Molecular characterisation and disease severity of leptospirosis in Sri Lanka

Kanchana Kumari Bandara^{1,2}, Manjula Weerasekera¹/+, Chinthika P Gunasekara¹, Nilantha Ranasinghe³, Chamil Marasinghe⁴, Neluka Fernando¹

¹Department of Microbiology ⁴Department of Medicine, Faculty of Medical Sciences, University of Sri Jayewardenepura, Sri Lanka ²Department of Basic Sciences, Faculty of Allied Health Sciences, General Sir John Kotelawala Defense University, Sri Lanka ³Base Hospital, Tangalle, Sri Lanka

Leptospirosis is a re-emerging zoonotic disease all over the world, important in tropical and subtropical areas. A majority of leptospirosis infected patients present as subclinical or mild disease while 5-10% may develop severe infection requiring hospitalisation and critical care. It is possible that several factors, such as the infecting serovar, level of leptospiraemia, host genetic factors and host immune response, may be important in predisposition towards severe disease. Different Leptospira strains circulate in different geographical regions contributing to variable disease severtly. Therefore, it is important to investigate the circulating strains at geographical locations during each outbreak for epidemiological studies and to support the clinical management of the patients. In this study immunochromatography, microscopic agglutination test and polymerase chain reaction were used to diagnose leptospirosis. Further restriction fragment length polymorphism and DNA sequencing methods were used to identify the circulating strains in two selected geographical regions of Sri Lanka. Leptospira interrogans, Leptospira borgpetersenii and Leptospira kirschneri atrains were identified to be circulating in western and southern provinces. L. interrogans was the predominant species eirculating in western and southern provinces in 2013 and its presence was mainly associated with renal failure.