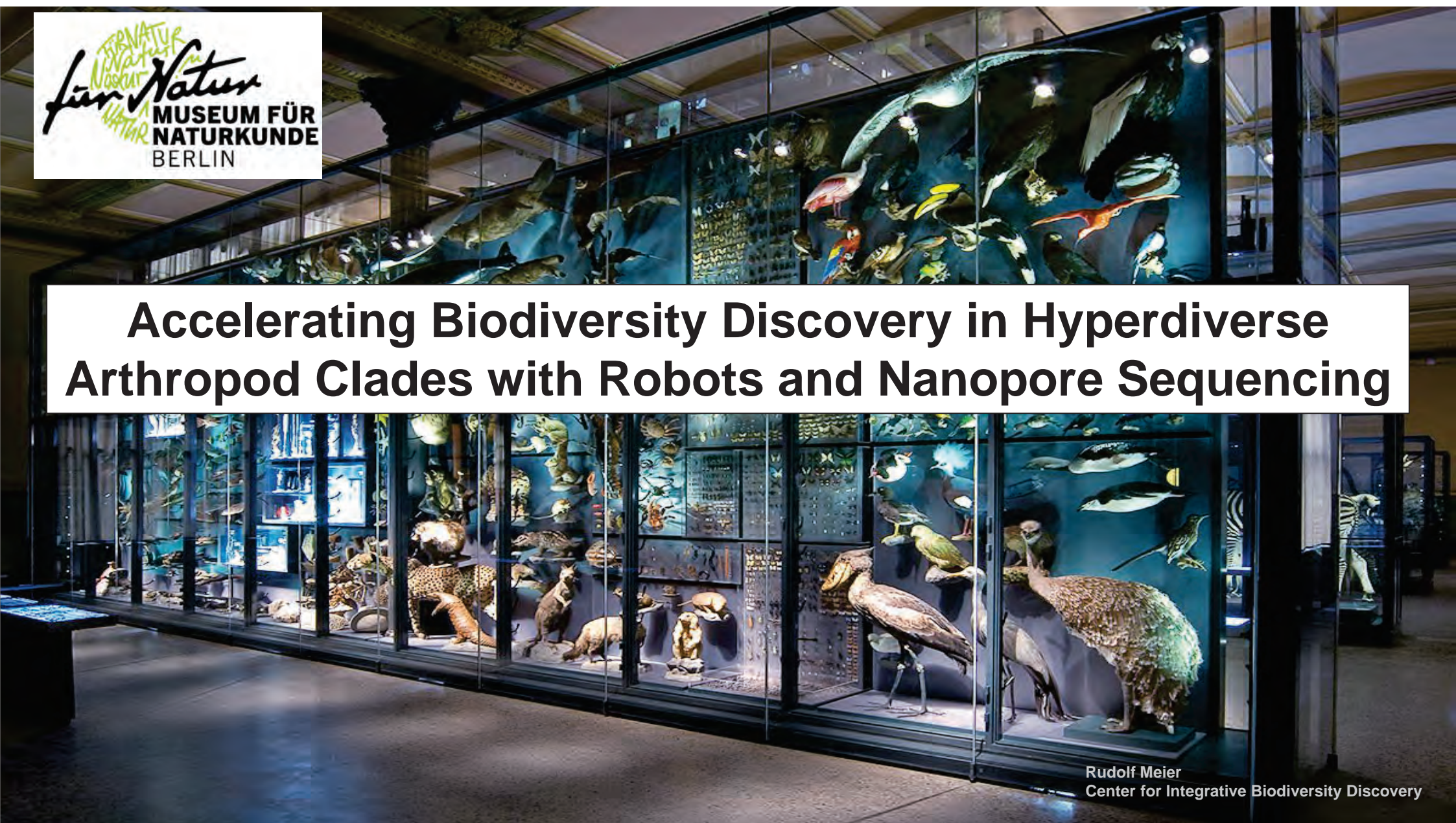




# Accelerating Biodiversity Discovery in Hyperdiverse Arthropod Clades with Robots and Nanopore Sequencing



Rudolf Meier  
Center for Integrative Biodiversity Discovery

# Our planet has a big biodiversity problem



1. Biodiversity Loss: threatens the survival of human societies
2. Lack of unbiased data to address the problem



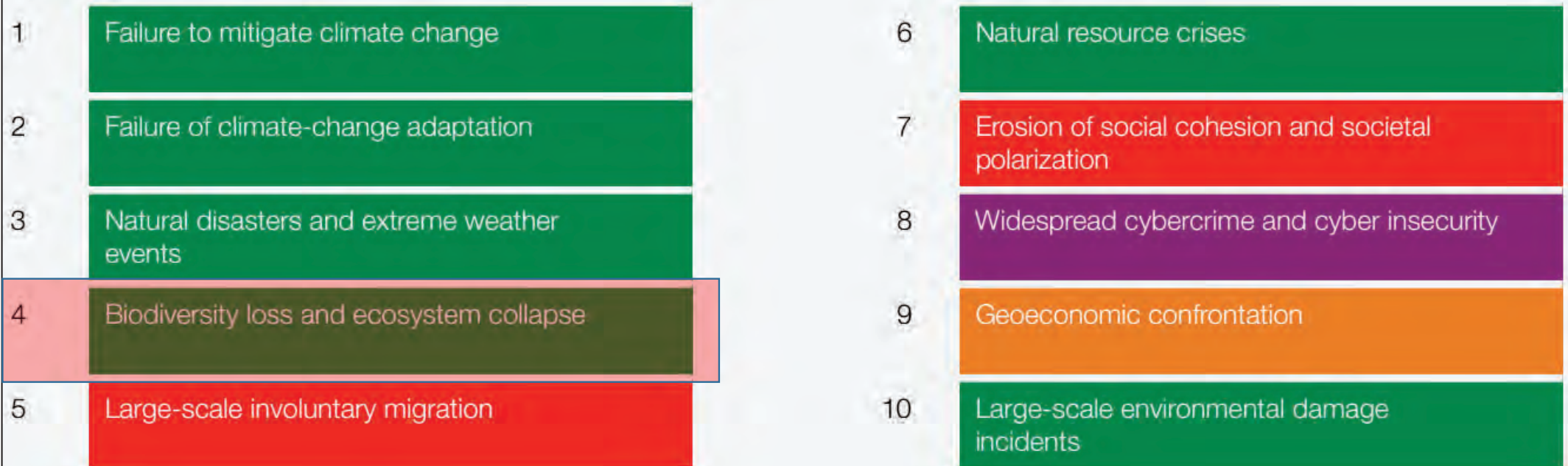
# The Global Risks Report 2023

## 18th Edition



- 1 Failure to mitigate climate change
- 2 Failure of climate-change adaptation
- 3 Natural disasters and extreme weather events
- 4 Biodiversity loss and ecosystem collapse
- 5 Large-scale involuntary migration

## Global risks ranked by severity over the long term (10 years)



Risk categories


- Economic
- Environmental
- Geopolitical
- Societal
- Technological

# Not only the World Economic Forum...



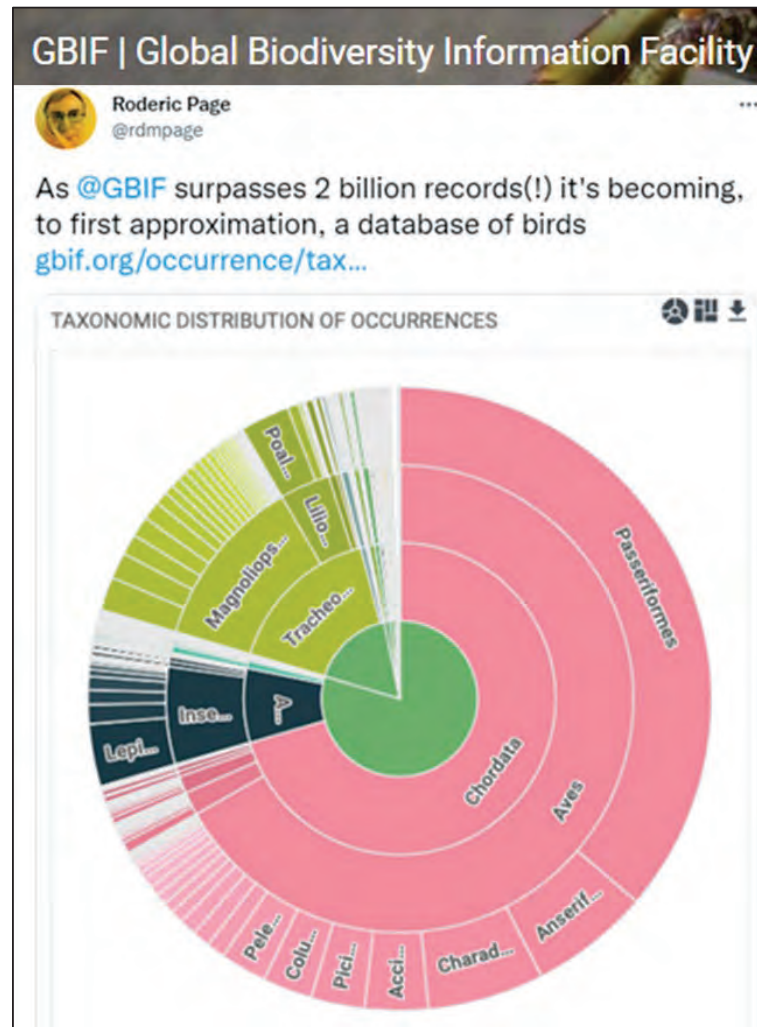
## 2. Lack of unbiased data to address the problem

GBIF | Global Biodiversity Information Facility





## 2. Lack of unbiased data to address the problem



## 2. Lack of unbiased data to address the problem

**Forbes**



FORBES > LEADERSHIP > LEADERSHIP STRATEGY




# Ugly Species Deserve Biodiversity Protections, Too

World Economic Forum Contributor 

[Follow](#)

Nov 2, 2020, 04:55am EST

 Listen to article 9 minutes 

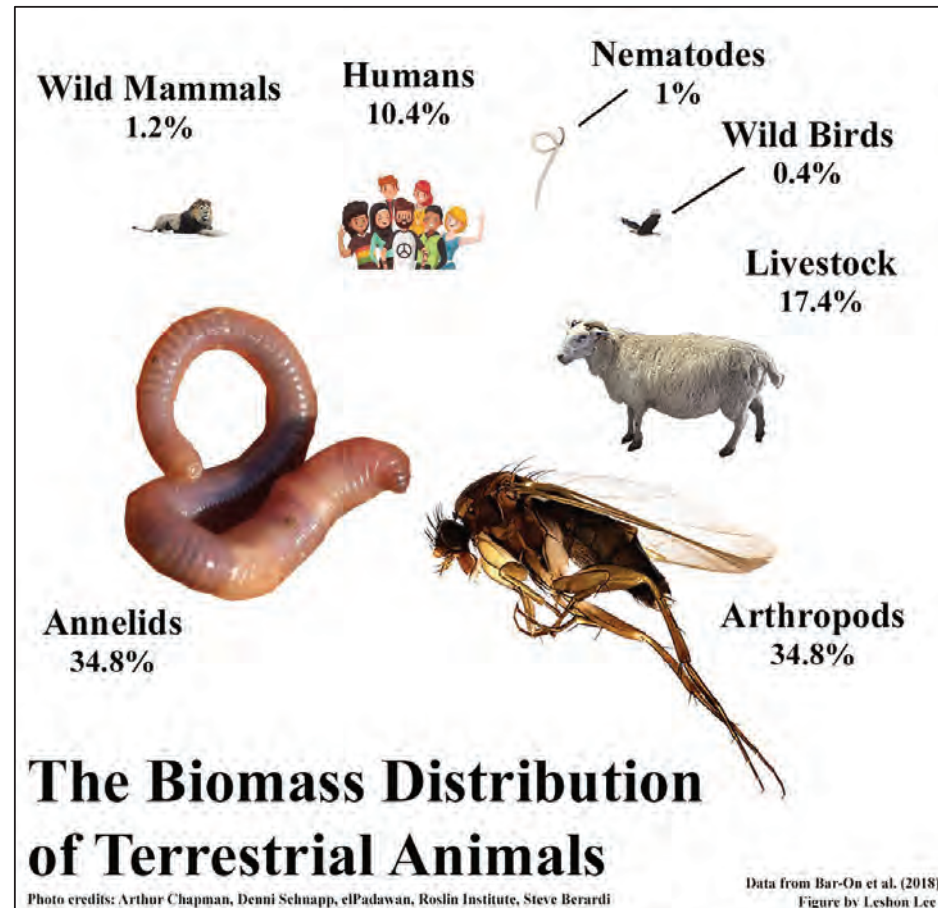


Hello, PHOTOHOLGIC/UNSPLASH



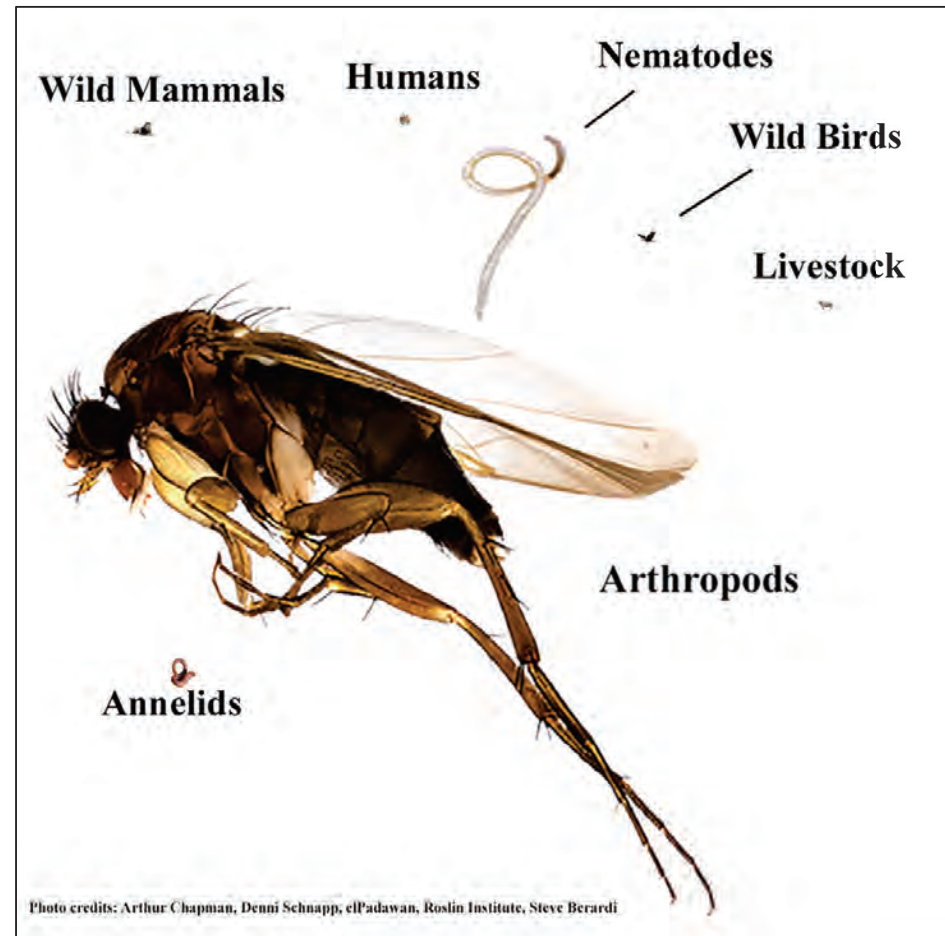
# “Ugly” species matter

**Taxon bias:**  
terrestrial animal biomass



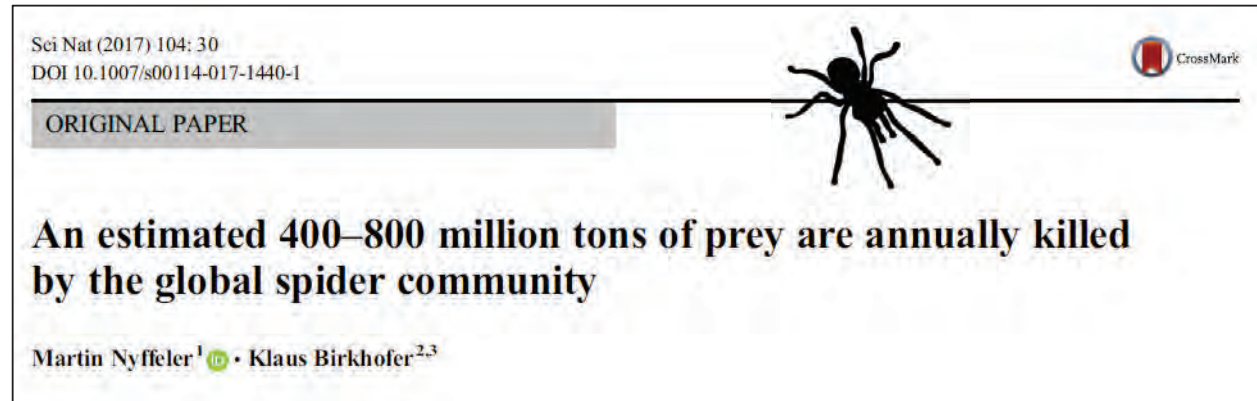
# “Ugly” species matter

**Taxon bias:**  
animal species diversity



# “Ugly” species matter

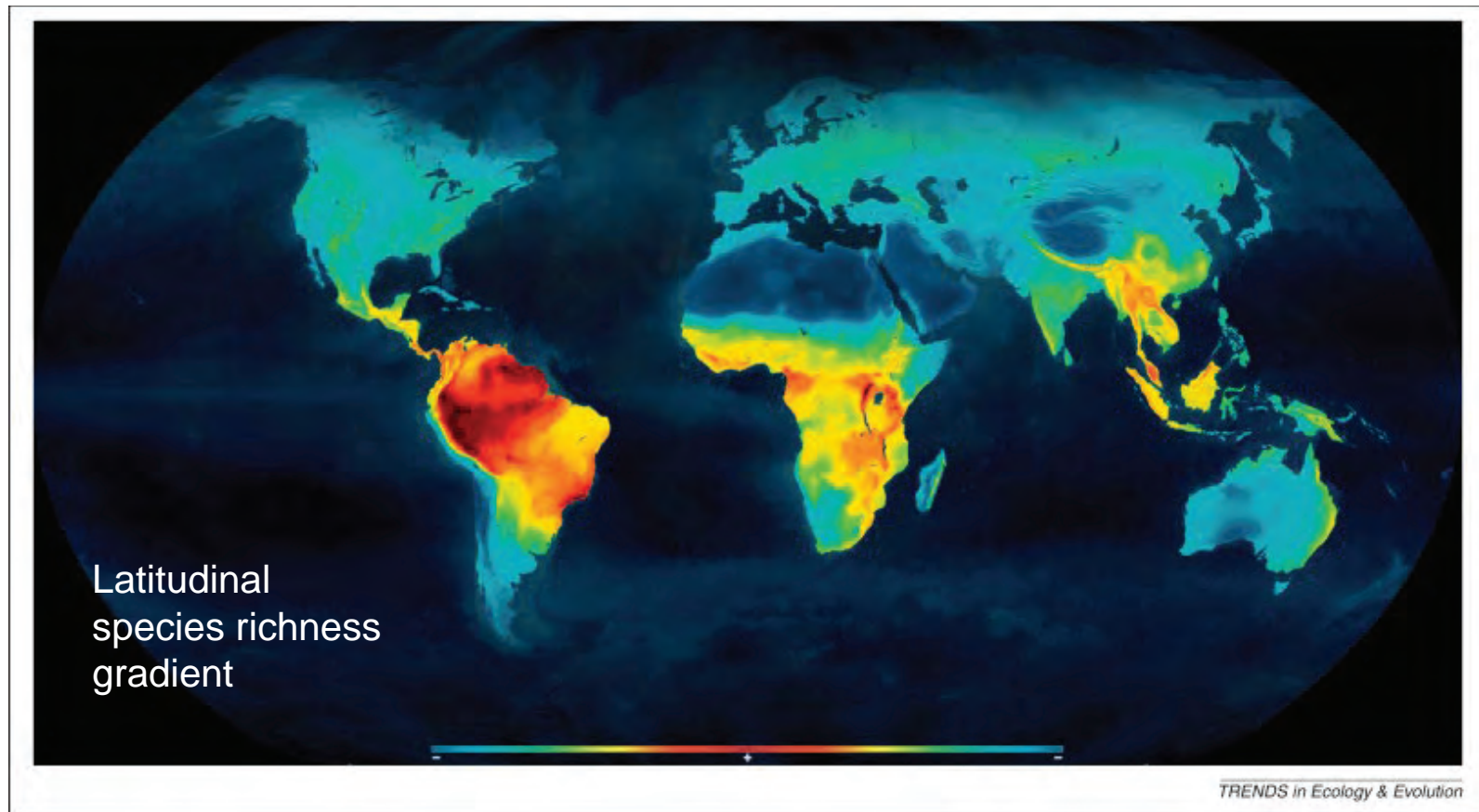
**Taxon bias:**  
Ecosystem impact



**Humans:** annual meat/fish consumption: 400 mio. tons



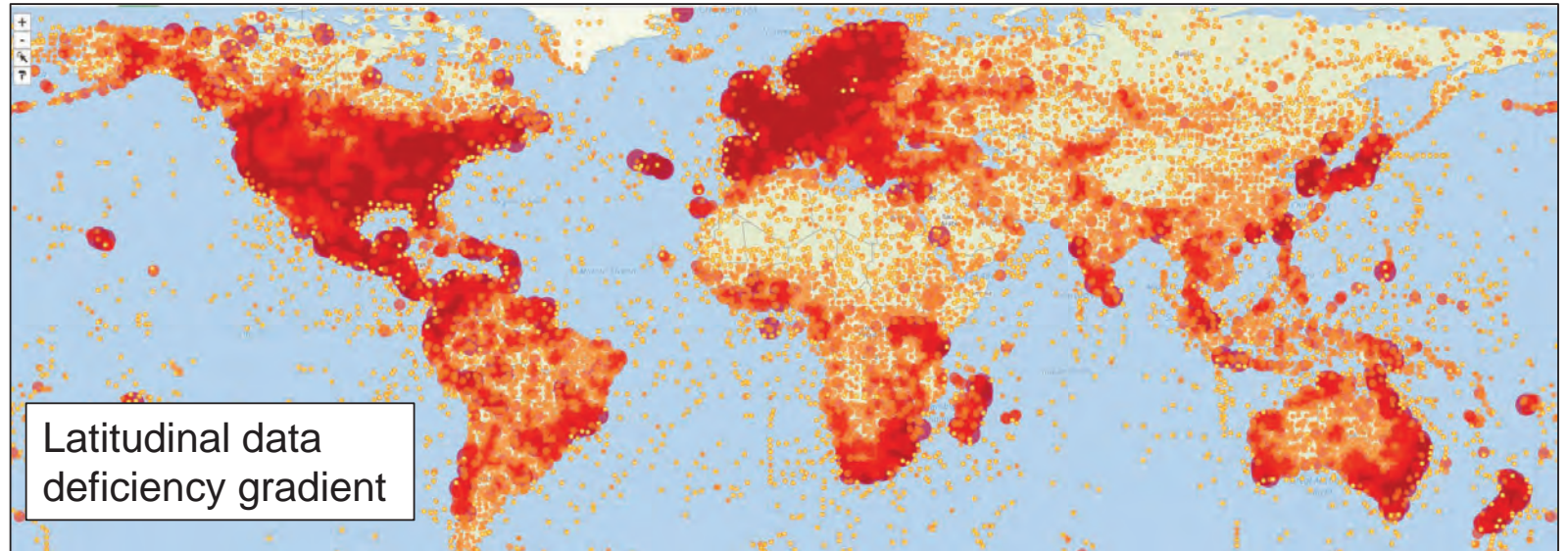
## Biodiversity data problem is worse at the global scale

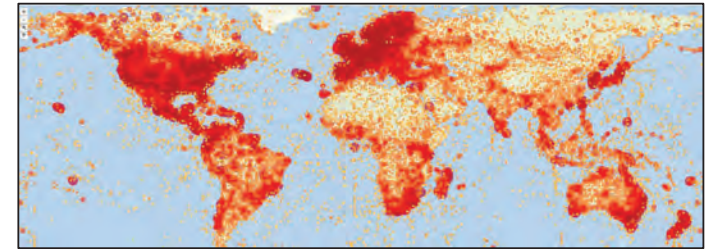


# GBIF insect data

## Insect data in GBIF:

- 169 million records
- 47 million specimens





Biodiversity: overcome the taxon and geographic biases

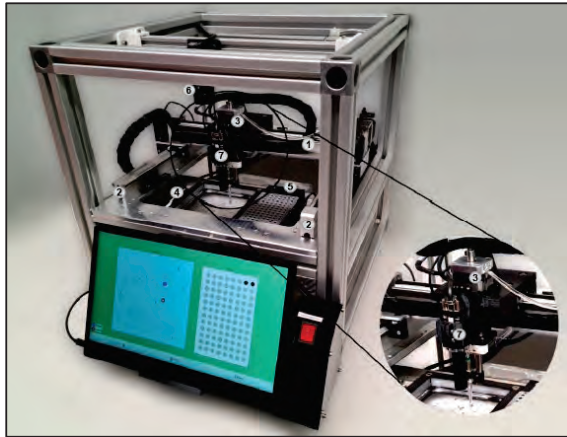
- we need cost-effective methods for collecting data
- methods have to be suitable for machines and humans alike



# Species Discovery Factory



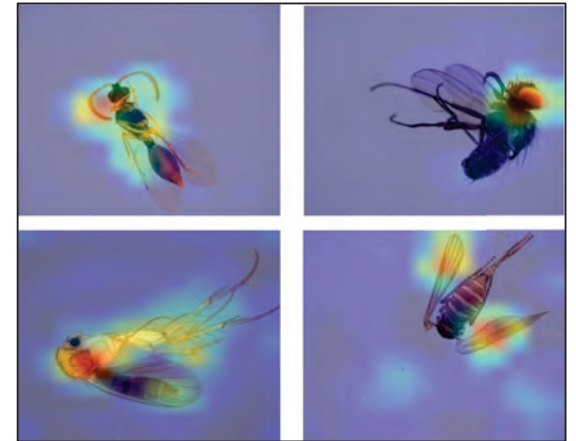
**1. DiversityScanner:**  
Digitization with robots

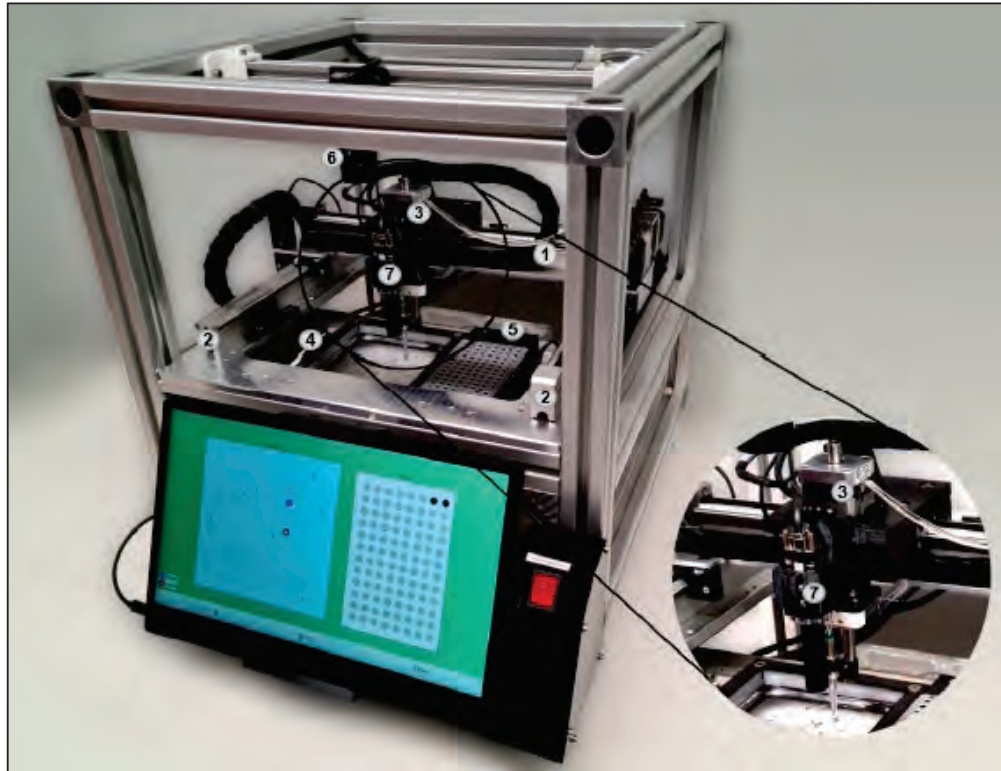


**2. Nanopore Sequencing:**  
Sorting with DNA barcodes



**3. Identified images:**  
biomonitoring with AI






- robot finds insects in a sample
- images each specimen
- identifies the specimen
- measures specimen
- prepares it for DNA sequencing

## MOLECULAR ECOLOGY RESOURCES

RESOURCE ARTICLE | [Open Access](#)

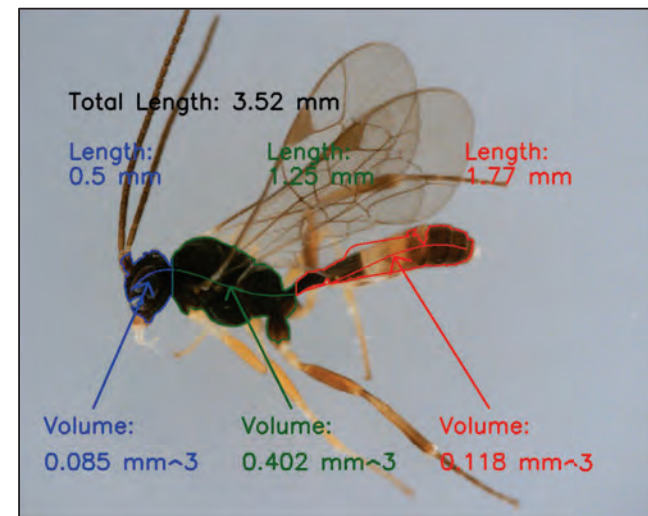
### DiversityScanner: Robotic handling of small invertebrates with machine learning methods

Lorenz Wühl, Christian Pylatiuk , Matthias Giersch, Florian Lapp, Thomas von Rintelen, Michael Balke, Stefan Schmidt, Pierfilippo Cerretti, Rudolf Meier 

First published: 04 December 2021 | <https://doi.org/10.1111/1755-0998.13567>

This article has been accepted for publication and undergone full peer review but has not been through the copyediting, typesetting, pagination and proofreading process, which may lead to differences between this version and the Version of Record. Please cite this article as doi:10.1111/1755-0998.13567

DiversityScanner  
Biomass estimation



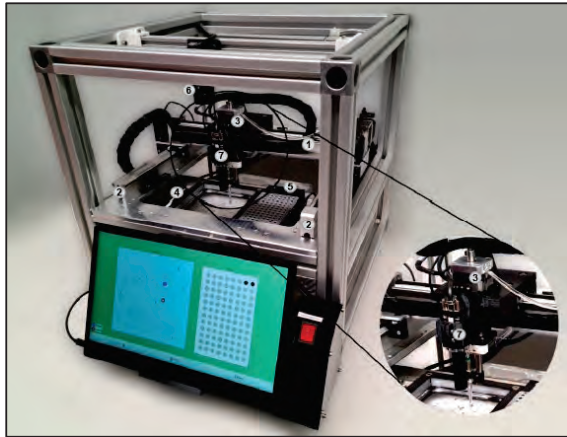




# Species Discovery Factory



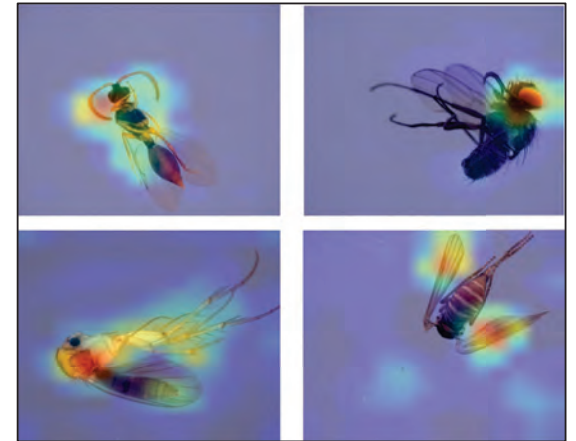
## 1. DiversityScanner: Digitization with robots



## 2. Nanopore Sequencing: Sorting with DNA barcodes



## 3. Identified images: biomonitoring with AI



## 2. Nanopore Sequencing



### Mobile labs for biodiversity discovery:

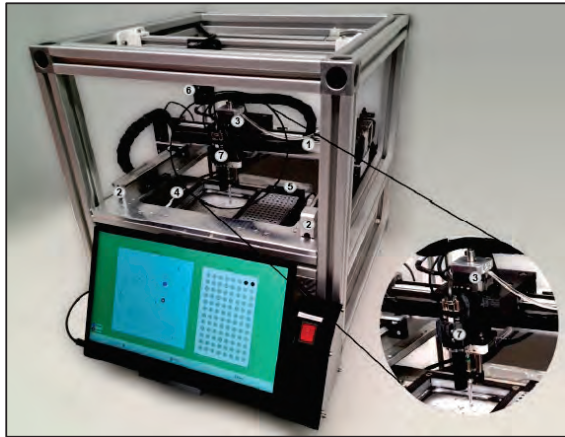
- simplified, fast, and robust techniques for specimen-based barcoding (10 cents/specimen)
- MinION: pocket-size sequencer



# Species Discovery Factory



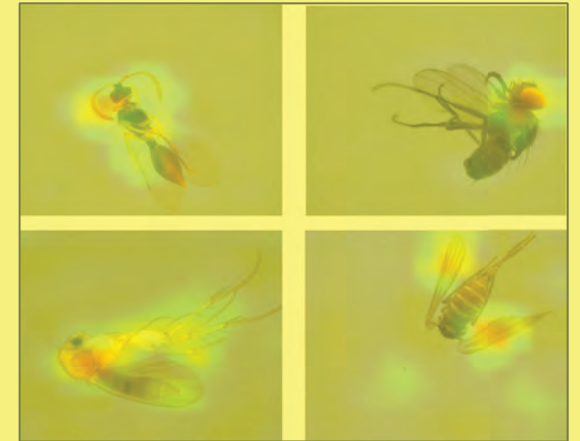
## 1. DiversityScanner: Digitization with robots



## 2. Nanopore Sequencing: Sorting with DNA barcodes

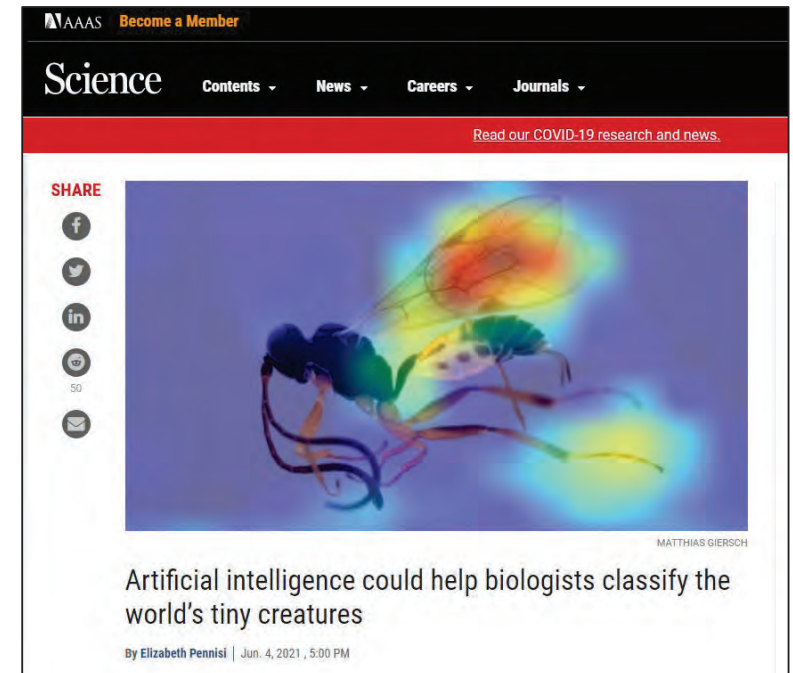
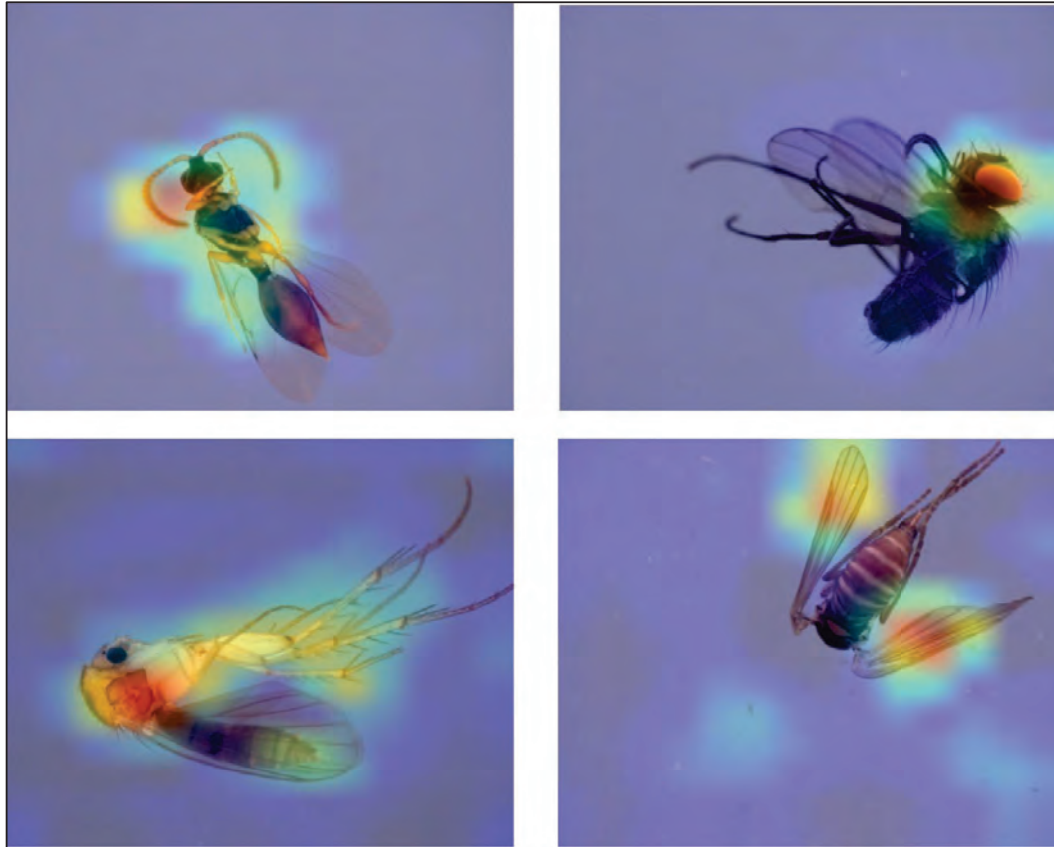


## 3. Identified images: biomonitoring with AI





### 3. Identification with AI: training of Convolutional Neural Networks



### 3. Identification with AI: training of Convolutional Neural Networks

Class (Taxon)	Result	Class (Taxon)	Result
Diptera Acalyptratae	91%	Diptera Psychodidae	89%
Diptera Calyptratae	83%	Diptera Sciaridae	92%
Diptera Cecidomyiidae	91%	Hemiptera Cicadellidae	100%
Diptera Chironomidae	97%	Hymenoptera Braconidae	82%
Diptera Dolichopodidae	86%	Hymenoptera Diapriidae	100%
Diptera Empididae & Hybotidae	87%	Hymenoptera Ichneumonidae	75%
Diptera Keroplatidae & Mycetophilidae	99%	Other	81%
Diptera Phoridae	97%		

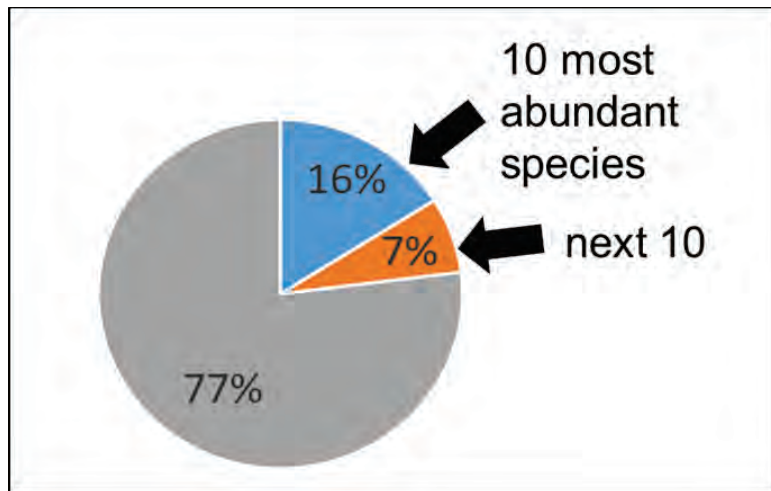
AI identification tools evolve

- common taxa benefit first
- evolve from family => genus => species

## For how many species do we need AI identification tools?

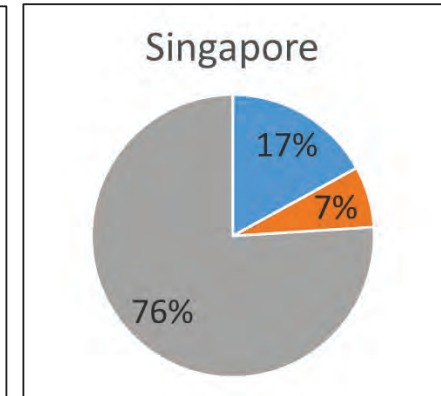
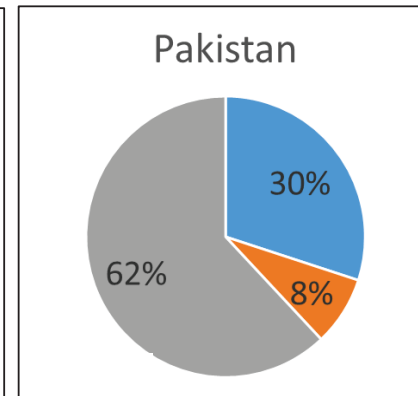
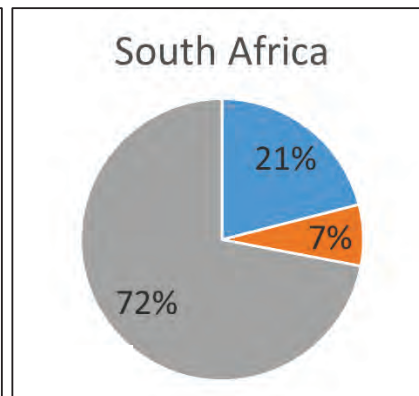
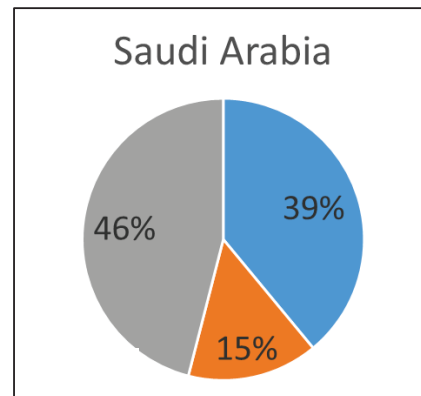
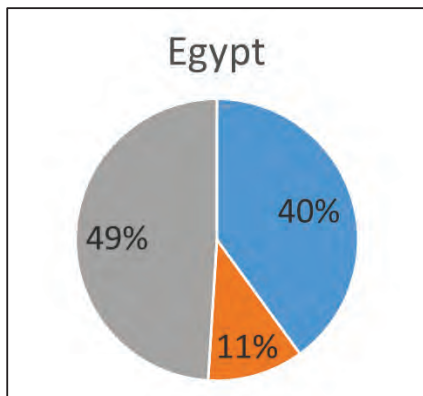
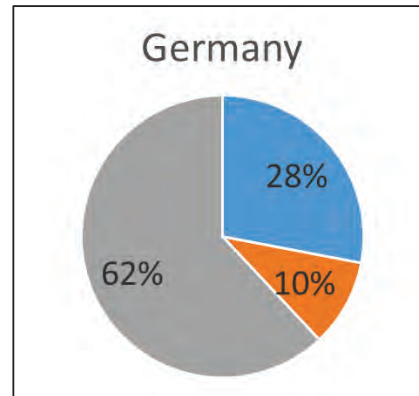
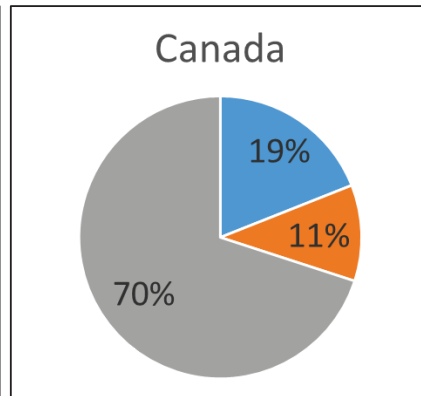
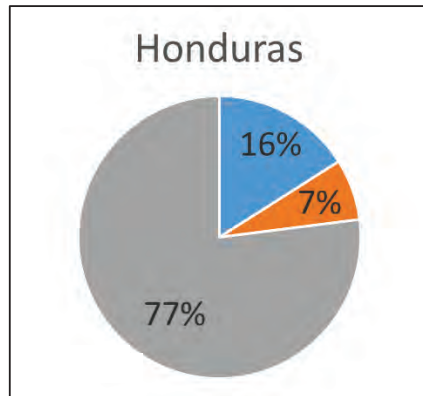
Honduras:

7229 specimens in >1700 species



=> 23% of all specimens can be covered by AI tools for 20 species

## For how many species do we need AI identification tools?





## How to get started?

### Natural environments:

Step 1: process 1000 specimens for 500 sites/sample (half a million specimens)  
=> discovery of 25,000 species

Step 2: image numerically important species and train AI tools  
=> AI tools for thousands of common species

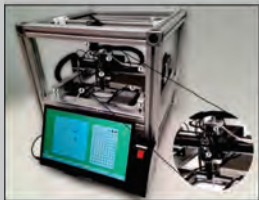
### Agricultural or forestry environments:

fewer important species => fewer specimens need to be processed

## Species Discovery Factory



**1. DiversityScanner:**  
Digitization with robots



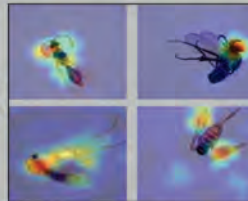
+

**2. Nanopore Sequencing:**  
Sorting with DNA barcodes



=

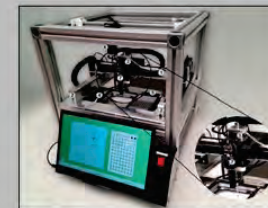
**3. Identified images:**  
biomonitoring with AI



## Species Biomonitoring Factory

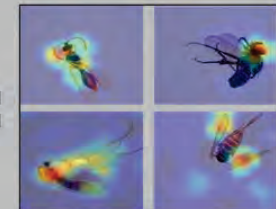


**1. DiversityScanner:**  
Digitization with robots



=

**2. Identified images:**  
biomonitoring with AI

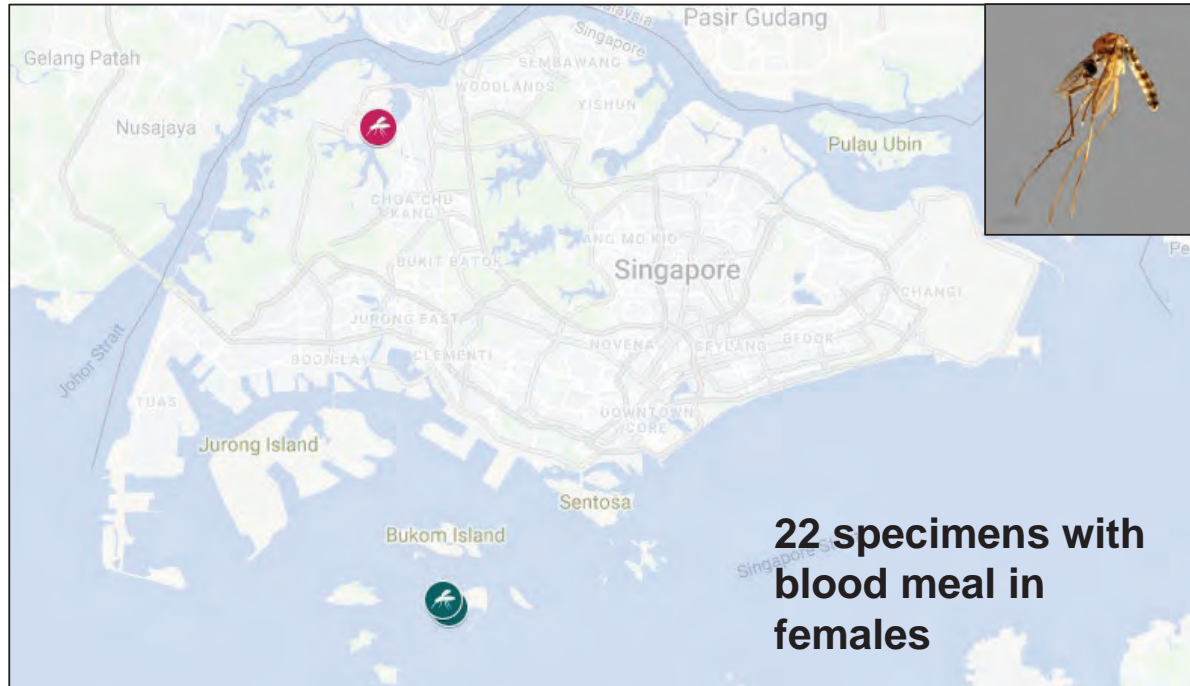


# Environmental impact of species: Molecular Ecology

- 4000 mosquitoes from Singapore processed
- => ca. 120 species detected
- => females for ca. 10 species had visible blood meals
- => metabarcoding of gut content

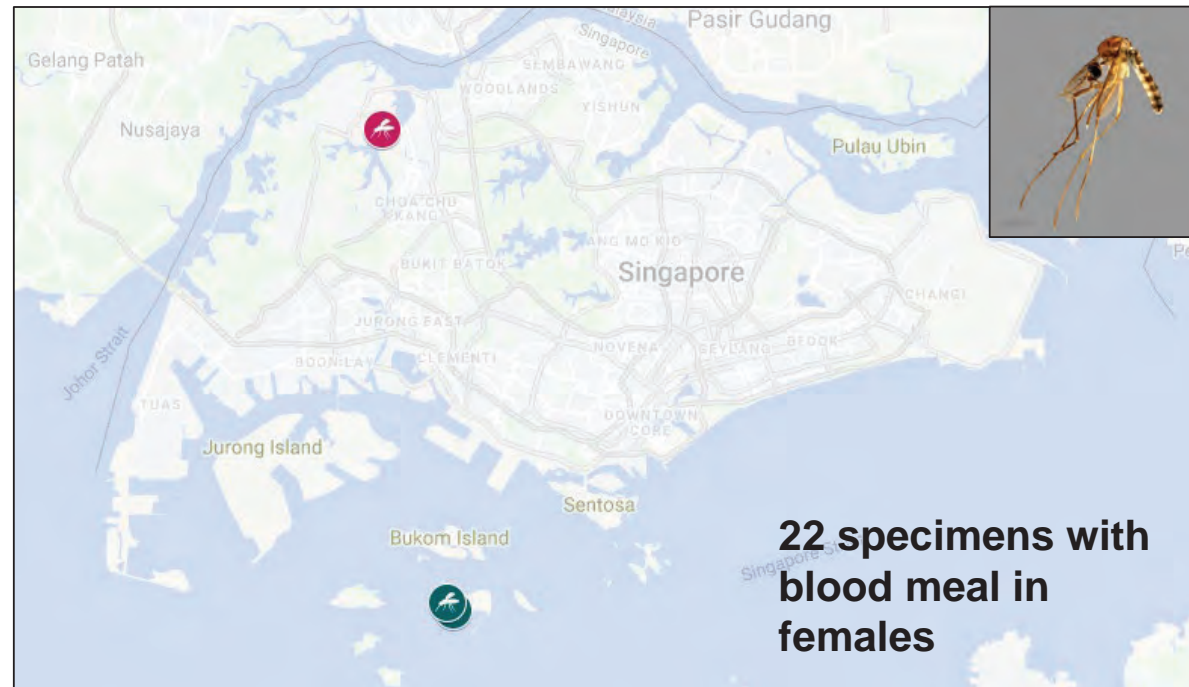


# *Aedes* sp. ZRCBDP00009685





## *Aedes* sp. ZRCBDP00009685



=> high potential as an agent transmitting viruses between distantly related hosts, but fortunately with restricted distribution

## *Uranotaenia* sp. ZRCBDP0009257





*Leong*  
*Tzi Ming*

# Conclusions

- The world expects us to monitor biodiversity holistically
  - A major challenge is not enough and biased data
  - Particularly data deficient are insects although they have high abundance and species diversity
  - Fortunately, automation and machine learning can overcome this challenge
- => cost-effective data generation tools for machines and humans are feasible