## Human Babesiosis Caused by *Babesia odocoilei*: A Confirmed Zoonosis

## John Donald Scott

Tick Research, Upper Grand Tick Research Centre, Canada

Article history Received: 13-11-2024 Revised: 20-11-2024 Accepted: 23-11-2024

Email: jkscott@bserv.com

Abstract: Babesia odocoilei has been gaining attention as a novel agent of tick-borne disease. Researchers allege that the American dog tick is a vector of *B. odocoilei*, but supportive evidence for this contention is absent. Likewise, *B. odocoilei* was allegedly found in two European patients, but supportive evidence for this postulate is lacking and it appears that the patients were infected with a close member of a sister group, *Babesia venatorum*. In contrast, a study by Scott *et al.* in two Canadian patients provides molecular evidence for the first-ever confirmed cases of human babesiosis caused by *B. odocoilei* anywhere in the world.

Keywords: Babesia odocoilei, Babesiosis, Ticks, Canada, Ixodes

Maggi *et al.* (2024) claim that *Babesia microti* is the predominant Babesia species in central and eastern North America, but this assumption is incorrect. Creditable scientific findings show that the intraerythrocytic piroplasmid *Babesia odocoilei* is the predominant *Babesia* species in Canada. Tick researchers found that the ratio of *B. odocoilei* to *B. microti* is 60-1 (Scott *et al.*, 2024) and *B. odocoilei* is the predominant *Babesia* species in *Ixodes scapularis* ticks nationwide. Maggi *et al.* (2024) state that *B. odocoilei* has been reported in Canada but incorrectly cited Herwaldt *et al.* (2003). In reality, Scott *et al.*, discovered the *B. odocoilei* archetype in Canadian ticks (Scott *et al.*, 2022).

Although Maggi *et al.* (2024) claim that the American dog tick, *Dermacentor variabilis*, is a primary vector of *B. odocoilei*, none of the references cited in their study substantiate *D. variabilis* as a vector of *B. odocoilei*. This single-celled, red blood cell parasite causes malaria-like, febrile symptoms in humans who have been bitten by *B. odocoilei*-infected *Ixodes* ticks. After a comprehensive review of the scientific literature, there are no peer-reviewed references that show transstadial passage (larva to nymph and/or nymph to adult) of *B. odocoilei* in *D. variabilis*. In reality, it appears that the western black-legged tick, *Ixodes pacificus* and the black-legged tick, *Ixodes scapularis*, are the primary vectors of *B. odocoilei* in North America.

Maggi *et al.* (2024) intimated that two male patients in Austria and Italy were the first patients to be infected with *B. odocoilei.* However, the babesial strains that were recovered from the two patients clustered with a related sister *Babesia* group, EU-1 (*Babesia venatorum*), and the amplicons from the two patients failed to meet the current molecular criteria for *B. odocoilei*. Maggi *et al.* (2024) mistakenly alluded to the supposition that *B. odocoilei* amplicons from Europe mimicked those from people in Canada and the United States, but the researchers mistakenly used a reference out of context and netted a miscitation. The archetypical reference for human infection with *B. odocoilei* is Scott *et al.* (2021).

Based on molecular characterization and patient symptomology, Scott *et al.* (2021) are the first research team to confirm that *B. odocoilei* causes human babesiosis. *Babesia* amplicons from these Canadian patients had 99.77 and 99.55% similarity with *B. odocoilei* type strains in GenBank. Phylogenetically, the amplicons from the two Canadian study participants are well within the accepted molecular range of a valid *B. odocoilei* species. These convincing findings verify that Scott *et al.* (2021) have described the first-ever confirmed cases of human babesiosis caused by *B. odocoilei*.

## Conflicts of Interest

The author has no competing interests.

## References

Herwaldt, B. L., Cacciò, S., Gherlinzoni, F., Aspöck, H., Slemenda, S. B., Piccaluga, P., Martinelli, G., Edelhofer, R., Hollenstein, U., Poletti, G., Pampiglione, S., Löschenberger, K., Tura, S., & Pieniazek, N. J. (2003). Molecular Characterization of a Non-*Babesia divergens* Organism Causing Zoonotic Babesiosis in Europe. *Emerging Infectious Diseases*, 9(8), 942–955. https://doi.org/10.3201/eid0908.020748



- Maggi, R. G., Calchi, A. C., Moore, C. O., Kingston, E., & Breitschwerdt, E. B. (2024). Human *Babesia* odocoilei and Bartonella spp. co-infections in the Americas. *Parasites & Vectors*, 17(1), 302. https://doi.org/10.1186/s13071-024-06385-4
- Scott, J. D., Sajid, M. S., Pascoe, E. L., & Foley, J. E. (2021). Detection of *Babesia odocoilei* in Humans with Babesiosis Symptoms. *Diagnostics*, *11*(6), 947.

https://doi.org/10.3390/diagnostics11060947

- Scott, J. D., McGoey, E., & Pesapane, R. R. (2022). Tick-Borne Pathogens Anaplasma phagocytophilum, Babesia odocoilei, and Borrelia burgdorferi Sensu Lato in Blacklegged Ticks Widespread across Eastern Canada. J Biomed Res Environ Sci, 3(10), 1249–1256. https://doi.org/10.37871/jbres1586
- Scott J. D., & Scott C. M. (2024) Human Babesiosis Caused by *Babesia odocoilei*: An Emerging Zoonosis. *Am J Biomed Sci & Res.* 24(6): 667-668. https://doi.org/10.34297/AJBSR.2024.24.003261