



Comment on “Molecular Characterization of Animal *Fasciola* spp. Isolates from Kermanshah, Western Iran”

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Dear Editor-in-Chief

In the previous work by Bozorgomid et al. (1), helpful information regarding the molecular epidemiology of fascioliasis in Kermanshah was demonstrated by characterizing the population genetic variability of *Fasciola* species. Although we think this study is very informative to researchers interested in the molecular epidemiology, the study contains Materials and Methods which are disputable.

The verification of the sequences of *Fasciola* species obtained in Bozorgomid et al. (1) has not been included in Materials and Methods section. Thus, the results of chimera detection should be provided as a further verification step since the sequences with undetected chimeras may be misinterpreted as novel species (2, 3). If chimeric DNA sequences are observed, they should be discarded for the further data analysis.

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The authors declare that there is no conflict of interests.

References

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