

DETECTION OF GENETIC VARIATION OF DNA GYRASE ENCODING FLUROQUINOLONE-RESISTANT ACINETOBACTER BAUMANNII

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Dissertation submitted in partial fulfilment of the requirements

for the degree of

Bachelor of Pharmacy (Hons.)

ACKNOWLEDGEMENT

First and foremost, I do declare my outmost gratitude to My Lord Allah, The All Mighty, The Most Merciful and The Most Graceful. Alhamdulillah, finally I managed to complete this thesis by the praise of Allah and His guidance and blessings. This project would not have been possible without the support of many people. I would like to express my deepest gratitude to my supervisor, Prof Dr. Teh Lay Kek, for her guidance, care, patience; and providing a conducive atmosphere for doing research. I would also like to thank Mr Mohamad Izwan Ismail for guiding my research since last semester and helping me especially in developing my skill and enhancing knowledge in biotechnology, biochemistry and genetics. Special thanks goes to iPROMISE UiTM members especially Prof Dr. Mohd Zaki Salleh, Director of iPROMISE for his guidance. Many thanks to Madam Nur Jalinna Abdul Jalil, Miss Noor Syakilla Azman, Miss Ismanura'in Ibrahim, and other post-graduate students in iPROMISE for helping me. My research would not have been possible without their helps. I would also like to thank my parents, and family members. They were always supporting me and encouraging me with their best wishes. Finally, I would like to thank my best friends, Zafirah Zul and Abdul Rahman Aiman Rosli. They were always there cheering me up and stood by me through the good times and bad.

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ABSTRACT

Mutations of the *gyrA* gene have been reported to contribute to *Acinetobacter baumannii* resistance towards quinolones. This study investigates the variation of DNA sequences of *gyrA* gene in 2 different isolates of *A.baumannii* which were the resistant and susceptible strains. The main aim of this project is to determine the genetic variation that encode resistance towards fluoroquinolones. In this study, primers that allow amplification of the region spanning *gyrA* was designed. The amplicons were amplified and subsequently purified before sequencing to determine the DNA sequences of the isolates. From bioinformatics analysis, there was a sequence change from cytosine to thymine in resistant strain of *A.baumannii*. The nucleotide change caused a change in amino acid codon from Serine to Leucine at position 21. Serine is hydrophilic in properties while Leucine are hydrophobic. Quinolones drugs having hydrophilic properties probably cannot bind to receptor due to hydrophobic changes. The mutation has probably caused the underlying resistance towards quinolones shown by the *A.baumannii*.

CHAPTER 1

INTRODUCTION

1.1 Acinetobacter baumannii

Nosocomial infections caused by *Acinetobacter baumannii* is reported to increase steadily (Vila et al., 1995; Vila et al, 2002). The control of *Acinetobacter baumannii* infections is challenging due to its versatile ability to develop resistance towards antimicrobial resistance (Cisneros & Rodriguez-Bano, 2002). Groups that are highly at risk to infection caused by *Acinetobacter baumannii* are immunocompromised patients, critically ill patients and patients that receive long-term treatment such as severe burns and traumatic injury (Buchan, Ledeboer, & Edmiston, 2011). Last few decades, the emergence of resistance towards multiple groups of antibiotics by *Acinetobacter baumannii* are due to broad use of antimicrobials within hospitals and also the chemical disinfectant used in sterilizing surgical environment (Adams-Haduch et al., 2008; Buchan et al., 2011; Shehata, 2012).

Groups of antibiotics that *Acinetobacter baumannii* are resistant to include β-lactams, carbapenems (e.g.,imipenem, meropenem and doripenem), aminoglycosides (e.g.,gentamicin, amikacin), ampicilin-sulbactam, extended-spectrum cephalosporin (e.g., ceftazidime) and quinolones (e.g., ciprofloxacin and levofloxacin) (Adams-Haduch et al., 2008; Hujer et al., 2009). When *Acinetobacter baumannii* are resistant to three or more groups of antimicrobial agents, it is classified as multidrug resistant (MDR) (Hujer et al., 2009).