

Short note [Nota corta]



First report of the pepper thrips *Thrips parvispinus* (Karny) in Costa Rica†

[Primer reporte del trips del pimiento *Thrips parvispinus* (Karny) en Costa Rica]

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SUMMARY

Background: The accidental invasive insects pose a growing threat to agricultural productivity and food security worldwide. **Objective:** To report for the first time the pepper thrips *Thrips parvispinus* in Costa Rica. **Methodology:** Specimens were collected from May to October 2024 in commercial crops *Capsicum annuum*, *Citrus sinensis*, *Coriandrum sativum*, *Cucumis sativus*, and ornamental *Croton* spp. in the western region of the country, encompassing both protected environments and greenhouses. The thrips identification was carried out using external morphological characteristics and partial sequencing of the mitochondrial cytochrome c oxidase subunit I (COI) gene. **Results:** Morphological and molecular analyses confirmed the species identity of *T. parvispinus*. **Implications:** The observed damage included feeding scars on young leaves, margin curling, necrosis of foliar tissue, and floral deformation; with relevant implications for the health of these crops in Costa Rica. **Conclusion:** The detection of this invasive species represents a phytosanitary concern in Central America and underscores the necessity for continuous monitoring and implementation of effective insect management strategies throughout the region.

Key words: invasive species; crop thrips; taxonomic identification; plant health; exotic species.

RESUMEN

Antecedentes: La introducción de especies de insectos invasivas conlleva una constante preocupación en la agricultura y la seguridad alimentaria mundial. **Objetivo:** Reportar por primera vez la presencia de el trips del pimiento *Thrips parvispinus* en Costa Rica. **Metodología:** Los muestreos de especímenes se realizaron entre mayo y octubre de 2024 en cultivos comerciales de *Capsicum annuum*, *Citrus sinensis*, *Coriandrum sativum*, *Cucumis sativus* y *Croton* spp., incluyendo ambientes protegidos e invernaderos. La identificación de los especímenes se llevó a cabo mediante análisis de morfología externa, complementados con secuenciación parcial del gen mitocondrial COI. **Resultados:** Los análisis morfológicos y moleculares confirmaron la identidad de la especie *Thrips parvispinus*. **Implicaciones:** Se documentaron daños típicos de alimentación de los insectos hacia hojas jóvenes, enrollamiento de márgenes, necrosis en láminas foliares y afectación floral, lo que podría tener consecuencias para la sanidad vegetal

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de dichos cultivos en Costa Rica. **Conclusión:** Este hallazgo constituye una alerta fitosanitaria para la región centroamericana, dada la capacidad invasiva de esta especie. Su presencia refuerza la necesidad de monitoreo constante y de correctas medidas fitosanitarias en los países de América.

Palabras clave: especies invasoras; trips en cultivos; identificación taxonómica; fitosanidad; especie exótica.

INTRODUCTION

Insects of the order Thysanoptera, commonly referred to as thrips, constitute a significant group due to their impact on a wide array of crops, either through direct physical damage or by transmitting viruses that adversely affect agricultural yields (Lewis, 1997; Maris *et al.*, 2003; Moudén *et al.*, 2017). Species in the genus *Thrips* are notable for their remarkable diversity and frequent occurrences in global phytosanitary records (Bragard *et al.*, 2019; Díaz-Montano *et al.*, 2011; He *et al.*, 2020). This genus is present in all continents except Antarctica, with the highest incidence in places located between 30° N and 60° N. However, their presence diminishes at higher elevations and in environments with limited floral availability (Gentile and Bailey, 1968). Generally, *Thrips* species are phytophagous (Nakahara, 1994), typically measuring no more than 5 mm in length (Hulagappa *et al.*, 2022; Hutasoit *et al.*, 2019), and exhibiting cryptic behavior (Morse and Hoddle, 2006), which has facilitated their global dispersal (Lewis, 1997).

Research on the genus *Thrips* in the Americas is limited, with most distributional data originating from the northern part of the continent (Nakahara, 1994). However, this information is less comprehensive than that available from Europe, as many species reported in the Americas have Holarctic European origins (Monteiro *et al.*, 2001; Mound and Marullo, 1996). Nakahara (1994) documented 62 *Thrips* species in North America, 43 of which are endemic. In contrast, only a few species are known from the Neotropical region, most of which have been introduced (Mound and Marullo, 1996). Three species have been identified in Argentina (De Borbón, 2008), four in Brazil (Cavalleri and Gonçalves, 2022; Monteiro *et al.*, 2001), and several in Colombia (Ebratt-Ravelo *et al.*, 2024), whereas nine species have been recorded in Central America (Goldarazena *et al.*, 2012; Mound and Marullo, 1996).

Globally, several *Thrips* species are recognized as significant agricultural pests, including *Thrips tabaci*, *Thrips palmi*, and *Thrips parvispinus* (Bragard *et al.*, 2019; Díaz-Montano *et al.*, 2011; Hulagappa *et al.*, 2022). *Thrips parvispinus* was first collected and described from the flowers of *Trichosanthes tricuspidata* in Thailand and Vietnam (Karny, 1922). This polyphagous species measures 0.8 to 1.2 mm in length, with adults typically exhibiting yellow to light brown coloration and darker-tinted wings (Pavithran

et al., 2024). Native to Southeast Asia, *T. parvispinus* has emerged as a serious pest in India and Indonesia, particularly affecting *Capsicum annuum* (sweet pepper or chili) crops (Gleason *et al.*, 2023; Hulagappa *et al.*, 2022; Tyagi *et al.*, 2015). In Europe, it was first detected in Greece in 1998 (Mound and Collins, 2000) and was recorded in northern Australia in 2005 (Mound and Masumoto, 2005). Additionally, it was reported in Spain in 2017 (Pijnakker, 2023), France in 2018 (European Food Safety Authority, 2019), the Netherlands in 2019 (National Plant Protection Organization of the Netherlands, 2019), Germany in 2021, and Belgium in 2022 (Pijnakker, 2023). In the Americas, *T. parvispinus* was first detected in Barbados (Seal *et al.*, 2023); however, in continental North America, it was first reported in Florida in 2020, causing damage to ornamental plants (Soto-Adames, 2020), then it was confirmed in Canada in 2023 (Gleason *et al.*, 2023) and recently, was recorded in Sinaloa, México (Payán-Arzapalo, 2025).

In recent years, the arrival of invasive thrips species in Costa Rica, such as *Megalurothrips usitatus*, has been documented (Rodríguez-Arrieta *et al.*, 2023). Prior to this report, there were no records of *T. parvispinus* in Costa Rica. This study is the first to document the occurrence of this species in association with commercially important crops in this country.

MATERIAL AND METHODS

Sampling

Thrips specimens were collected from various locations in Alajuela province: Sarchí, San Pedro (10°06'41.7"N, 84°19'58.8"W; May 2nd, 2024) in a protected *Capsicum annuum* (chili pepper) cultivation environment; San Carlos, Muelle (10°27'55.6"N, 84°27'38.8"W; August 20, 2024) in a greenhouse with *Citrus sinensis* (orange); Zarcero, Brisas (10°13'45.7"N, 84°23'44.4"W; September 1st, 2024) in a greenhouse with *C. annuum*; San Ramón, Bajo Rodríguez (10°18'25.1"N 84°31'55.4"W; September 26, 2024) in an open field with *Croton* sp. and Bajo Zúñiga (10°08'20.3"N, 84°29'53.1"W; September 9, 2024) in an open field with *C. annuum*, *Cucumis sativus* (cucumber) and *Coriandrum sativum* (coriander); Palmares (10°03'38.6"N, 84°26'17.7"W; October 9, 2024) in a greenhouse with *C. annuum*. The elevation at the collection sites ranged from 200 m a.s.l. in San Carlos, Muelle, to 1,000–1,500 m a.s.l. in Sarchí, Zarcero, Palmares, and San Ramón. ArcGIS Maps (Esri, 2024) were used to illustrate these

locations (Figure 1). A total of 318 adult and both sexes specimens were sampled and collected directly from the host plants (53 specimens per location) that were carried out to the laboratory stored in 90% of ethylic alcohol in glass test tubes with cap.

Light Microscopy

Thrips specimens were identified at the Centro de Investigación en Estructuras Microscópicas (CIEMic), Universidad de Costa Rica (UCR), using conventional light microscopy with a Zeiss® Axiolab microscope at magnifications ranging from 5X to 63X. Specimens were prepared and cleared following the standard mounting techniques described by Mound and Marullo (1996), and taxonomic identification using the morphological keys provided by Mound and Masumoto (2005). Post-processing of the images was performed with HeliconFocus version 8.2.6 (HeliconSoft, 2022) with depth map stacking, and final composition of the figures was completed in Photoshop version 25 (Adobe Inc., 2024). Voucher specimens were deposited in the Thysanoptera Collection at CIEMic-UCR.

DNA Extraction, Sequencing, and Genetic Inference

Genomic DNA was extracted using the DNeasy Blood and Tissue Kit (QIAGEN®) following the manufacturer's instructions. A partial fragment of the mitochondrial cytochrome c oxidase subunit I (COI)

gene was amplified using the primers LCO1490 (5'-GGTCAACAAATCATAAAGATATTGG-3') and HCO2198 (5'-TAAACTTCAGGGTGACCAAAAAATCA-3') (Folmer *et al.*, 1994). PCR reactions were carried out in 25 µl volumes containing 3–50 ng of DNA (1–2 µl), 0.25 µl of Taq DNA polymerase (DreamTaq, Thermo Scientific, 5 U/µl), 2.5 µl of DreamTaq Buffer (10X), 2.5 µl of dNTP mix (10 mM each), and 1.25 µl of each primer (10 µM). The thermal cycling profile consisted of an initial denaturation at 95°C for 1 min, followed by 35 cycles of denaturation at 95°C for 1 min, annealing at 58°C for 1 min, extension at 60°C for 1 min, and a final extension at 72°C for 10 min. The PCR products were purified and sequenced by MacroGen, Inc. (Seoul, South Korea) using the same primers used for amplification. Complementary strands were assembled and edited using BioEdit v7.2.5. Sequence alignment was performed using MUSCLE in MEGA v.11.0.13 (Tamura *et al.*, 2021) with sequences of 570 bp. Phylogenetic analysis was conducted using the Maximum Likelihood (ML) method within the same software. Branch support was assessed through bootstrap analysis with 1,000 replicates, and the Kimura 2-parameter (K2P) nucleotide substitution model was applied. Sequences from specimens collected in Costa Rica were deposited in GenBank (NCBI, National Center for Biotechnology Information) under accession numbers PQ624786, PQ624787, PQ624788, PQ624789, PQ624790, and PV335501.



Figure 1. Locations reporting the first detection of *Thrips parvispinus* in Costa Rica, Central America.

RESULTS AND DISCUSSION

At all locations, many of the specimens collected were female *Thrips parvispinus* (282 females and 36 males). The body coloration of these specimens was light brown on the head and thorax, in contrast to the dark brown on the abdomen (Figure 2A). This coloration exhibits variability in tone across different body sections; specifically, segments III and the basal halves of segments IV–V of the antennae are lighter in color (Figure 2B). Like other members of the genus *Thrips*, this species is characterized by seven-segmented antennae with bifurcated sensoria on

segments III and IV, absence of ocellar seta I (Figure 2C), and two pairs of long posteroangular setae on the prothorax (Figure 2D). Abdominal sternites III–VI possess ten discal setae, whereas sternite VII lacks these setae (Figure 2E). The forewings are shaded in the distal three-quarters and remain clear only at the base. Morphologically, *T. parvispinus* resembles *T. orientalis*, particularly because of the complete row of setae along the first forewing vein (Figure 2F), a feature that is uncommon among other species in this genus. In contrast, *T. orientalis* exhibits uniformly light brown forewings and possesses fewer discal setae (seven or fewer) on abdominal sternites III–VI.



Figure 2. External morphology of the species *Thrips parvispinus*. (A) Macropterous female; (B) seven-segmented antenna; (C) head lacking ocellar setae I; (D) prothorax with long posteroangular setae; (E) abdominal sternites V–VII; (F) forewing with a complete row of setae along the first vein.

Visual damage was observed in young host plant leaves, where leaf margins exhibited curling and deformation or even senescence of new shoots (Figs. 3A, 3B). This thrips was found on both sides of the leaf (Figure 3C) and throughout the plant; however, damage was most evident on the abaxial leaf, where discoloration, necrotic spots, and epidermal destruction were observed (Figs. 3D, 3E). *T. parvispinus* was also observed in the plant flowers (Figure 3F).

Phylogenetic analysis confirmed the morphological identification and validated the species identity

through COI gene sequences compared to the NCBI database (Figure 4). The resulting phylogenetic tree revealed the formation of two clades, demonstrating a clear separation between *T. parvispinus* and other *Thrips* species such as *T. orientalis* and *T. palmi*. Additionally, the subdivision within the *T. parvispinus* clade indicates a degree of geographic structuring among haplotypes, with genetic similarities observed in sequences from India and Indonesia, whereas sequences from Costa Rica and the United States appeared to share a common haplotype (Ahmed *et al.*, 2024).

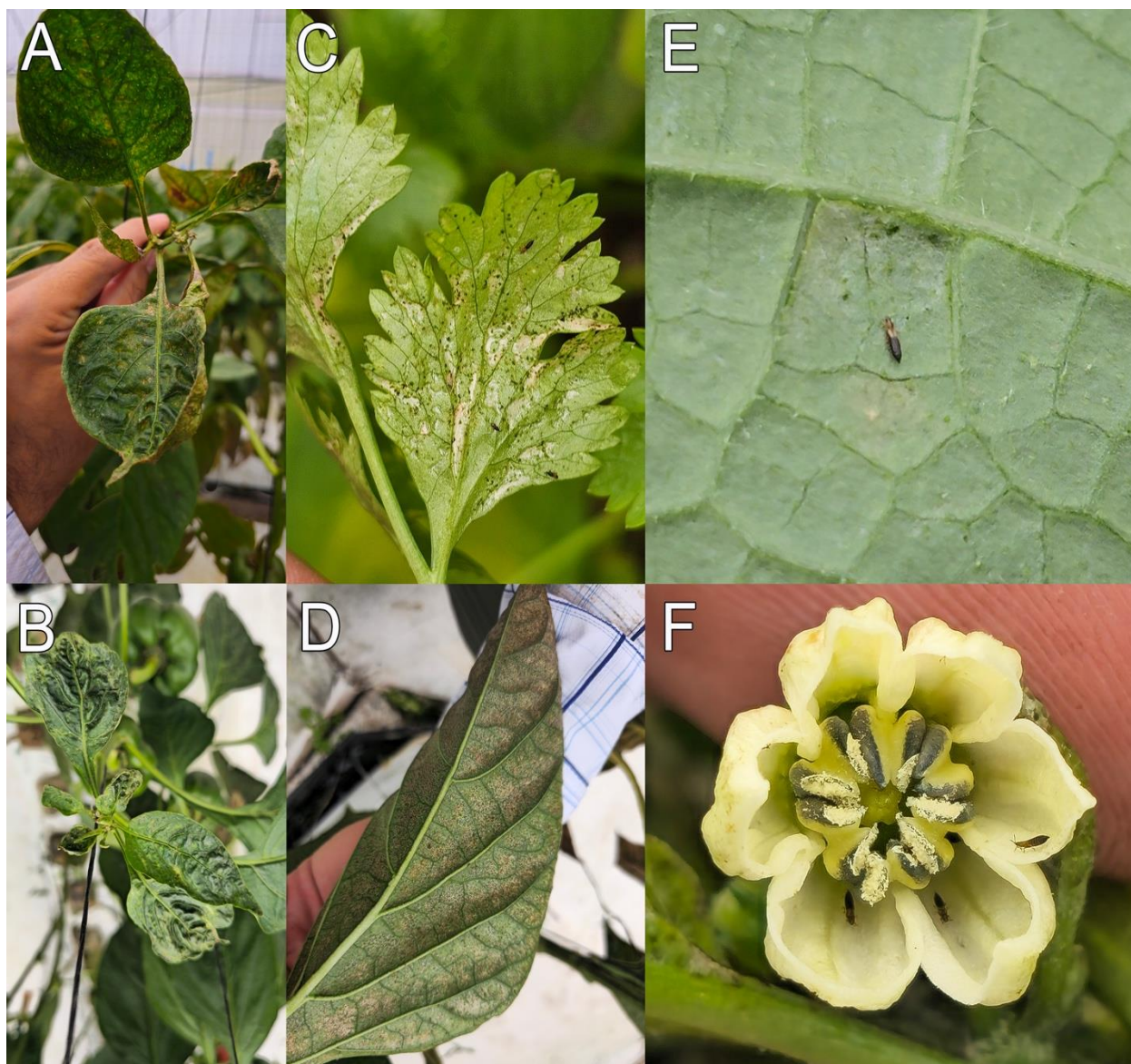


Figure 3. Damage inflicted by *Thrips parvispinus* on host plants. (A, B) Leaf curling and deformation of new shoots in *C. annuum*; (C) feeding activity observed on the leaf surface of *C. sativum*; (D) mature leaves displaying extensive abaxial tissue destruction in *C. annuum*; (E) feeding damage evident on the abaxial leaf surface in *C. sativum*; and (F) presence of the species on the flowers of *C. annuum*.

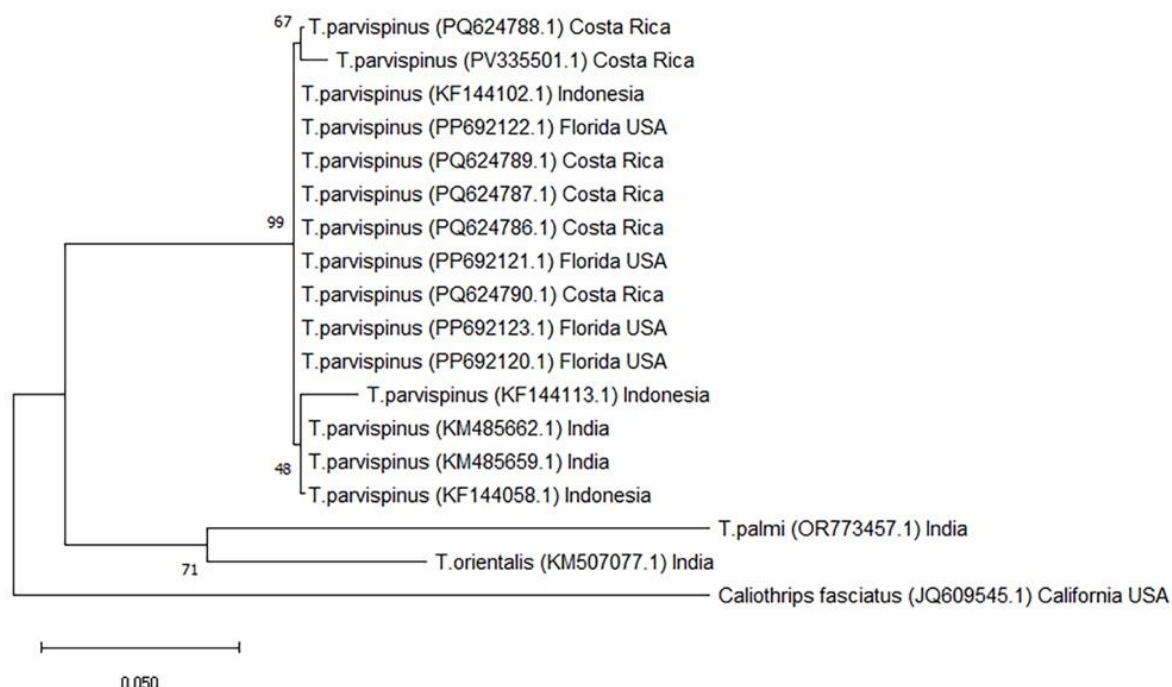


Figure 4. Phylogenetic relationships of *T. parvispinus* from Costa Rica. Maximum Likelihood tree constructed from partial COI gene sequence alignments. Bootstrap support based on 1,000 replicates was calculated using the Kimura 2-parameter model. Accession JQ609545.1, designated as *Caliothrips indicus* from California (U.S.A.), served as the outgroup.

Previous studies have demonstrated that *T. parvispinus* feeds on approximately 43 plant species from 19 families worldwide (Ataide *et al.*, 2024). In North America, in addition to *Capsicum*, it has been documented in *Gardenia*, *Hoya*, *Anthurium*, and *Mandevilla*, highlighting its significance in agricultural production (Ahmed *et al.*, 2024). The larvae primarily pierce and scrape the abaxial surface of the leaves, extracting cellular sap and resulting in a yellowish to brownish appearance in the affected areas (Thorat *et al.*, 2022). Damage to mature leaves manifests as streaks, leaf curling, or necrotic zones, while newly emerged leaves exhibit drying and wilting (Sridhar *et al.*, 2021). In flowers, adults cause damage by scraping petals and feeding on pollen, which leads to wilting and flower drop, affecting pollination and reducing fruit productivity (Sireesha *et al.*, 2021).

The presence of this species may also be a warning about its potential for viral vectoring in Central America, a phenomenon well documented for *T. parvispinus* in Asia (Rachana *et al.*, 2022). Furthermore, the successful establishment of these invasive species in new regions may result from the exchange of agricultural materials among trading countries coupled with limited phytosanitary measures (Ahmed *et al.*, 2024).

CONCLUSION

The recent detection of *Thrips parvispinus* in Costa Rica underscores its capacity for range expansion within the Americas, which has previously been reported in the United States, Canada, and Mexico. Its association with commercially important crops confirms its polyphagous nature in both protected and open-field agricultural systems. By identifying the specimens through external morphology and mitochondrial COI gene analysis, we confirmed its identity and detected genetic similarities with previously reported populations in North America, suggesting a potential introduction route linked to regional horticultural trade. This finding serves as a phytosanitary alert for the Central American region and emphasizes the urgent need to implement monitoring programs for invasive thrips.

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Disclosure statement. The authors declare no potential conflicts of interest.

Compliance with ethical standards. The authors confirm that this investigation was conducted under current ethical procedures. No human or animal was used in this study.

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Author contribution statement (CRediT). J.A. Rodríguez-Arrieta – Conceptualization, Writing and editing, Funding acquisition, Methodology, Data curation, Validation, Supervision. A. Barrientos-Castro – Resources, Methodology, Formal analysis, Visualization. J. Masis-Arce – Writing -original draft, Software. I. Varela-Benavides – Funding acquisition, Conceptualization, Supervision.

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