

UNIVERSITI TEKNOLOGI MARA

**EFFECTS OF HUMAN
APOLIPOPROTEIN A1, B AND E IN
MODULATING THE
INTERNALISATION AND
REPLICATION OF DENGUE VIRUS
SEROTYPES 1 AND 2 IN HUH-7
CELL LINES *IN VITRO***

RADZI IKHSAN BIN AHMAD

Thesis submitted in fulfilment
of the requirements for the degree of
Doctor of Philosophy
(Medicine)

Faculty of Medicine

June 2025

ABSTRACT

Dengue virus (DENV) causes a range of diseases, from mild fever to severe dengue, with internalisation into target cells being a critical step in infection. While previous studies have revealed a direct correlation between severe dengue disease and changes in plasma lipid profiles using clinical data, the underlying mechanisms remain unclear. Thus, this study takes a reverse approach by examining the effects of lipid profile modulation *in-vitro* to uncover these mechanisms. Specifically, this study aims to compare DENV internalisation in Huh-7 cells in the presence of apolipoproteins A1 (ApoA1), B (ApoB), and E (ApoE). Tetrazolium reduction assays were used to assess the cytotoxicity of apolipoproteins on Huh-7 cells. DENV-1 or DENV-2 at an M.O.I. of 1, along with 2 µg/mL of apoA1, apoB, or apoE, was introduced into confluent Huh-7 cells and incubated for one hour at 37 °C with 5% CO₂. Unbound virus was then removed. For direct measurement of viral attachment, the cells were lysed, and the cell lysate was collected. For indirect measurement, fresh media was added to the cells, followed by incubation for another 72 hours. The supernatant was collected to measure the virus released from Huh-7 cells. SR-B1 and LDLR knockdown was achieved by transfecting 60% confluent Huh-7 cells with specific siRNA for 48 hours. The indirect measurement experiments were then repeated after siRNA treatment. Collected samples were subjected to qPCR for viral load determination. Differences between groups were analyzed using ANOVA and an independent t-test. DENV2 RNA levels increased with each apolipoprotein treatment compared to controls, with ApoB showing the highest and significant increase (p=0.031), while DENV1 viral load showed no significant differences. After 72 hours, DENV2 RNA levels rose significantly in ApoA1- and ApoB-treated cells (p=0.008 and p=0.015) compared to ApoE, with ApoA1 showing the highest enhancement (101.31%). siRNA knockdown of the scavenger receptor class B type 1 (SR-B1) by 51% reduced the ApoA1-mediated increase in DENV infection by 33% (p=0.023). Similarly, a 39% knockdown of LDL receptor (LDLR) decreased ApoA1- and ApoB-enhanced infection by 58.2% (p=0.014) and 60.5% (p=0.002), respectively. In conclusion, ApoA1 and ApoB significantly enhance DENV internalisation, likely by facilitating initial attachment to cell surface receptors, with SR-B1 and LDLR potentially playing a role in DENV infectivity. These findings contribute to a deeper understanding of how lipid pathways facilitate viral entry and suggest that targeting apolipoprotein interactions and receptor pathways could inform therapeutic strategies against DENV infection.

ACKNOWLEDGEMENT

Firstly, I wish to thank God for granting me the opportunity to embark on my PhD journey and for enabling me to complete this long and challenging path successfully. My deepest gratitude and thanks go to my supervisor, Assoc. Prof. Dr. Thuhairah Hasrah Abdul Rahman from the Faculty of Medicine, UiTM Sungai Buloh Campus, for her invaluable guidance, encouragement, and unwavering support throughout this research and the preparation of my dissertation. I also extend my sincere thanks to the members of my supervisory committee, Assoc. Prof. Dr. Fadzilah Mohd Nor and Assoc. Prof. Dr. Wang Seok Mui, for their professional assistance, insightful suggestions, and extensive discussions that significantly enriched my research process.

I am deeply appreciative of the Institute of Medical Molecular Biotechnology (IMMB), Faculty of Medicine, UiTM Sungai Buloh, for providing excellent research facilities. My gratitude also goes to all the staff of IMMB, particularly for their guidance and technical consultation throughout my study. A heartfelt thanks to my laboratory members for their shared thoughts, continuous support, and motivation, which have been a source of inspiration throughout this journey.

Finally, I wish to express my special appreciation to my beloved parents, as well as my siblings for their prayers, financial support, and unwavering motivation. Thank you for your patience, understanding, and for always standing by me, even during my most challenging times. Without your support, this research and thesis would not have been possible.

To everyone who has indirectly contributed to this research, your kindness and assistance mean the world to me. Thank you very much. Alhamdulillah.

TABLE OF CONTENTS

	Page
CONFIRMATION BY PANEL OF EXAMINERS	ii
AUTHOR'S DECLARATION	iii
ABSTRACT	iv
ACKNOWLEDGEMENT	v
TABLE OF CONTENTS	vi
LIST OF TABLES	x
LIST OF FIGURES	xi
LIST OF SYMBOLS	xiii
LIST OF ABBREVIATIONS	xiv
CHAPTER 1 INTRODUCTION	1
1.1 Research Background	1
1.2 Problem Statement	6
1.3 Research question	7
1.4 Hypothesis	8
1.5 Research Objectives	8
1.6 Significance of Study	8
1.7 Scope of Study	9
1.8 Justification of Study	10
1.8.1 Dengue Viruses Serotype 1 and 2	10
1.8.2 Cells Selection	10
CHAPTER 2 LITERATURE REVIEW	11
2.1 General Overview of Dengue Fever	11
2.2 Dengue through the Ages	11
2.3 Dengue Dynamics	13
2.4 Dengue and its Global Grip and Human Toll	15

CHAPTER 1

INTRODUCTION

1.1 Research Background

Dengue fever is one of the most prevalent mosquito-borne diseases caused by the dengue virus (DENV) and spread through the bite of female mosquitoes of the *Aedes* type, principally *Aedes aegypti* (Heilman et al., 2014). The infection produces symptoms including the sudden onset of high fever, nausea, vomiting, severe headache, muscle and joint pains, and a characteristic skin rash (Heilman et al., 2014). However, in some cases, it can lead to life-threatening complications such as dengue haemorrhagic fever and dengue shock syndrome (Alejandria, 2015; Tuiskunen Bäck & Lundkvist, 2013). Typically, it occurs in tropical and subtropical areas of the world especially in urban and semi-urban areas (Liu et al., 2021). The DENV is a single positive-stranded RNA virus of the family *Flaviviridae*; genus *Flavivirus* (Tuiskunen Bäck & Lundkvist, 2013). There are four serotypes of DENV (DENV1, DENV2, DENV3, and DENV4), classified based on their antigenic distinctions. Infection with one serotype usually provides lifelong, serotype-specific immunity, while offering only partial immunity to the other serotypes (Holmes & Burch, 2000). Subsequent infection with a different serotype increases the risk of severe complications (Holmes & Burch, 2000).

According to the Centre for Disease Control (CDC), almost 50% of the global population lives in areas at risk for DENV infection (CDC, 2024). According to the World Health Organization (WHO), the global incidence of dengue increases dramatically every year (WHO, 2024). About 67 to 136 million people are infected by dengue each year and approximately 10,000 to 20,000 cases are fatal (Bhatt et al., 2013). It is estimated that 3.9 billion people are at risk of DENV infection in 128 countries (Brady et al., 2012). This includes Africa, America, Eastern Mediterranean, Southeast Asia, and the Western Pacific region (Guo et al., 2017). Malaysia is not an exception with previous data showing an increased incidence of dengue cases in Malaysia since the first major outbreak in 1973 (Wallace et al., 1980). Figure 1.1