



Research article

Mechanical vector potential of houseflies (*Musca domestica*) for mites

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Abstract

Houseflies (*Musca domestica*), which live in common areas with humans, can be mechanical vectors of many pathogens. Mites are an extremely diverse community of organisms that occupy various microhabitats, and many species are harmful to human health. This study aimed to detect the presence of mites in the washing water of houseflies and assess the potential role of these flies in mite transmission. Houseflies collected from households within the boundaries of the Tatvan district in Bitlis were included in this study. Houseflies were gathered from 80 households, with 43 located in urban areas and 37 in rural regions, within the study's scope. After washing with PBS, the collected wash water was meticulously scrutinized under a light microscope to detect the presence of mites. Mites were found in 15 (18.7%) of the 80 samples, 8 (21.6%) of the rural samples, and 7 (16.3%) of the urban samples ($P=0.544$). It was determined that 40% of the identified mites belonged to the order Trombidiformes, 40% to Astigmata, and 20% to Mesostigmata. In conclusion, we believe that house flies act as vectors, play a role in the spread of mite populations, and can carry certain mites into homes that can affect the health of humans.

Keywords: *Acaridae; Bitlis; home dust; mesostigmata; mite; stroge mite*

1. Introduction

Houseflies (*Musca domestica*) are one of the most widespread species of flies worldwide, associated with humans, and can complete their entire life cycle close to human settlements (Adenusi et al., 2018; Park et al., 2019). They pose a threat to public health by mechanically vectoring pathogenic microorganisms. Flies that come into contact with garbage, sewage, and various contaminated environments can transfer pathogens to food via their mouthparts and other body surfaces, thereby mechanically transmitting disease agents (Ali et al., 2024; Lhoak and Al-Awadi, 2024). Mites can also be transported through mechanical transmission, and insect-mediated dispersal is a common mode of spread among mites

(Michalska et al., 2023).

Mites represent a highly diverse assemblage of organisms that occupy a variety of distinctive microhabitats. Some species inhabit the nasal passages of mammals, the tracheae of insects, human and other mammalian hair follicles (such as *Demodex* spp.) (Rychlik et al., 2025), the skin of mammals (such as scabies mites (Aydemir et al., 2025), feathers of birds, and skin scales of reptiles. In addition, many species are plant parasites, and some are significant pests of crops, such as citrus, apple, strawberry, bean, and wheat (Arlian and Platts-Mills, 2001; Kilinc et al., 2024). Mites found in house dust can trigger allergic reactions in humans, leading to allergic diseases such as asthma and atopic dermatitis (Wan et al., 2024). Food and storage mites can infest foodstuffs such as cereals, flour, nuts,

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<https://doi.org/10.51753/flsrt.1674545> Author contributions

Received 12 April 2025; Accepted 25 July 2025

Available online 30 August 2025

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and cheese, potentially causing food damage and allergic responses in humans (Bergmann, 2022; Akdemir et al., 2023).

This study aimed to evaluate the role of houseflies in the spread of mites by investigating the presence of mites in the water used to wash houseflies.

2. Materials and methods

In this study, house flies collected from 80 households, including 43 from the district center of Tatvan in Bitlis, Türkiye, and 37 from rural areas, were examined between June and July 2024. These flies were captured from inside the houses, with each household contributing between 10 and 15 specimens that were then placed in a 15 ml tube. Subsequently, they were transported to the Parasitology Laboratory at Van Yüzüncü Yıl University's Faculty of Medicine for examination of mite presence.

The collected samples were examined morphologically using a stereomicroscope (Olympus SZ61, Japan), and the species were identified based on the characteristics described in the literature (Nihei and Carvalho, 2007). Flies identified as *Musca domestica* were included in the study.

The methods described by Hamoo and Alnuri (2019) and Förster et al. (2009) were modified and applied for mite identification. To each centrifuge tube, 10 ml of PBS (1M, pH = 7.01) was added, and the contents were vortexed for approximately one minute. After vortexing, the flies were removed using a tweezer and discarded. The remaining liquid was centrifuged at 5000 rpm for five minutes. The supernatant was removed using a Pasteur pipette, leaving approximately one ml of liquid at the bottom of the tubes. After homogenizing the remaining liquid using a Pasteur pipette, a drop was placed on a microscope slide, covered with a coverslip, and examined under a light microscope using 10X and 40X objectives. Mites were identified using published identification keys (Spieksma, 1990; Colloff and Spiekma, 1992).

A Z-ratio test was used to compare the prevalence of mites between urban and rural households. Data tabulation was performed using the Minitab software package (version 14).

3. Results

In this study, mites were detected in 15 out of 80 samples (18.7%), including 8 of 37 samples from rural households (21.6%) and 7 of 43 samples from urban (central) households (16.3%). There was no statistically significant difference in mite detection rates between urban and rural dwellings ($p=0.544$). Of the detected mites, 6 (40%) belonged to the order Trombidiformes and 3 (20%) to the order Mesostigmata (Fig. 1). Additionally, three mites belonging to the order Astigmata were classified under the family Histiostomatidae (Fig. 2), and another three mites were identified as members of the family Acaridae (Fig. 3).

4. Discussion

Houseflies typically feed on waste such as feces and animal manure, as well as decaying organic material such as carrion, in their natural behavior pattern. When they feed on these substances, pathogens can adhere to their mouthparts, wings, and legs, allowing them to potentially transport these microorganisms into human settings (Khamesipour et al., 2018; Gul et al., 2024). The house fly is considered a potential mecha-



Fig. 1. A mite belonging to the order Mesostigmata.



Fig. 2. A mite belonging to the family Histiostomatidae.

nical vector for over 200 pathogens (D'Arco et al., 2024). The isolation of pathogens from the body of houseflies strongly suggests their role as mechanical vectors (Khamesipour et al.,

2018). Previous studies have found fungi (Ysquierdo et al., 2017), bacteria (Bahrndorff et al., 2014), viruses (Haddow et al., 2017), and parasites (Eke et al., 2016) in houseflies, indicating their potential to transmit various diseases through vectors. In this study, mites were detected in the washing water of 18.7% of flies. Therefore, we believe that house flies play a role as mechanical vectors in the spread of mites, similar to other pathogens.



Fig. 3. A mite belonging to the family Acaridae.

Approximately 40,000 mite species have been identified to date (Akpınar et al., 2015). Mites found in human environments are generally classified as either house dust mites or storage mites (Wong et al., 2011). Exposure to house dust mites has been shown by various studies to be positively associated with the development and severity of allergic diseases such as asthma, allergic rhinitis, atopic dermatitis, and atopic eczema (Cevik et al., 2024; Klain et al., 2024; Wan et al., 2024).

Storage mites, often overlooked in daily clinical practice, represent a more common respiratory allergen in children than previously assumed (Kaya et al., 2025). Species belonging to the families Pyroglyphidae, Glycyphagidae, and Acaridae are commonly observed in house dust (Akyazi et al., 2019). Many species from the families Glycyphagidae and Acaridae are known to occur as storage mites (Huppertz et al., 2020). Red

mites (Trombiculidae), belonging to the order Trombidiformes, are small larval mites that cause red mite dermatitis (trombiculosis). The main hosts of these mites are wild animals, and more than 50 species have been reported to infest humans, with approximately 20 species capable of causing diseases in both humans and animals (Sykes, 2022). Mites of the Histiogastriidae family typically inhabit the surface of moist organic matter, where they are presumed to subsist on bacteria and other microorganisms using their highly specialized palps and chelicerae. Nevertheless, a case report by El-Arfaj et al. (2007) documented an instance in which a histiogastriid mite was implicated in causing human autoacariidosis. In this study, mites belonging to the family Acaridae, which includes species recognized as both house dust and storage mites, were detected in house flies collected from the dwellings. Additionally, mites from the orders Mesostigmata and Trombidiformes, as well as from the family Histiogastriidae, were also identified. We suggest that this taxonomic diversity may be attributed to the ability of the flies to move freely between indoor and outdoor environments.

Many species of the family Acaridae are known to affect human health. *Acarus siro*, commonly found in flour and cheese, is a significant pest of stored foods. Contact with this mite can lead to dermatitis and allergies, whereas ingestion may cause intestinal and urinary tract diseases and malnutrition (Barbosa et al., 2022). *Tyrophagus putrescentiae* is one of the most important pests of stored food, infesting a wide range of products, including cereals, flour, nuts, dried fruits, cured meats, and cheese. It is also the most clinically relevant allergen-producing storage mite species associated with allergic respiratory diseases (Zhou et al., 2022). *Lepidoglyphus destructor* has also been shown to cause allergies in humans (Matos Semedo et al., 2024). Based on these findings, we suggest that the role of house flies in the transmission of mites from the Acaridae family is of significant concern for human health.

The major limitation of this study is that the species of mites could not be determined, as the mites in the preparations could not be identified using Hoyer's solution.

In conclusion, houseflies serve as mechanical vectors for the dissemination of mite populations and possess the capability to transport mites that may affect human health. While the role of houseflies as mechanical vectors for microorganisms has been extensively investigated, their potential as mite carriers has been explored in only a limited number of studies. This constitutes a significant knowledge gap, as mites on flies could facilitate the spread of parasitic infestations among animals, contaminate food products, and influence epidemiological dynamics in both the public health and veterinary domains. Understanding this interaction is crucial, particularly for developing comprehensive control strategies in environments where houseflies and mites coexist in substantial numbers.

Conflict of interest: The authors declare that they have no conflict of interests.

Informed consent: The authors declare that this manuscript did not involve human or animal participants and informed consent was not collected.

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Cite as: Yildiz, R., Afsar, M., Barlik, F., Aykut, M., Aydemir, S., & Ekici, A. (2025). Mechanical vector potential of houseflies (*Musca domestica*) for mites. *Front Life Sci RT*, 6(2), 131-134.