาเภสัชวิทยา คณะแพทยศาสตร์ มหาวิทยาลัยขอบแก่น

ลกสารที่รับรอง

- : I. แบบเสนอเพื่อขอรับการพิจารณาจริยธรรมการวิจัยในมนุษย์ version I.O ฉบับถงวันพี่ 18 กมภาพันธ์ 2552
  - 2. โครงร่างการวิจัย version 1.0 ฉบับลงรันที่ 18 กุมภาพันธ์ 2552
  - แบบบันพึกข้อมูลการวิจัย version 1.0 ฉบับลงวันพี่ 18 ภูมภาพันธ์ 2552

ได้ผ่านการรับของจากคณะกรรมการจริยธรรมการวิจัยในมนุษย์มหาวิทยาลัยขอนแก่น โดยอีด หลักเกณฑ์ตามคำประกาศเธลชิงก็ (Declaration of Helsinki) และแนวทางการปฏิบัติการวิจัยทางคลินิกที่ดี (ICH GCP)

ให้ไว้ ณ วันที่ 3 มีนาคม พ.ศ. 2552

(รองศาสตราชารย์จิรามรณ์ ศรีนิครินทร์

ปวะทานคณะกรรมการจริยธรรมการวิจัยในมนุษย์ ประจำสาขาวิชาทางชีวเวชสาสตร์และการวิจัยทางการแพทย์

มหาวิทยาลัฮขอนแก่น

สำคัญที่ 3.4.01 : 05/2552

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วันหมดอน 2 มีมาคม พ.ศ. 2553

คณะกรรมการจริยธรรมการวิจัยในมนุษย์มหาวิทยาลัยขอนแก่น ห้อง 1733 ชั้น 13 อาคารสมเด็จพระสรีนครินทราบรบราชนที อนุสรณ์ คณะแทพแสสตร์ โทร. (043) 366616-17

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# APPENDIX B LIST OF COMMUNICATIONS

#### **Communications:**

#### 1. POSTER PRESENTATION:

- 1.1 "Genetic polymorphism of NAD(P)H-quinone oxidoreductase-1 in Thais and its association with cholangiocarcinoma" in *The 3rd Asian Pacific Regional ISSX Meeting*. The imperial queen's park Hotel, Bangkok; 10-12 May 2009.
- 1.2 "NAD(P)H-quinone oxidoreductase-1 (NQO1) C609T genotypes in the Thai population" in *The 31st Pharmacological and Therapeutic Society of Thailand Meeting*. Khon Kaen University; 18-20 March 2009. (*Best presentation Award*)

#### 2. PROCEEDING:

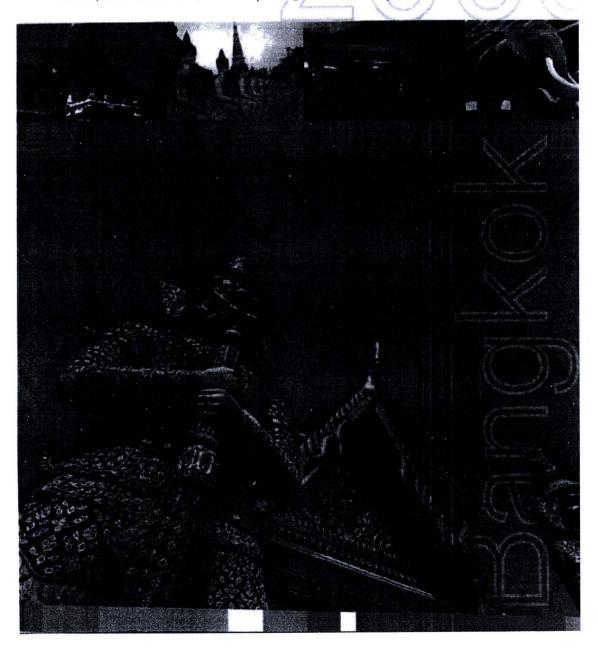
2.1 Zeekpudsa P, Prawan A, Kukongviriyaoan V and Bhudhisawasdi V. NAD(P)H-quinone oxidoreductase-1 (NQO1) C609T genotypes in the Thai population. *Thai J Pharmacol.* 2009; 31(1): 57-60.



## 3rd Asian Pacific Regional Meeting

Understanding Xenobiotics for Better Drug Development and Therapy

The Imperial Queen's Park Hotel | Bangkok, Thailand | 10 - 12 May 2009



#### Abstracts

125. POLYMORPHISMS IN THE PROMOTER REGION OF UGT1A9 GENE IN THAIS

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The human UDP-glucuronosyltransferase, UGT1A9, catalyzes glucuronidation of various endobiotics and xenobiotics. Genetic polymorphisms of *UGT1A9* can influence detoxifying capacities and have considerable effect on the metabolisms of numerous drugs. The purpose of this study was to investigate the polymorphisms in the 5D-flanking region of *UGT1A9* gene in That population. Genomic DNA from 93 healthy unrelated voluntaers was amplified by the polymerase chain reaction (PCR) and DNA sequencing was performed to determine mutations. There was a novel single nucleotide polymorphism (SNP) in *UGT1A9* promoter region, heterozygous -688A>C, with frequency of 0.2473. In addition, three known polymorphisms were found, -440T>C, -331C>T and one base insertion of thymidine resulting in -118A(T)<sub>10</sub>AT (*UGT1A9\*1b*), with frequencies of 0.9785, 0.9677 and 0.5323, respectively. In conclusion, this is the first study to demonstrate the genetic variations in the promoter region of *UGT1A9* gene in That population.

## 126. GENETIC POLYMORPHISM OF NAD(P)H-QUINONE OXIDOREDUCTASE-1 IN THAIS AND ITS ASSOCIATION WITH CHOLANGIOCARCINOMA

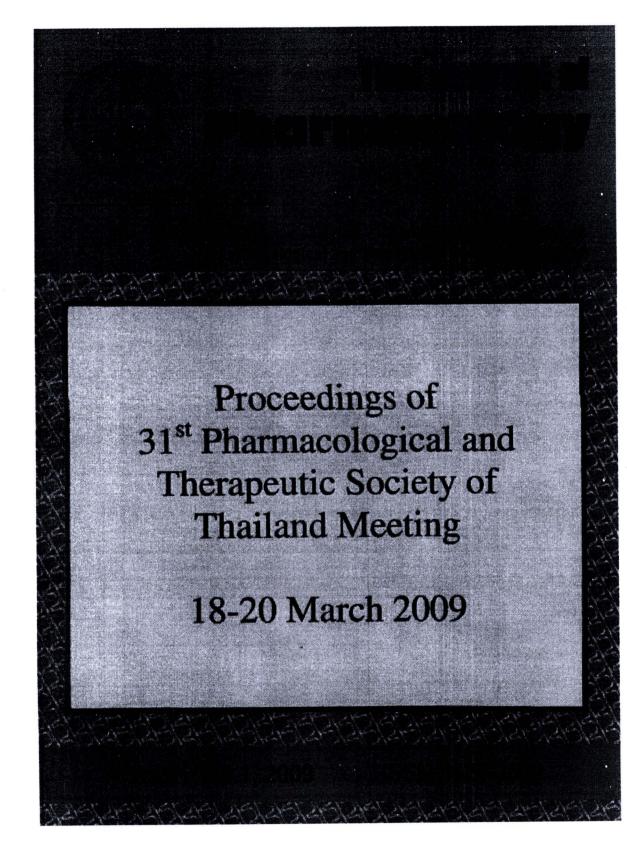
<u>Pornsin Zeekpudsa</u>, Auemduan Prawan, Veerapol Kukongvirlyapan and Vajarabhongsa Bhudhisawasdi Department of Pharmacology, Faculty of Medicine, Liver Fluke and Cholangiocarcinoma Research Center, Khon Kaen University, Khon Kaen, Thailand, 40002

NAD(P)H-quinone exidereductase-1 (NQO1) is a detextlying/antioxidant enzyme that plays a critical role in cellular defense against reactive oxygen species and toxic quinone derivatives, which in turn confer cytoprotection, inhibition of mutation and carcinogenesis. The gene coding for NQO1 has a genetic polymorphism (C->T) at nucleotide position 609 (amino acid codon 187) of the NQO1 cDNA. This mutation has been associated with a decreased enzymatic activity, and increased risk of chemically-induced cytoxicity and susceptibility to various forms of cancer. However, the role of NQO1 polymorphism in relation to carcinogenesis of cholangiocarcinoma (CCA), the most common liver cancer in the Northeast of Thalland, is unknown. Present study, we genotyped the NQO1 C609T polymorphism by PCR-RFLP in 210 CCA patients and 189 healthy control subjects matched for age, sex, and ethnicity. Among 189 Northeastern Thai healthy controls investigated, the NOO1 609T was present at 44% and the frequency distributions of C/C, C/T and T/T genotype were 32%, 53% and 15%, respectively. T allele frequency in the CCA patients was comparable with the healthy controls (p=0.4). In analysis for association of NQO1 genotype and survival time of the CCA patients at the time of diagnosis, the variant NQO1 genotypes (C/T and T/T) were not associated with lower survival time when compared with C/C genotype (p=0.092). Our findings suggested that NQO1 C609T polymorphism may not directly represent a genetic risk factor for CCA, however, the present study cannot exclude NQO1 as a possible modifier for CCA development. Further study in a larger population and biological function of NQO1 gene is required to verify the role of NQO1 in CCA.

### 127. CNS PENETRANT OR NOT? PITFALLS WITH IN VIVO DETERMINATION USING BRAIN TO BLOOD

Zigiang (Zack) Cheng¹, Jinqiang Zhang¹, Rongxia Liu¹, Yiwen Wu¹, Zhen Ge¹, Yi Li¹, Zong-ping Zhang³, Yu Yang¹, Yan Chen¹, Hong Lu¹, Hans Hu², Barry Wang², Jason Meng², Raymond Zhao² and Eric Yang¹¹¹Drug Metabolism and Pharmacokinetics, GlaxoSmithkline, Shanghal, China, 201203, ²NeuroInflammation Discovery Performance Unit, GlaxoSmithkline, Shanghal, China, 201203

Purpose: The total brain to blood ratio (8/8) at single time point obtained from in vivo experiment has traditionally been used to determine whether a molecule is a CNS penetrant. Despite the effectiveness for



Original article

#### NAD(P)H-Quinone Oxidoreductase-1 (NQO1) C609T Genotypes in the Thai population

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#### Abstract

NAD(P)H-quinone oxidoreductase-1 (NQO1) is a detoxifying/antioxidant enzyme that plays a critical role in cellular defense against reactive oxygen species and toxic quinone derivatives, which in turn confer cytoprotection, inhibition of mutation and carcinogenesis. A polymorphism in human NQO1 at nucleotide position 609 (amino acid codon 187) is associated with diminished NQO1 enzymatic activity, and increased risk of chemically-induced cytoxicity and susceptibility to various forms of cancer. The purpose of this study was to determine the NQO1 polymorphism in the Thai population. We genotyped the NQO1 C609T polymorphism by PCR-RFLP in 189 unrelated healthy Thai subjects. The frequency of NQO1 C609 or wild type allele was 58.6%, where those of C/C, C/T and T/T genotypes were 32%, 53% and 15%, respectively. The frequency of NQO1 C609 allele in Thais was closely related to those observed in the East Asian (Oriental) population. Additionally, Thais exhibited a relatively low frequency of NQO1 C609 allele compared to Caucasian and African-American populations. Since, this is the first report on the NQO1 polymorphism in Thai population, data from this study can be use to further evaluate the impact of NQO1 polymorphism on susceptibility to chemically-induced toxicity and cancer risk.

Keywords: NAD(P)H-quinone oxidoreductase-1 (NQO1), Polymorphism, Thai population

#### Introduction

NAD(P)H:quinone oxidoreductase 1 (NQO1) is a detoxifying/antioxidant enzyme that plays an important role in protecting cells against chemically induced oxidative stress, cytotoxicity, mutagenicity, and carcinogenicity. NQO1 protects cells from oxidative damage by preventing the generation of reactive oxygen species and reactive electrophiles from certain environmental carcinogens (1). A genetic polymorphism (C→T) at nucleotide position 609 (amino acid codon 187) of the human NQO1 cDNA was show to reduce NQO1 enzyme activity, which may diminish the protection provided by NQO1. Variation in NQO1 enzyme activities has been suggested to influence an individual ability to metabolize carcinogenic agents and thus to be causally linked to cancer risk (2). Over years, a number of studies on the genetic polymorphism of NQO1 gene in many populations have been documented (4-7). However, no study has been reported on the genetic polymorphism of NQO1 gene in the Thai population. In the present investigation, efforts have been made to understand the genetic distribution of NQO1 polymorphism in Thai population.

57

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