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# epidemiology and molecular characterization of biarribal vibuses in adules

BOOMPA SUANTAL

MASIER OF SCIENCE IN RECROMOLOGY

THE GRADUATE SCEOOL CHIANG MAI UNIVERSITY DECEMBER 2011



# EPIDEMIOLOGY AND MOLECULAR CHARACTERIZATION OF DIARRHEAL VIRUSES IN ADULTS

### **BOONPA SUANTAI**



# A THESIS SUBMITTED TO THE GRADUATE SCHOOL IN PARTIAL FULUILLMENT OF THE REQUIREMENTS FOR THE DEGREE OF MASTER OF SCIENCE IN MICROBIOLOGY

THE GRADUATE SCHOOL
CHIANG MAI UNIVERSITY
DECEMBER 2011

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20 December 2011

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I would like to express my sincere gratitude to Professor Dr. Niwat Maneekarn, my thesis advisor, for his professional guidance and excellent support throughout my study. In addition, I would also like to express my great appreciation to Dr. Pattara Khamrin, my co-advisor, for her excellent guidance during my Master Thesis experiments. Additionally, I would like to give special thanks for all staffs in the Department of Microbiology, Faculty of Medicine, Chiang Mai University who were always provide me with valuable advices and assistances.

Besides my advisors, I would like to thank the examining committee, Associate Professor Dr. Leera Kittigul for her valuable points of suggestions and comments to make my thesis manuscript in a better shape.

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I would also like to express my appreciation to all of my teachers who have been teaching and supporting me in every aspects during my study.

Last but not least, I would like to give my deep gratitude to my parents and sister for their love, attention, understanding, and supporting me spiritually throughout my life.

Boonpa Suantai

Thesis Title

Epidemiology and Molecular Characterization of

Diarrheal Viruses in Adults

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### **ABSTRACT**

Viral gastroenteritis is a common public health problem that causes morbidity and mortality worldwide, particularly in developing countries. The purpose of this study was to investigate the epidemiology and molecular characterization of the viruses detected in fecal samples collected from adult patients with diarrhea. A total of 332 fecal specimens collected between January to December, 2008 were screened for the presence of group A and C rotaviruses, sapovirus, Aichi virus, human parechovirus, noroviruses GI and GII, enterovirus, adenovirus and astrovirus by reverse transcription (RT) and multiplex-PCR methods. The genogroups and genotypes of these viruses were further identified by multiplex-PCR using genotype-specific primers and by nucleotide sequencing. Out of 332 fecal specimens tested, the overall detection rate of diarrheal viruses was 4.2%. Adenovirus and enterovirus were the most predominant viruses detected with the prevalent rate of 1.2%, followed by Aichi virus at 0.9%, and norovirus GII at 0.6%. In addition, mixed infection of norovirus GII and human parechovirus was also detected in one fecal specimen

(0.3%). However, rotaviruses of both groups A and C, astrovirus, norovirus GI and sapovirus were not detected in this study. Based on nucleotide sequence and phylogenetic analyses, the results indicated that all of 3 norovirus GII detected in the present study belonged to GII/4 genotype. Additionally, 3 strains of NoV GII/4 can be subdivided into 2 distinct variant groups, 2006a and 2006b variants. For adenoviruses, 3 genotypes were found to be circulated in this area, including AdV24, AdV25, and AdV40. For enterovirus, 4 genotypes of enterovirus were identified, including echovirus 30, enterovirus 99, poliovirus genotype 3 (PV3), and coxsackievirus A20 (CV-A20). Interestingly, Aichi viruses of both genotypes A and B were also detected in this surveillance. In conclusion, this study provided the molecular epidemiological data of wide variety of diarrheal viruses circulating in adults with diarrhea in Chiang Mai, Thailand.

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โมเลกุลของเชื้อไวรัสก่อโรคอุจจาระร่วงในผู้ใหญ่

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# บทคัดย่อ

E 16251

โรคกระเพาะอาหารและลำไส้อักเสบหรือโรกอุจจาระร่วงที่เกิดจากการติดเชื้อไวรัสยังเป็น
ปัญหาที่สำคัญทางค้านสาธารณสุขทำให้เกิดการเจ็บป่วยและการตายที่พบได้บ่อยทั่วโลกโดยเฉพาะ
อย่างยิ่งในประเทศที่กำลังพัฒนา วัตถุประสงค์ของงานวิจัยนี้เป็นการศึกษาระบาดวิทยาและการ
ตรวจหาคุณลักษณะเฉพาะในระดับโมเลกุลของเชื้อไวรัสก่อโรกอุจจาระร่วงในผู้ใหญ่ โดยนำ
ตัวอย่างอุจจาระจากผู้ใหญ่ที่มีอาการอุจจาระร่วง จำนวนทั้งหมด 332 ตัวอย่าง ที่เก็บในช่วงเคือน
มกราคมถึงเดือนธันวาคม พ.ศ. 2551 มาตรวจคัดกรองหาเชื้อไวรัสโรตา กรุ๊ป A และ C ไวรัสษาโป
ไวรัสไอชิ ไวรัสพารีโก ไวรัสโนโร GI และ GII ไวรัสเอ็นเทอโร ไวรัสอะดิโน และไวรัสแอสโทร
โดยวิธี revese transcription (RT) และ multiplex-PCR หลังจากนั้นนำไวรัสที่ตรวจพบไปจำแนก
genogroups และ genotypes ต่างๆ โดยวิธี multiplex-PCR โดยใช้ primer ที่จำเพาะและโดยการทำ
nucleotide sequencing เพื่อถอดรหัสพันธุกรรมของเชื้อ ผลการตรวจตัวอย่างจำนวน 332 ตัวอย่าง
พบเชื้อไวรัสที่เกี่ยวข้องกับโรกอุจจาระร่วงในผู้ใหญ่ร้อยละ 4.2 โดยพบเชื้อไวรัสอะดิโนและไวรัส
เอ็นเทอโร สูงสุดโดยแต่ละเชื้อพบร้อยละ 1.2 รองลงมาคือ ไวรัสไอชิ ร้อยละ 0.9 ไวรัสโนโร GII
ร้อยละ 0.6 นอกจากนี้ยังพบการติดเชื้อร่วมระหว่างไวรัสโนโร GII กับไวรัสพารีโคในหนึ่งตัวอย่าง
ตรวจคิดเป็นร้อยละ 0.3 อย่างไรก็ตาม การศึกษาในครั้งนี้ตรวจไม่พบไวรัสโรตา กรุ๊ป A และ C

E46251

ไวรัสแอสโทร ไวรัสโนโร GI และไวรัสซาโปเลย ผลจากการตรวจวิเคราะห์ลำดับนิวคลีโอไทด์ และการวิเคราะห์วิวัฒนาการของตัวเชื้อโดยการวิเคราะห์ทาง phylogenetic พบว่าเชื้อไวรัสโนโร GII ทั้งหมด 3 ตัวอย่าง ที่พบในการศึกษาในครั้งนี้จัดอยู่ใน GII/4 genotype ซึ่ง 3 สายพันธุ์ของเชื้อ ไวรัสโนโรที่พบสามารถจำแนกย่อยออกเป็น 2 variants คือ variants 2006a และ 2006b สำหรับเชื้อ ไวรัสอะคิโนที่พบในการศึกษานี้พบว่ามีอยู่ทั้งหมด 3 สายพันธุ์ คือ AdV24, AdV25, AdV40 และ เชื้อไวรัสเอ็นเทอโรพบ 4 สายพันธุ์ คือ echovirus 30, enterovirus 99, poliovirus 3, และ coxsackievirus A20 นอกจากนี้ยังพบว่าเชื้อไวรัสไอชิที่แยกได้จากการศึกษาในครั้งนี้มีทั้งชนิดที่ เป็น genotype A และ B โดยสรุป การศึกษาในครั้งนี้ทำให้ได้ข้อมูลทางด้านระบาดวิทยาของเชื้อ ไวรัสหลากหลายชนิดที่ก่อโรคอุจจาระร่วงในผู้ใหญ่ในจังหวัดเชียงใหม่ของประเทศไทย

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# **ABBREVIATIONS**

% Percent

°C Degree Celsion

μg Microgram

μl Microliter

AdV Adenovirus

AGE Acute gastroenteritis

AiV Aichi virus

AstV Astrovirus

AUS Australia

BGD Bangladesh

BRZ Brazil

bp Basepair

CDC Center for Disease control and Prevention

CHN China

cDNA Complementary deoxyribonucleic acid

DEU Germany

DNA Deoxyribonucleic acid

dNTPs Deoxynucleoside triphosphates

dsRNA Double-stranded Ribonucleic acid

EDTA Ethylenediaminetetraacetic acid

EIA Enzyme immunoassay

ELISA Enzyme-Linked Immunosorbent Assay

**EM** 

Electron microscopy

G

Genogroup

g

Gram

HuCVs

Human caliciviruses

**HPeV** 

Human parechovirus

**FRA** 

France

**IEM** 

Immune electron microscopy

IND

Indonesia

JPN

Japan

kp

Kilobasepair

kDa

Kilodaton

KOR

Korea

M

Molar

mAbs

Monoclonal antibody

mg

Milligram

min

Minute

ml

Milliliter

mM

Millimolar

mRNA

Messenger ribonucleic aacid

MW

Molecular weight

**NEP** 

Nepal

nm

Nanometer

ng

Nanogram

NoV

Norovirus

#### xvii

NoV GI

Norovirus genogroup I

NoV GII

Norovirus genogroup II

**NSPs** 

Nonstructural proteins

**NZL** 

New Zealand

**ORFs** 

Open reading frames

**PAGE** 

Polyacrylamine gel electrophosis

**PBS** 

Phosphate buffer saline

**PCR** 

polymerase chain reaction

**RNA** 

Ribonucleic acid

**RNase** 

Ribonuclease

rpm

Revolutions per minute

RT

Reverse transcriptase

RT-PCR

Revese-transciption polymerase chain reaction

rVLP

Recombination virus-like particle

RV

Rotavirus

SaV

Sapovirus

sec

Second

SG

Subgroup

SIG

Singapore

ssRNA

Single-stranded ribonucleic acid

**TBE** 

Tris-borate EDTA

THA

Thailand

TUN

Tunisia

# xviii

UK

United Kingdom

US

United State

VIE

Vietnam

VPs

Viral proteins