# Molecular Survey of Babesia gibsoni and Hepatozoon canis Infection in Dogs from Jeju Island 

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Purpose: Babesia gibsoni and Hepatozoon canis are a tick-borne protozoan that infected dogs and have been reported throughout the world. Manifestration of their infection varies from being sub-clinical in apparently healthy dogs to severe illness(body weight loss, anemia, lethargy and death). In order to survey the infection of babesia and hepatozoon spp., we have screened 184 dogs in Jeju island.

Materials and Method: Blood samples were collected from 184 dogs in Jeju island. We have examined the CBC, Giemsa staining and DNA analysis. We extracted the genomic DNA from blood using the genomic DNA extraction $\operatorname{kit}(G-D E X \| b, i N t R O N ~ B i o, ~ K o r e a) . ~ F i v e ~ p a i r s ~$ of primer sets were designed to amplify 18 s rRNA DNA fragments of babesia and hepatozoon spp.. And nested PCR, PCR-RFLP and PCR product sequencing analysis were performed for the detection and differentiation of the them.

Results: Of the 184 dogs, $14(7.6 \%)$ were B. gibsoni, $16(8.7 \%)$ were H. canis, and 6 dogs( $3.3 \%$ ) were co-infection.

Conclusion: There have been few clinical case reports for the Hepatozoon canis in Korea. More epidemiological data obtained using molecular methods is needed to understand canine Hepatozoon spp..

Key words: Babesia gibsoni, Hepatozoon canis, Dog, Jeju

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