Distribution of major and potential malaria vectors in Mannar and Trincomalee Districts and systematics of anophelines in Sri Lanka

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P.A.D.H.N. Gunathilaka B.Sc (Kelaniya)

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Molecular Medicine Unit,
Faculty of Medicine,
University of Kelaniya,
Sri Lanka.

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VII. ABSTRACT

Background: Malaria was a major public health problem in Sri Lanka until recent past. Entomological surveillance is used to evaluate control programmes and facilitate interventions.

Objectives: To study on spatial, temporal, ecological, host preferences, and breeding habitats of major and potential malaria vector mosquitoes in selected sentinel sites in the Districts of Mannar (n= 3) and Trincomalee (n= 5) and to revise existing morphological keys of Sri Lankan anopheline mosquitoes.

Methodology: Densities of *Anopheles* were surveyed for a period of 25 months (June, 2010 to June, 2012) using World Health Organization recommended entomological techniques. Quality of water in breeding habitats was tested for abiotic variables. A multiplex, real-time Polymerase Chain Reaction (PCR) assay was developed to detect the host preference of *Anopheles*. Morphological identification keys for Sri Lankan anophelines were revised.

Results: A total of 161,891 anophelines representing 17 species were recorded from the Districts of Mannar (n=74,181) and Trincomalee (n=87,710). The most abundant species was *Anopheles subpictus* (n=91,370). *An. culicifacies* was recorded only in Trincomalee (1.34%). *An. culicifacies* and *An. subpictus* have adapted to breed in a wide range of water bodies including polluted water in urban settings with low dissolved oxygen (<3 mg/l) and high salinity ($21,105 \pm 1,344$ mg/l). Host preference

revealed that all anophelines preferred bovine as the host than humans. Anopheline

morphological identification keys for both larvae and adults were revised.

Discussion: The adaptation of An. culicifacies and An. subpictus to breed in polluted

water may lead the emergence of urban malaria in Sri Lanka, a phenomenon that has

not been reported on a regular basis as yet. Therefore, continuous monitoring of the

breeding of An. culicifacies in polluted water and to what extend it impacts on

malaria elimination programmes is to be established. The presence of human blood,

in some anopheline species indicates the possibility of transmitting human malaria.

The revised morphological identification keys for anophelines can be used as a rapid

and convenient guide to identify malaria vectors.

Conclusion: This investigation adds new information to the knowledge on malaria

entomology. Therefore, this study would be beneficial to implement new vector

control approaches.

Keywords: Entomological, programme, malaria, mosquitoes, Anopheles.

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