

Genetic population differentiation of the blue swimming crab *Portunus pelagicus* (Portunidae) in Thai waters revealed by RAPD analysis

S. Klinbunga^{1,2}, V. Yuvanatemiya³, S. Wongphayak³, K. Khetpu⁴, P. Menasveta^{2,5} and B. Khamnamtong^{1,2}

¹Aquatic Molecular Genetics and Biotechnology Laboratory, National Center for Genetic Engineering and Biotechnology, National Science and Technology Development Agency, Klong 1, Klong Luang, Pathumthani, Thailand
²Center of Excellence for Marine Biotechnology, Faculty of Science, Chulalongkorn University, Bangkok, Thailand
³Faculty of Marine Technology, Burapha University, Chanthaburi Campus, Chanthaburi, Thailand
⁴Program in Biotechnology, Faculty of Science, Chulalongkorn University, Bangkok, Thailand
⁵Department of Marine Science, Faculty of Science, Chulalongkorn University, Bangkok, Thailand

Corresponding author: B. Khamnamtong E-mail: bavornlak@biotec.or.th

Genet. Mol. Res. 9 (3): 1615-1624 (2010) Received May 14, 2010 Accepted June 12, 2010 Published August 17, 2010 DOI 10.4238/vol9-3gmr886

ABSTRACT. Genetic diversity and population differentiation of the blue swimming crab, *Portunus pelagicus*, in Thailand were analyzed by RAPD analysis. One hundred and twelve RAPD fragments were generated from 109 individuals of *P. pelagicus* using OPA02, OPA14, OPB10, UBC122, and UBC158 primers. The percentage of polymorphic bands in each geographic sample and that of each primer across overall samples were 72.7-85.0 and 92.0-100%, respectively. Large numbers of polymorphic bands found in the RAPD analysis suggested high genetic diversity of Thai *P. pelagicus*. The mean genetic distance between

samples across all primers was 0.0929-0.2471. Significant geographic heterogeneity was observed across samples overall and between all pairs of geographic samples (P < 0.01 for θ and P < 0.0001 for the exact test), indicating strong genetic differentiation of *P. pelagicus* in Thai waters, despite its high potential of dispersal. Limited gene flow levels (0.44-1.19 individuals per generation) of Thai *P. pelagicus* were also observed. A fine scale level of differentiation suggested that *P. pelagicus* from each geographic sample in Thai waters should be regarded as a separate genetic population and treated as a different exploited stock.

Key words: Genetic diversity; Population structure; RAPD; *Portunus pelagicus*; Species-specific markers

Genetics and Molecular Research 9 (3): 1615-1624 (2010)