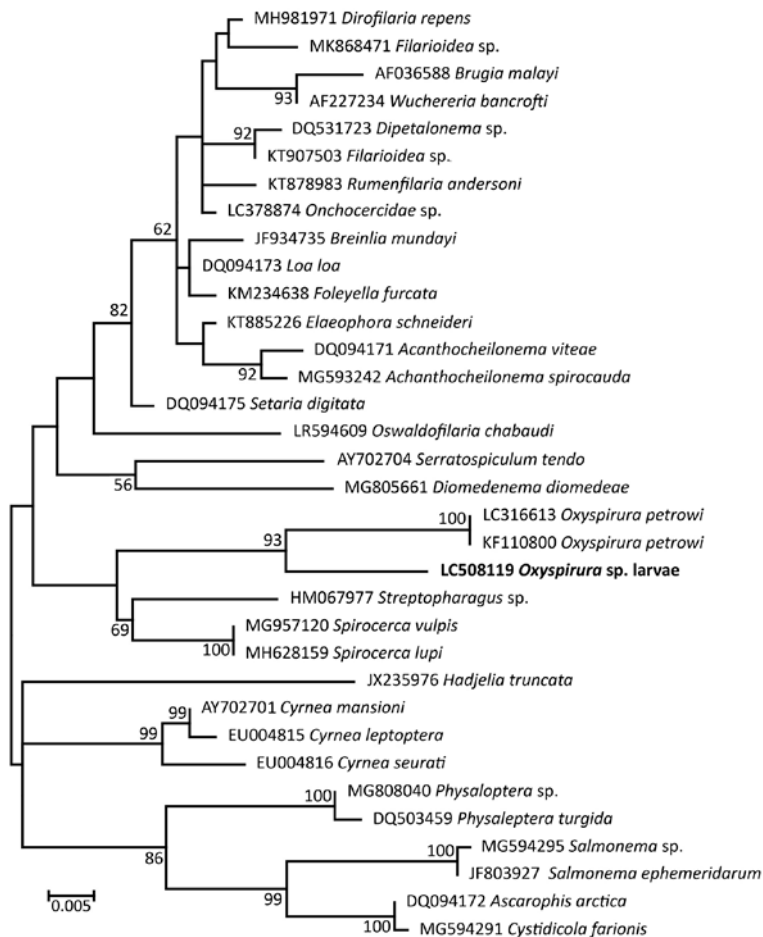


Pruritic Cutaneous Nematodiasis Caused by Avian Eyeworm *Oxyspirura* Larvae, Vietnam

Appendix



Appendix Figure. Molecular phylogenetic tree reconstructed from 18S rDNA sequences of *Oxyspirura* larvae from skin lesions of a patient, Vietnam. The evolutionary history was inferred by using the maximum likelihood method based on the Kimura 2-parameter model. A discrete Gamma distribution was used to model evolutionary rate differences among sites (+G). The rate variation model allowed some sites to be evolutionarily invariable (+I). All positions containing gaps and missing data were eliminated. Bootstrap values >50% are shown above branches. Bold text indicates nucleotide sequences obtained in this study. Scale bar indicates nucleotide substitutions per site.