Abstract

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Molecular and morphological identification of *Brevipalpus* spp. (Acari: Tenuipalpidae) in *Citrus sinensis*, Costa Rica*

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The genus *Brevipalpus* presents interspecific variability since its description. Therefore, the question arises as to which species are found on citrus in Costa Rica. Molecular and morphological techniques are complementary tools that allow a more precise identification at the species level, which is important, some species of the genus Brevipalpus have been associated with virus transmission, such as *Citrus leprosis virus*.

In this research, 16 orange (*Citrus sinensis* L.) farms were sampled, distributed in the Central, Northern Huetar, and Brunca Region. The species *Brevipalpus yothersi*, *B. azores*, *B. papayensis* were identified by morphological characters and a new species of this genus (*Brevipalpus* sp. nov.) was proposed, which will be named and described in a forthcoming paper. Partial sequencing of the COX 1 gene identified the species *B. yothersi*, *B. azores*, and *Brevipalpus* sp. nov. *Brevipalpus yothersi* was found in the Central and Brunca Regions, while *B. azores*, *B. papayensis* and *Brevipalpus* sp. nov. were found only in the Brunca Region.

Keywords: Molecular identification, flat mites, citrus, COX1, populations diversity