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Isolation and characterization of Leptospira interrogans from clinically suspected leptospirosis patients from selected hospitals in Western Province, OP 2

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Background: Culture isolation of human pathogenic Leptospires is a challenge. Isolation of Leptospires is confirmatory of the disease but is difficult and laborious to perform. However, culture has an epidemiological value mainly to identify responsible reservoir animals. The last report of isolation of Leptospires from an infected patient in Sri Lanka was published in 1973.

Objectives: To isolate and identify circulating Leptospira species from clinically suspected

Methods: Whole blood from 100 leptospirosis suspected patients were subjected to DNA extraction followed by secY PCR. Acute leptospirosis was presumptively diagnosed by detecting IgM using an immunochromatographic assay. Few drops of whole blood were inoculated into semi-solid Ellinghausen-McCullough-Johnson-Harris medium for culture and incubated at 30°C. Positive cultures were selected by presence of motile spiral bacteria under dark field microscopy and subjected to nested PCR targeting the flaB gene of pathogenic Leptospira and sequencing for phylogenetic analysis. Culture positive patients sera were subjected to microscopic agglutination test (MAT) to identify the infective Leptospira serogroups using a panel of 10 representative leptospiral serogroups.

Results: Among 100 patients, secY PCR found to be positive in 27(27%), while 26 patients gave positive IgM results. Only 2 patients were found to be culture positive (2%)using dark field microscopy. PCR of both cultures were positive for flaB gene from pathogenic Leptospires. The phylogenetic distance of flaB sequences suggested Leptospira interrogans while MAT identified infective serogroups

Conclusion: For early detection of pathogenic Leptospires, molecular based assays are more suitable as Sejroe and Canicola. than culture isolation. Serogroup Sejroe and Canicola of Linterrogans have more affinity to animal origin such as cattle and dogs respectively. Thus, in the current study we stress the importance of cross contamination of infected animals in human habitats.

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