

Toxoplasma gondii in Slaughtered Sheep in High- and Low-Humidity Regions in the South of Iran: Molecular Prevalence and Genotype Identification

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Toxoplasma gondii is one of the most common meat-borne zoonoses that infect all warm-blooded animals and humans. Sheep (*Ovis aries*) is one of the main reservoirs of *T. gondii* worldwide, and the infections induce various sequels, such as abortion and stillbirth. The present study aimed to identify the effects of humidity on the prevalence of *T. gondii* in sheep in high- and low-humidity regions. Heart samples from 200 slaughtered sheep (140 samples from a high-humidity region and 60 samples from a low-humidity region) were collected from Hormozgan Province (south of Iran). The samples were tested by nested PCR targeting the RE gene. Genotyping was performed by the PCR-RFLP method using the SAG3 and GRA6 genes. Some isolates were sequenced and recorded in the GenBank. *T. gondii* DNA was detected in 10.71 percent of the samples from the highly humid region, whereas no positive samples were detected in the low-humidity region. Genotyping revealed that all isolates belonged to the *T. gondii* type III genotype. Our study indicated that humidity is an important factor for the prevalence of *T. gondii* in sheep. Additionally, our study also showed the dominance of type III strain of *T. gondii* in sheep in the south of Iran.

Keywords: *Toxoplasma gondii*, Genotype Identification, Slaughtered Sheep