

OCCURRENCE OF POWDERY MILDEW CAUSED BY *Podosphaera* sp. ON INVASIVE WEED *Ageratum houstonianum* IN NEPAL

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ARTICLE HIGHLIGHTS

- First report of *Podosphaera* sp. on invasive *Ageratum houstonianum* in Nepal.
- Pathogen identified by unique cask-shaped conidia, chasmothecia and fibrosin bodies.
- Potential biocontrol agent discovered for managing this invasive weed.
- This weed can act as a pathogen reservoir; monitoring is essential to protect nearby crops.

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ABSTRACT

Ageratum houstonianum is an invasive annual weed in Nepal that grows in diverse habitats such as fallow areas, crop fields, and roadsides. This study reported a powdery mildew disease on this weed species, which causal agent was identified as *Podosphaera* sp. Pathogen identification was based on symptomatology and micromorphological observations. The pathogen was characterized by slightly dark, branched and septate hyphae with nipple-like appressoria; straight and unbranched conidiophore; hyaline and cask-shaped conidia with fibrosin-like bodies. Immature chasmothecia with mycelium-like appendages were also observed. The asci were ellipsoidal with a thin oculus. Based on the literature, the fungus was first reported in Nepal on *Ageratum houstonianum*. This report expands the known geographical distribution and host range of this powdery mildew pathogen in South Asia and highlights the pathogen as a potential biocontrol agent for this invasive weed. Additionally, *A. houstonianum* is common in agroecosystem, thus, it can serve as a reservoir of *Podosphaera* sp. facilitating spread to crop plants. Therefore, careful utilization, monitoring, and management of the pathogen are essential both to control the host weed and to limit the spread of pathogen to crops.

Keywords: blue goat weed, chasmothecia, Erysiphaceae, host range, *Sphaerotheca*

INTRODUCTION

Powdery mildew is a fungal disease caused by various species of ascomycetous fungi and affects a wide range of agricultural crops, weeds, and wild plants. Powdery mildew fungi are obligate pathogens that depend on living host cells for growth and development (Panstruga 2003). According to Adhikari (2022), 13 genera and 53 species of powdery mildew-causing fungi were reported on 76 flowering plant species from different regions of Nepal.

Ageratum houstonianum Mill. (Asteraceae) is an annual herbaceous plant native to Mexico and Central America (Johnson 1971) that has spread

across most of the tropical to temperate regions of Asia, Europe and Australia (CABI 2022). This weed is widely recognized as one of the most problematic species in Nepal, with extensive distribution across mid-hill and lowland regions, including roadsides, gardens, protected areas, agricultural land, community forests, and fallow land (Shrestha *et al.* 2019). It reduces agricultural productivity, depletes soil fertility, and competes with native plants. It also causes cattle poisoning and reduces fodder availability (Bhatta *et al.* 2022; Pal *et al.* 2011). Furthermore, the weed alters soil physicochemical characteristics, competes for resources, releases allelochemicals, and interferes with pollination (Shrestha *et al.* 2018; Shrestha *et al.* 2019).

Globally, *A. houstonianum* is affected by several diseases, including downy mildew (caused by fungus *Plasmopara halstedii*), powdery mildew (caused by *Podosphaera xanthii*), leaf blight (induced by *Botrytis cinerea*), and root rot (caused by *Phytophthora* and *Pythium*) (Bose *et al.* 2025; Baiswar *et al.* 2008; Pisani *et al.* 2019; Reeves *et al.* 2022; CAES 2022). However, reports of these diseases in Nepal are very limited.

In October 2020, a severe outbreak of powdery mildew on *A. houstonianum* was observed in a sal (*Shorea robusta*) forest located in Kailali District of far-western Nepal. Accordingly, this study was conducted to identify the causal organism responsible for the disease and to evaluate its potential as biocontrol agent for invasive weeds like *A. houstonianum*. In addition, the weed may act as a reservoir for powdery mildew pathogens, posing indirect risks to nearby crops and native flora (Yeh *et al.* 2021; Singh *et al.* 2023).

MATERIALS AND METHODS

Field Survey

Barhaban Collaborative Forest, Gauringanga Municipality-6, Kailali District of Sudurpashchim Province, Nepal (28°48'34.34" N, 80°40'37.63" E; altitude 204 m asl) was surveyed in October 2020. During transect walks along roadsides and trails across the forest, forest edges and within the forest interior, *Ageratum houstonianum* was widely distributed as an invasive weed across all surveyed habitats. Powdery mildew symptoms were frequently observed on the weed population at multiple locations.

Collection of Disease Samples

At each representative site (roadsides, trails, and forest edges), *A. houstonianum* plants were inspected (n = 30). From each site, five symptomatic leaves were collected, placed in sterile paper bags, properly labelled with sample ID and collection date, and stored in a gel icebox for transport. The photographs of infected plants and symptomatic leaves were also taken in situ. Leaf samples and herbarium specimens (voucher no. WS-KL-1252 & WS-KL-1266) were transported to the Plant Pathology Laboratory at the Forest Research and Training Center (FRTC), Department of Forest, Government of Nepal, Kathmandu.

Microscopy and Measurements

Powdery mass from diseased leaf surface was transferred onto microscope slides using the Scotch tape method (Larone 1995) and mounted in lactophenol cotton blue (LPCB) staining. Slides were examined under 100X, 400X and 1000X magnifications using an OPTIKA Trinocular Microscope B-293 to observe mycelia, appressoria, chasmothecia, conidia, asci, and conidiophores. All micromorphological characteristics were documented and photographed, while measurements were taken for each structure. The fungus observed was identified using the keys of Braun and Cook (2001) and Amano (1986). Molecular confirmation was not performed due to resource limitations. Morphometric measurements were made using ImageJ software (version 1.48).

RESULTS AND DISCUSSION

Powdery mildew symptoms on *Ageratum houstonianum* were documented at the study site. Infected leaves exhibited multiple oval to irregular patches of grayish-white powdery growths. As colonies matured, they coalesced to form dense white masses across the leaf lamina (Figure 1B, C). Microscopic examination showed that the powdery growth on the upper leaf surface comprised epiphytic mycelia and conidia (Figure 1C). Symptoms were less conspicuous on the lower leaf surface. Severe infection caused leaf necrosis, defoliation, and plant collapse (Figure 1A).

Powdery mildew disease caused by *Podosphaera xanthii* has been previously reported on *Ageratum conyzoides* in India, Pakistan and China (Nadeem *et al.* 2018; Mukhtar *et al.* 2013; Mukhtar & van Peer 2017). The pathogenic genus *Podosphaera* associated with powdery mildew disease on *Ageratum houstonianum* was first reported in India by Baiswar *et al.* (2008) and later confirmed as *Podosphaera xanthii* by Bose *et al.* (2025). These findings highlight the widespread occurrence of *Podosphaera* species as causal agents of powdery mildew in *Ageratum*.

Hyphae of currently reported pathogen were found to be septate, slightly dark, branched, 4.01 – 5.91 μm (n = 10) wide with nipple-like appressoria (Figure 1F). Conidiophores were straight and unbranched, foot-cells cylindrical with basal septa, 50.75 – 117.72 $\times$ 4.24 – 7.57 μm (n = 8), followed by one to three shorter cells (Figure 1D). Conidia were single borne or sometimes in chains, hyaline, barrel shaped (blunt rectangular type or somewhat

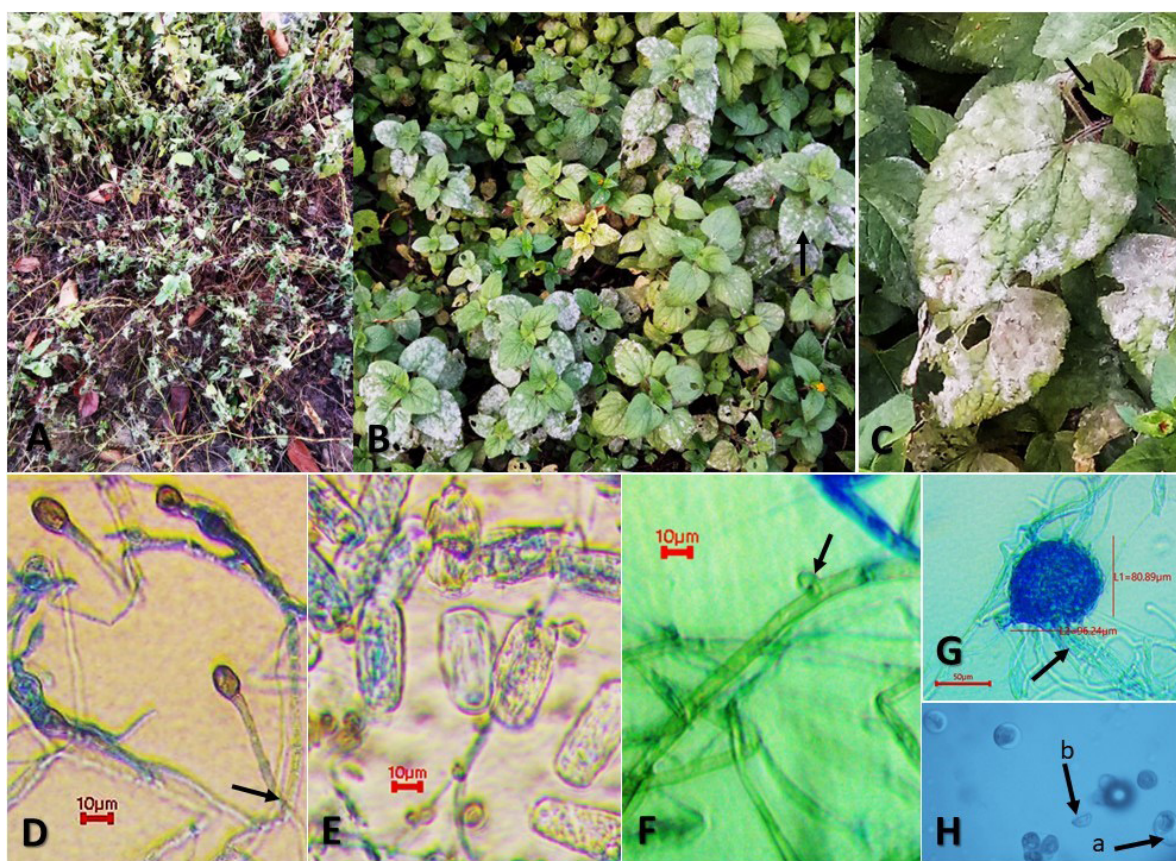


Figure 1 Powdery mildew on *Ageratum houstonianum*.

Notes: A = Destruction of *Ageratum houstonianum* due to severe infection by *Podosphaera* sp; B = Infected plants with initial infestation (arrow); C = Characteristic symptoms of disease on mature leaves with necrotic lesion and unaffected young leaves (arrow); D = Conidiophores with single conidium and basal septum (arrow); E = Conidia; F = Septate mycelium with nipple-like appressorium (arrow); G = an immature chasmothecium with lateral mycelial appendages (arrow); H = Asci showing thin oculus (arrow a) and deformed ascus without ascospore (arrow b).

ellipsoidal too), $31.95 - 43.52 \times 14.72 - 18.63 \mu\text{m}$ ($n = 10$) sized with fibrosin-like bodies (Figure 1E). Immature chasmothecia (formerly cleistothecia) of size $64.37 - 135.64 \mu\text{m}$ ($n = 15$) in diameter with appendages of mycelium-like bodies were also observed (Figure 1G). The asci were ellipsoidal with diameter of $45.61 - 63.59 \mu\text{m}$ ($n = 5$) with a thin-layer ascus wall i.e., oculus (Figure 1H, arrow a). Some deformed empty asci were also observed (Figure 1H, arrow b), while no ascospores were observed. These measurements were compared with three most common species of *Podosphaera* causing powdery mildew frequently in plants (Table 1). The measurements were consistent with the taxonomic description of *Podosphaera xanthii* (Baiswar *et al.* 2008; Braun & Cook 2012).

Based on micromorphological characters, the pathogen on *A. houstonianum* was identified as *Podosphaera*, with *Sphaerotheca* as a known synonym. The micromorphological measurements were

consistent with *Podosphaera xanthii* sensu lato. However, molecular analyses were not carried out in this study due to resource limitations. Therefore, DNA sequencing is essential and recommended to confirm species-level identification of the pathogen.

Previous studies have confirmed *Podosphaera xanthii* as the pathogen causing powdery mildew disease on *Ageratum conyzoides* (congeneric member of *A. houstonianum*) across South to East Asia (Aumentado *et al.* 2023), including Pakistan (Mukhtar *et al.* 2013), India (Amano 1986; Thite *et al.* 2017), China (Mukhtar & van Peer 2017) and Thailand (Meeboon *et al.* 2018). Powdery mildew on *A. houstonianum* caused by *Podosphaera* has been reported only from India (Baiswar *et al.* 2008), and was recently confirmed as *Podosphaera xanthii* by Bose *et al.* (2025). This study also resolved the taxonomic ambiguity with *Podosphaera fusca*, as both names were shown to represent the same species.

Table 1 Comparison of the micromorphological characters of the *Podosphaera* sp.

Species	Hyphae and appressoria	Conidiophore	Conidia	Chasmothecia	Asci	Oculus	References
<i>Podosphaera</i> sp	Hyphae superficial; nipple-like appressoria, 4.01 – 5.91 µm;	50.75 – 117.72 × 4.24 – 7.57 µm	Ellipsoidal or blunt rectangular; 31.95 – 43.52 × 14.72 – 18.63 µm; fibrosin-like bodies present	64.37 – 135.64 µm diameter; appendages mycelium-like	1 – 3 per chasmothecium, 45.61 – 63.59 µm diameter	Distinct	Present study
<i>Podosphaera xanthii</i>	Hyphae superficial; nipple-like appressoria, 6 – 10 µm	40 – 80 µm long	Ellipsoid, 20 – 35 × 12 – 20 µm; fibrosin-like bodies present	80 – 120 µm diameter; appendages mycelium-like	2 – 3 per chasmothecium, 40 – 65 × 25 – 35 µm	Distinct 5 – 8 µm	Choi <i>et al.</i> (2022)
<i>Podosphaera leuotricha</i>	Hyphae superficial; appressoria 6 – 9 µm	50 – 90 µm long	Cylindrical to ellipsoid, 18 – 30 × 10 – 15 µm	90 – 130 µm diameter; appendages flexuous	6 – 8 per chasmothecium, 35 – 55 × 20 – 30 µm	Thickened, 5 – 7 µm	Strickland <i>et al.</i> (2021)
<i>Podosphaera aphanis</i>	Hyphae dense; appressoria 5 – 8 µm	40 – 70 µm long	Ellipsoid, 18 – 28 × 10 – 15 µm	70 – 110 µm diameter; hypha-like appendages	4 – 6 per chasmothecium, 30 – 50 × 18 – 25 µm	Visible, 4 – 6 µm	Nadeem <i>et al.</i> (2018)

Podosphaera xanthii is a polyphagous, obligate biotroph capable of colonizing diverse hosts belonging to the Asteraceae and Cucurbitaceae (Polonio *et al.* 2019). Infection caused by this pathogen on *A. houstonianum* indicates host plasticity and adaptive potential that may facilitate host range expansion (Yeh *et al.* 2021). On one hand, infection of *A. houstonianum* may reduce weed vigor and potentially alter competitive interactions within invaded plant communities. On the other hand, survival of the pathogen on multiple hosts may stabilize local populations and serve as inoculum reservoirs (Singh *et al.* 2023). This may enable the fungus to bridge the seasonal gaps in host availability and persist across crop rotations as well as geographic ranges (Thines *et al.* 2023).

A recent revision of the checklist of powdery mildew in Nepal (Adhikari 2022) did not include this pathogen on *A. houstonianum*. To the best of our knowledge and based on the available literature, this represents the first record of powdery mildew disease on this weed in Nepal. This finding expands the known host range of *Podosphaera* and contributes to the understanding of powdery mildew diversity in the region.

The detection of *Podosphaera* on *A. houstonianum* in Nepal raises two practical concerns. First, the survival of pathogen at the landscape level may complicate crop protection strategies. The genus *Podosphaera* is well known for its ability to survive in diverse ecological niches, including wild hosts, weeds, ornamentals, and volunteer plants, which can act as inoculum reservoirs (Amano 1986; Thines *et al.* 2023). Second, invasive plants nearby agrosystems require regular monitoring for disease management (Thite *et al.* 2017), as they may serve as inoculum reservoir (Singh *et al.* 2023). To manage these potential risks, efforts should focus on molecular confirmation of the pathogen through DNA sequencing (Polonio *et al.* 2019) and systematic surveys of both the pathogen and host plants across different habitats.

CONCLUSION

This study reported the first record of powdery mildew disease caused by *Podosphaera* sp. on *Ageratum houstonianum* in Nepal. The micromorphological characteristics are consistent with *Podosphaera xanthii* sensu lato, for which molecular confirmation is recommended. *Ageratum houstonianum* is recognized globally as one of the most prevalent invasive plant species, commonly

found along roadsides and crop field margins. *Podosphaera* sp. may have potential as a biocontrol agent for controlling invasive *A. houstonianum*; however, the host plant may also serve as inoculum reservoir, linking wild and crop hosts and bridging seasonal gaps in host availability. Therefore, integrated monitoring and weed disease management strategies are recommended to reduce potential risk to susceptible crops.

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