
2022

[**The Haptomonad Stage of *Critidia acanthocephali* in *Apis mellifera* Hindgut**](#)

2021

[**Bee Trypanosomatids: First Steps in the Analysis of the Genetic Variation and Population Structure of *Lotmaria passim*, *Critidia bombi* and *Critidia mellifica***](#)

[**Detection of Microsporidia in Pollinator Communities of a Mediterranean Biodiversity Hotspot for Wild Bees**](#)

[**Residual Tau-Fluvalinate in Honey Bee Colonies Is Coupled with Evidence for Selection for *Varroa destructor* Resistance to Pyrethroids**](#)

[**First description of *Lotmaria passim* and *Critidia mellifica* haptomonad stages in the honeybee hindgut.**](#)

[**A Case Report of Chronic Stress in Honey Bee Colonies Induced by Pathogens and Acaricide Residues.**](#)

[**A SNP assay for assessing diversity in immune genes in the honey bee \(*Apis mellifera L.*\)**](#)

[**Sporicidal evaluation of different substances against *Nosema ceranae* for surface disinfection**](#)

[**An Optimized Extraction Procedure for Determining Acaricide Residues in Foundation Sheets of Beeswax by Using Gas Chromatography-Mass Spectrometry**](#)

Dataset of the next-generation sequencing of variable 16S rRNA from bacteria and ITS2 regions from fungi and plants derived from honeybees kept under anthropogenic landscapes.

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