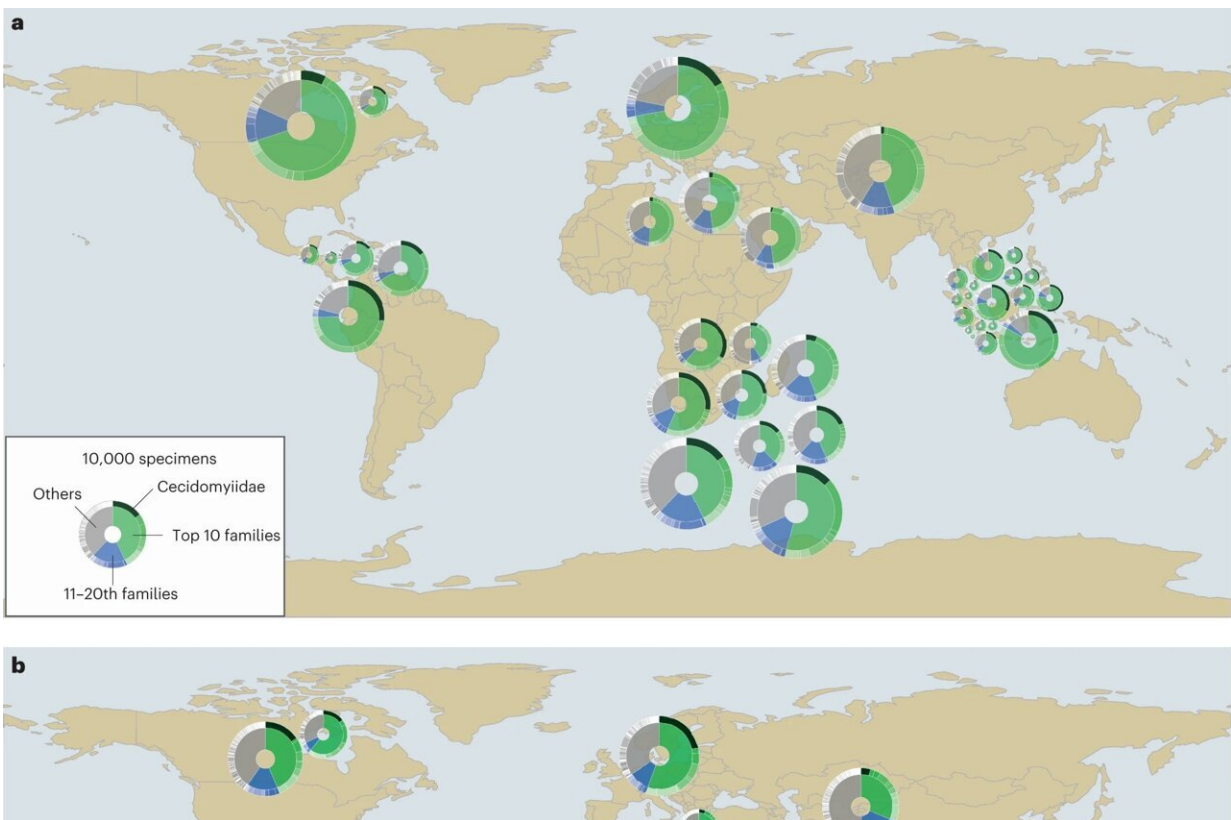


Biodiversity discovery: Unknown species ('dark taxa') drive insect diversity

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Congruence in the relative contribution of insect families to specimen abundance and species richness. **a,b**, Each chart shows the taxonomic composition of a sample obtained by an individual Malaise trap at a specific site. The inner circle represents the proportion of biodiversity in the top 10 (green), next 10 (blue) and remaining families (grey). The outer ring shows what proportion of biodiversity belongs to the top 10, next 10 and the remaining families. Black is used to illustrate the extraordinary diversity of Cecidomyiidae (Diptera). All charts are scaled relative to number of specimens (**a**) and species (**b**) at each site. Map

made with Natural Earth. Supplementary Fig. 2 provides precise geolocations for each site. Credit: *Nature Ecology & Evolution* (2023). DOI: 10.1038/s41559-023-02066-0

Biodiversity loss ranks among the top three risks to humanity, as stated in the 2023 World Economic Forum Global Risks Report.

Understanding biodiversity's basic building blocks is essential to monitor changes, identify influencing factors, and implement appropriate policies. However, much of terrestrial animal diversity, including insects, remains unknown or "dark taxa."

For example, the global [biodiversity](#) information portal GBIF has nine times more information on birds than [insects](#) and arthropods, despite birds only accounting for 0.2% of biodiversity. Prof. Rudolf Meier, head of the Center for Integrative Biodiversity Discovery at the Museum für Naturkunde Berlin, highlights the importance of this study in addressing this deficit for effective nature conservation. He emphasizes the need to learn more about insects, as their combined biomass and biodiversity far surpass that of all vertebrates, and they are crucial for survival.

To determine the global taxonomic composition of flying insects, researchers used Malaise traps. These standardized traps are widely employed in global biomonitoring programs. However, analyzing samples is challenging when examining individual insects. Thanks to recent advances in sequencing technologies, biodiversity can now be estimated with "DNA barcodes."

Researchers used DNA barcodes to assign 225,261 specimens to 25,000 species to 458 families. Dr. Amrita Srivathsan, the study's lead author, noted the surprising finding that 10–20 families dominate flying insect communities worldwide. This is remarkable as samples were collected

from various climates and habitats like [tropical rainforests](#), montane forests, cedar savannas, scrub forests, thorn fields, mangroves, and swamps, with only Australia and Antarctica not being sampled.

Describing insects is a major challenge in understanding life on Earth, with over 80% still undescribed. The authors emphasize that a large fraction of terrestrial animal biodiversity remains unknown to science and will continue to be, unless "dark taxa" become a priority target in biodiversity research.

The work is published in the journal *Nature Ecology & Evolution*.

More information: Rudolf Meier, Convergence of dominance and neglect in flying insect diversity, *Nature Ecology & Evolution* (2023).

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