Evaluation of Amblyomma americanum vector competence for Anaplasma marginale

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Bovine anaplasmosis is a tick-transmitted, production-limiting disease and a major obstacle to profitable beef cattle production in the United States. Changes in climate, ecosystems, and increases in animal transport have contributed to the expansion of various disease-transmitting tick species, including the Lone Star Tick (LST) (Ambylomma americanum), a species now commonly found on cattle. The intracellular rickettsial pathogen and agent of anaplasmosis, Anaplasma marginale (Am), is primarily transmitted by Dermacentor tick species in the U.S. The role of LST in the transmission of bovine anaplasmosis is currently unknown; however, the frequency of LST infestation on cattle warrants examination into whether LST contributes to Am transmission. The objective of this study was to examine the vector competence of LST for Am using a combination of field-surveillance and controlled laboratory transmission experiments. For the field-surveillance component, host-seeking LST were collected from a pasture used by a cattle herd naturally-endemic for Am and tested for Am using real-time PCR. The vector competence of LST for Am was specifically evaluated by comparing the ability of two geographically-distinct LST strains to acquire and transmit Am compared to a known Am vector, Dermacentor variabilis, using an experimental tick-calf Am transmission model. The Am bacterial levels were monitored throughout the transmission process in calf blood and tick midgut and salivary gland tissues to specifically tract Am. The results will provide a combination of field and laboratory evidence to illuminate the potential LST contribution to Am transmission in the U.S., information which will influence disease management strategies.

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