## Species diversity of acetic acid bacteria at Khanom-Mu Ko Thale Tai National Park, Nakhon Si Thammarat Province

Pattaraporn Yukphan<sup>1,\*</sup>, Taweesak Malimas<sup>1</sup>, Winai Chaipitakchonlatarn<sup>1</sup>, Wanchern Potacharoen<sup>1</sup>, Somboon Tanasupawat<sup>2</sup> and Yuzo Yamada<sup>1</sup>

<sup>1</sup>National Center for Genetic Engineering and Biotechnology, Pathumthani, Thailand, \*e-mail: pattaraporn@biotec.or.th, <sup>2</sup>Chulalongkorn University, Bangkok, Thailand

During the course of study of the species diversity of acetic acid bacteria at Khanom-Mu Ko Thale Tai National Park. Nakhon Si Thammarat Province, one hundred and eighty-three strains were isolated from one hundred and seventy-nine collected samples. Seventy-nine strains were selected to analyze the 5' ends of 16S rDNA sequences for identification. In phylogenetic trees based on these sequences, seventyone strains had their identification confirmed as acetic acid bacteria. A phylogenetic tree based on the sequences showed that nineteen strains should be classified into two new genera in the family Acetobacteraceae. The other fifty-two strains were assigned to six known species of three genera, Acetobacter, Asaia and Gluconobacter, based on 16S rRNA gene sequences, namely A. tropicalis, As. bogorensis, As. lannensis, G. cerinus, G. frateurii and G. oxydans. However, additional phenotypic and genetic characteristics have to be further characterized before proposing the new taxa, especially DNA-DNA hybridization compared with the type strains of closely related taxa.

