

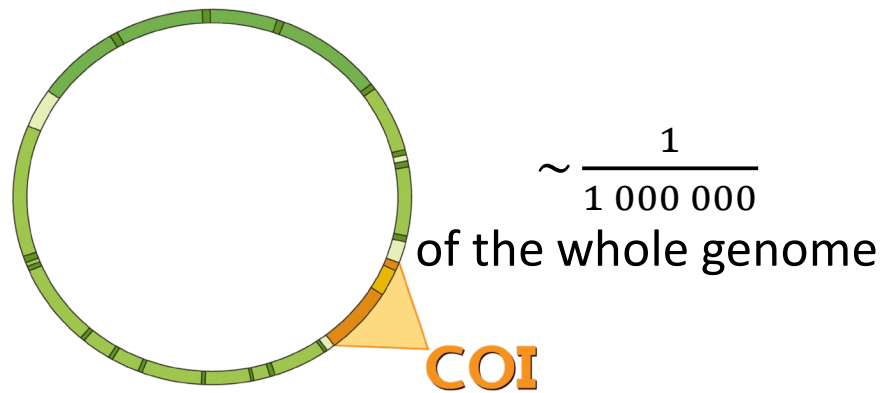
DNA Barcoding of  
Amazonian Plants and  
Insects:

Decoding Diversity and  
Interactions



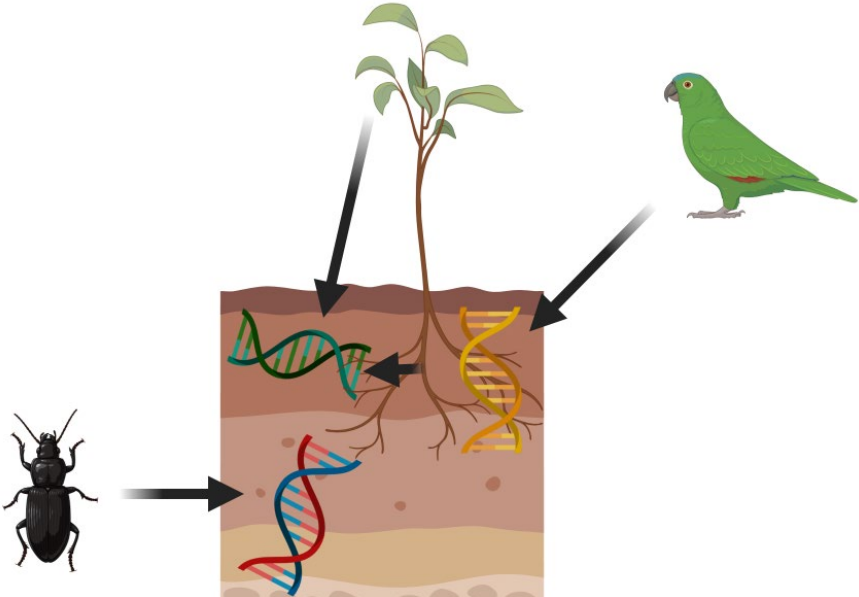


# DNA Barcoding: Introduction to Technology and Techniques

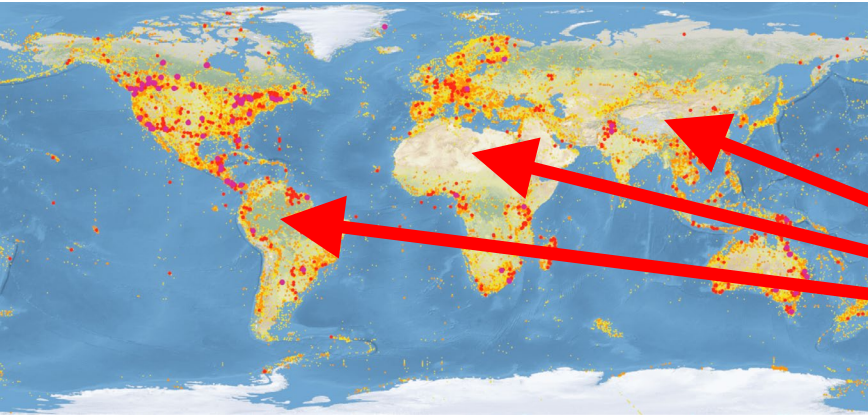
Vision: A world where we read the biosphere with DNA



658 bp of the mitochondrial DNA: COI

sampling strategies:	
	barcoding
	metabarcoding
	eDNA

# DNA Barcoding: Introduction to Technology and Techniques



internationally standardized protocols:

- 16M barcodes publicly available
- growth more than 3M barcodes / year
- challenge: specimen acquisition
- challenge: reference database required



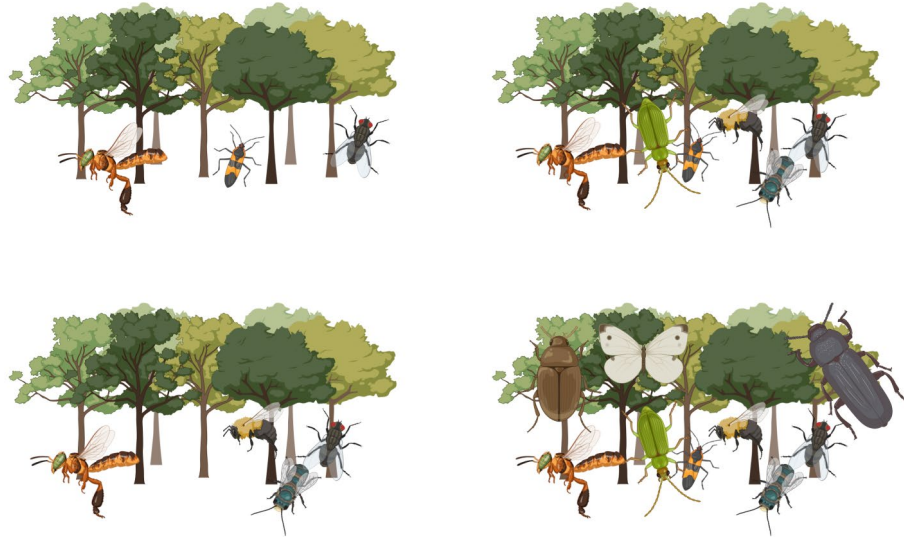
fast-paced technological advancements:

- portable DNA sequencer
- costs of sequencing / specimen < 0.1\$
- challenge: technical complexity

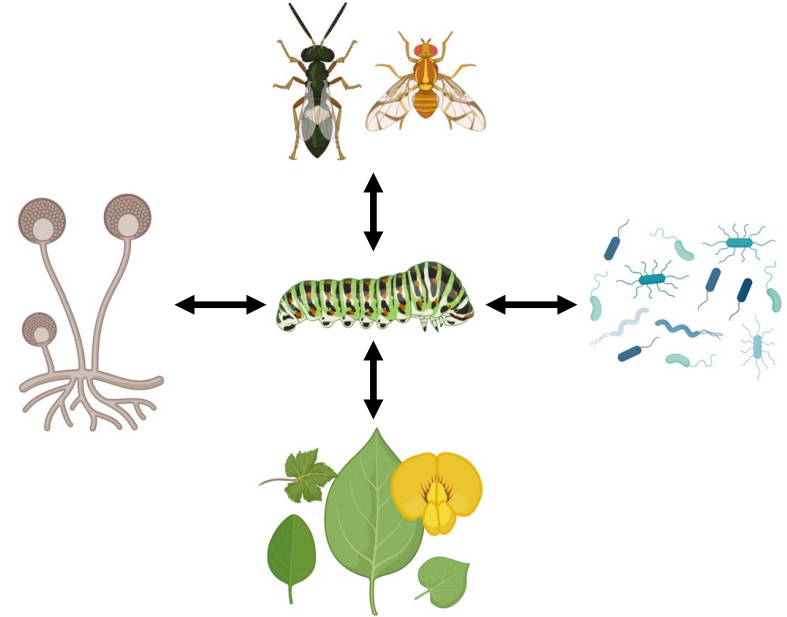
# DNA Barcoding: Potential Applications for Biodiversity Monitoring

## Biodiversity Baselines:

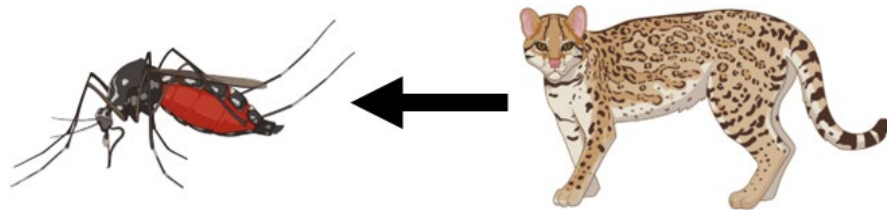
species inventories to identify priority areas



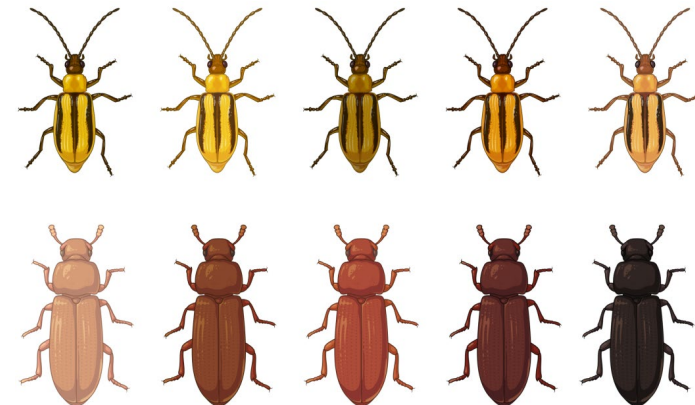
species interactions (e.g. parasitism)



elusive species: indirect/non-invasive detection  
(e.g. mosquito blood)



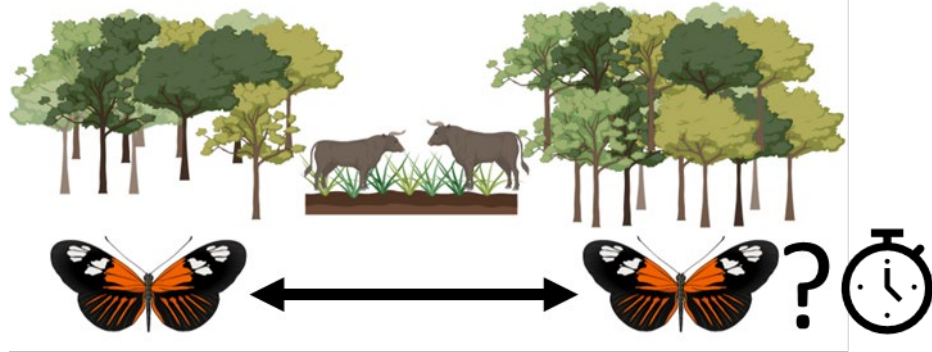
ecosystem resilience: taxonomic diversity



# DNA Barcoding: Potential Applications for Biodiversity Monitoring

## Temporal Variation:

connectivity: population loss by habitat fragmentation



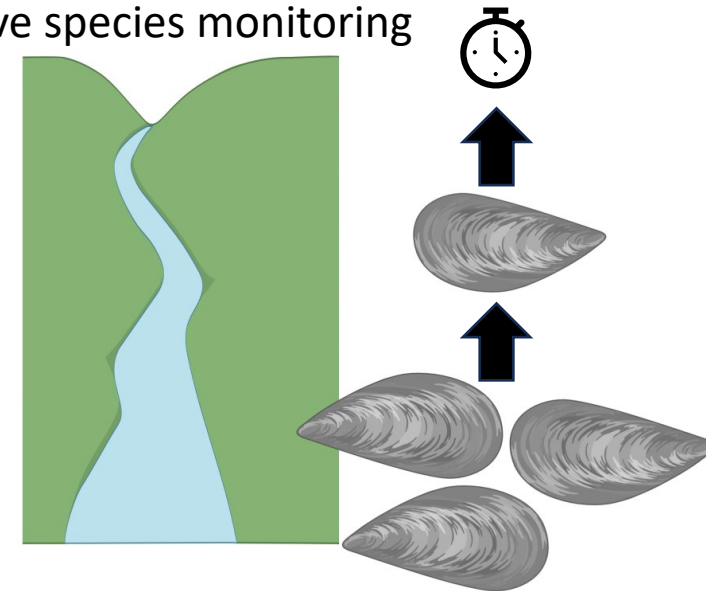
elevation shift due to climate change



impact of extreme weather events



invasive species monitoring



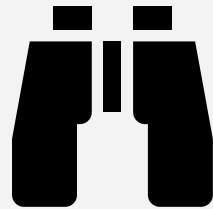
# DNA Barcoding: Case Studies and Practical Examples

## Biodiversity Inventory:

Malaise traps from 220m to 2900m



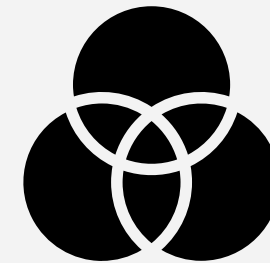
> 300 000 insects sorted, imaged & sequenced



~60% of BINs (species hypothesis) new to BOLD

>50% of BINs represented by single specimen

>80% of BINs by 3 or fewer specimens



<5% overlap: Manu Station and Wayquecha

<1% overlap: Wayquecha and Los Amigos

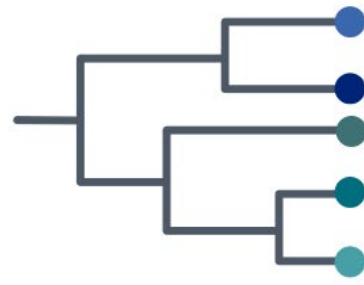
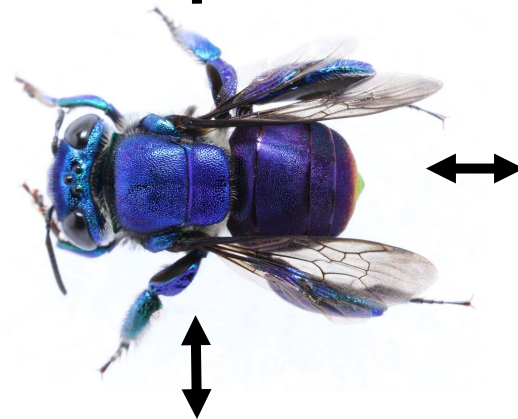
# DNA Barcoding: Case Studies and Practical Examples

## Euglossini (orchid bees):

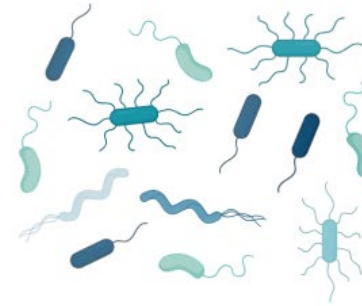
350m – 1450m, 2 years  
~ 2200 specimens



- transport mode of phytopathogenic fungi
- flower yeasts
- diversity discovery: 190 families in 80 orders



- phylogeny
- cryptic species
- potential as indicator species



- microbiome distinct from honey bees
- bacterial strain discovery



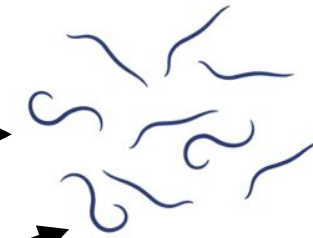
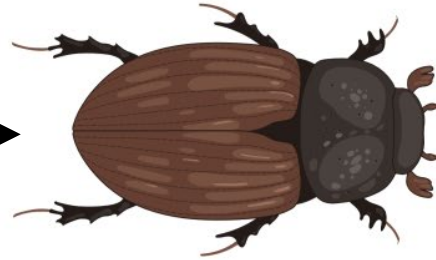
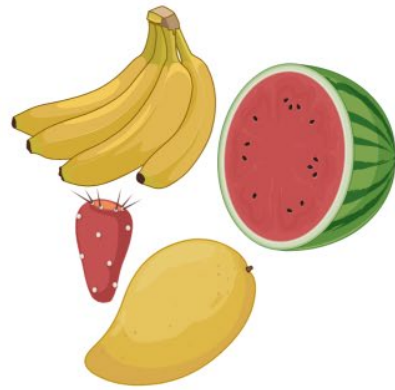
- approx. 300 plant genera, 90 families
- pollination networks
- plant distribution

# DNA Barcoding: Case Studies and Practical Examples

## Scarabaeinae (dung beetles):

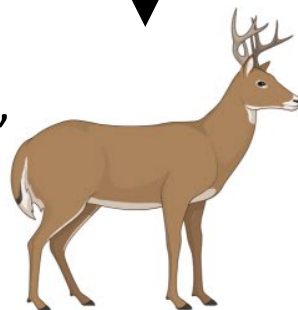
~ 1200 specimens

- fruits eaten by  
dung beetle  
and/or herbivore



- nematode  
species  
discovery  
- lifecycle or  
DNA traces?

- mammal/bird  
detection: Jaguarundi,  
Jaguar, Deer, Tapir,  
Paccari, Paca,  
Monkeys, Tinamous



- diet of herbivores





# DNA Barcoding: Case Studies and Practical Examples

