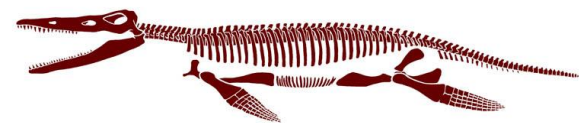


The advent of digital technology and its promise for biodiversity research

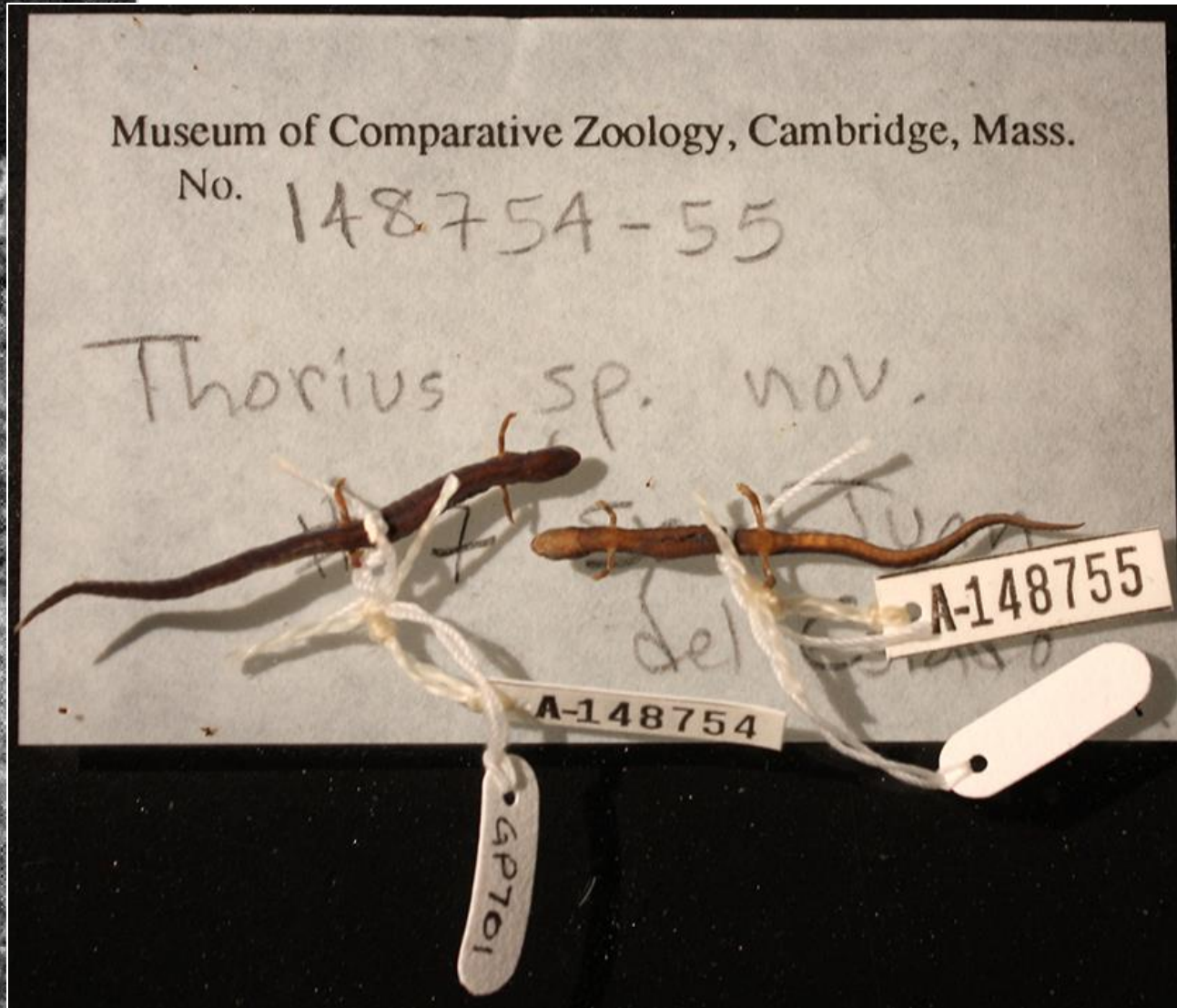
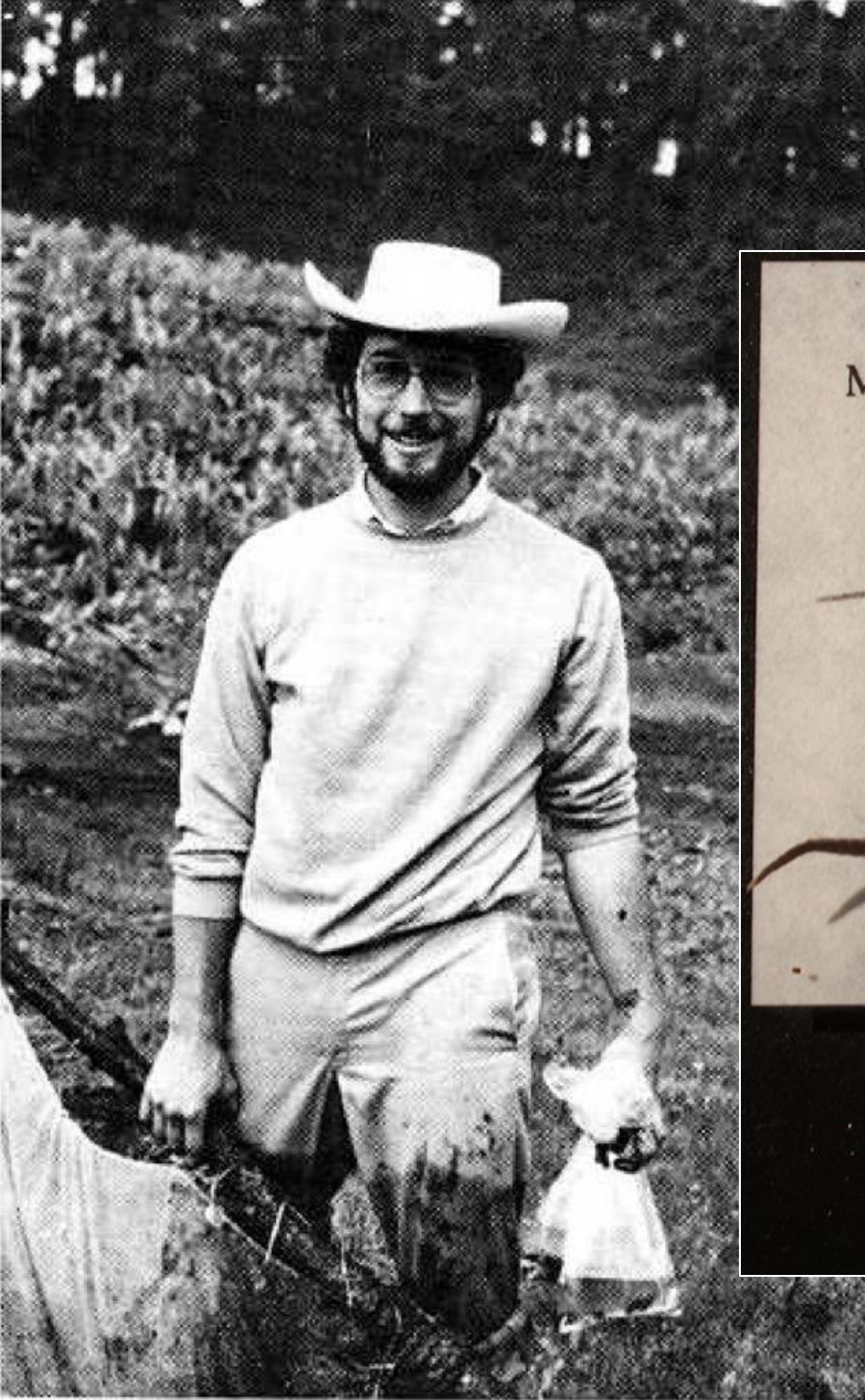
James Hanken

MUSEUM OF COMPARATIVE ZOOLOGY



HARVARD UNIVERSITY

ca. 1974



State of the Art, 1974

Veracruz	Thorius pennatulus narismagnus	Veracruz	Thorius pennatulus narismagnus
118142	Volcan San Martin, 3800'	8-1694	11
118143	Volcan San Martin, S slope, 3575'	85441	14
115164	S Sl Volcan San Martin Tuxtla, 3800'	88833	3
121134	SE slope Volcan San Martin, ca. 1100 m.	115163	2
121267	Volcan San Martin, SE slope, ca. 1190 m.		8
121268	Same as above, ca. 1005 m.		
121269	" " " " ca. 880 m.		
121270	" " " " ca. 1000 m.		
121271	" " " " ca. 1110 m.		

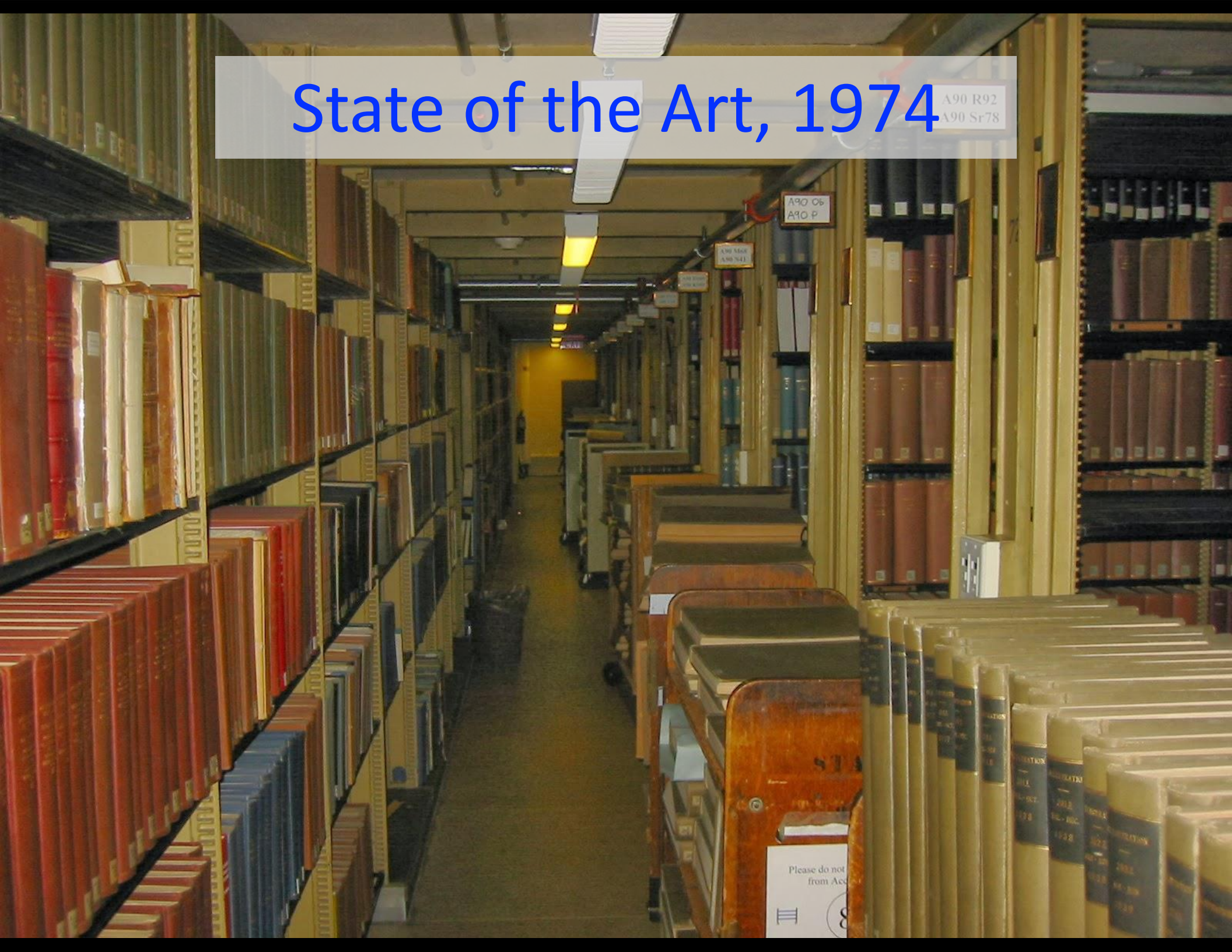
Oaxaca	Thorius sp.
126294	Ixtlan de Juarez, 13.1 mi NE, 9900'
126295	Cerro San Felipe, 10,500'
126296	Ixtlan de Juarez, 13.8 mi NE, 9800'
126297	Oaxaca, 14.1 mi NE and 2.7 mi E, 8900'
126845	Cofradia, ca. 8 mi. SW San Vicente Laohixio, ca. 9000'
132095	Llano de los Flores; 2700 m
132096-	"Cerro Pelon" SW slope, 13 mi. NE of Llano de los Flores (by road); 2700 m
132101	

Oaxaca	Thorius macdougalli
118913	Llano de las Flores, 12 mi. N of Ixtlan de Juarez, 9200'.
119705	Llano de las Flores, 2870 m. (S-2502-2505) 183

Oaxaca	Thorius narisovalis
119702	3.5 mi. S of Ranch el Pinto, 8350'. (S-2477) 5
119703	Rancho Mira Cumbre, 8600'. 2

Oaxaca	Thorius pulmonaris
117247	Cerro San Felipe
119701	3.5 mi. S of Rancho el Pinto, 8350'.
121886	ca. 10mi. NE of Ciudad Oaxaca

State of the Art, 1974



A90 C6
A90 P

A90 M6
A90 S41

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1-100 of 8,845

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Pro tip! Click on a table row to see the full occurrence details.



Identification	Taxonomy	Location	Recorded By	Sex	Date	Map	Media
MCZ Herp A-136428	Amphibia: <i>Thorius longicaudus</i>	Mexico, Oaxaca: 19 km S of Sola de Vega	James F. Lynch, David B. Wake, Theodore J. Papenfuss		1946-11-01		
MCZ Herp A-137376	Amphibia: <i>Thorius boreas</i>	Mexico, Oaxaca: Cerro Pelon, 50 km N Guelatao [VERBATIM ELEVATION:2850...	Gabriela Parra-Olea		1997-10-08		
MCZ Herp A-137378	Amphibia: <i>Thorius maddougalli</i>	Mexico, Oaxaca: 1 km E Cerro Machin	Gabriela Parra-Olea		1999-08-07		
MCZ Herp A-137380	Amphibia: <i>Thorius narisovalis</i>	Mexico, Oaxaca: Cerro San Felipe, 9.3 km W La Cumbre [VERBATIM ELEVATI...	Gabriela Parra-Olea		1999-08-04		
MCZ Herp A-137819	Amphibia: <i>Thorius longicaudus</i>	Mexico, Oaxaca: Mexico Hwy. 131, 19 km S (by road) Sola de Vega	James F. Lynch, David B. Wake, Theodore J. Papenfuss		1974-11-18		
MVZ Amphibian and reptile specimens 106399	Amphibia: <i>Thorius pennatulus</i>	Mexico, Veracruz: S-facing slope of River Canyon, 2 km N Teocelo, on r...	Collector(s): Samuel S. Sweet	undetermined	1972-12-19		
MVZ Amphibian and reptile specimens 106409	Amphibia: <i>Thorius pennatulus</i>	Mexico, Veracruz: Barranca, 2 km N of Teocelo	Collector(s): Theodore J. Papenfuss	undetermined	1972-12-19		

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Must have media
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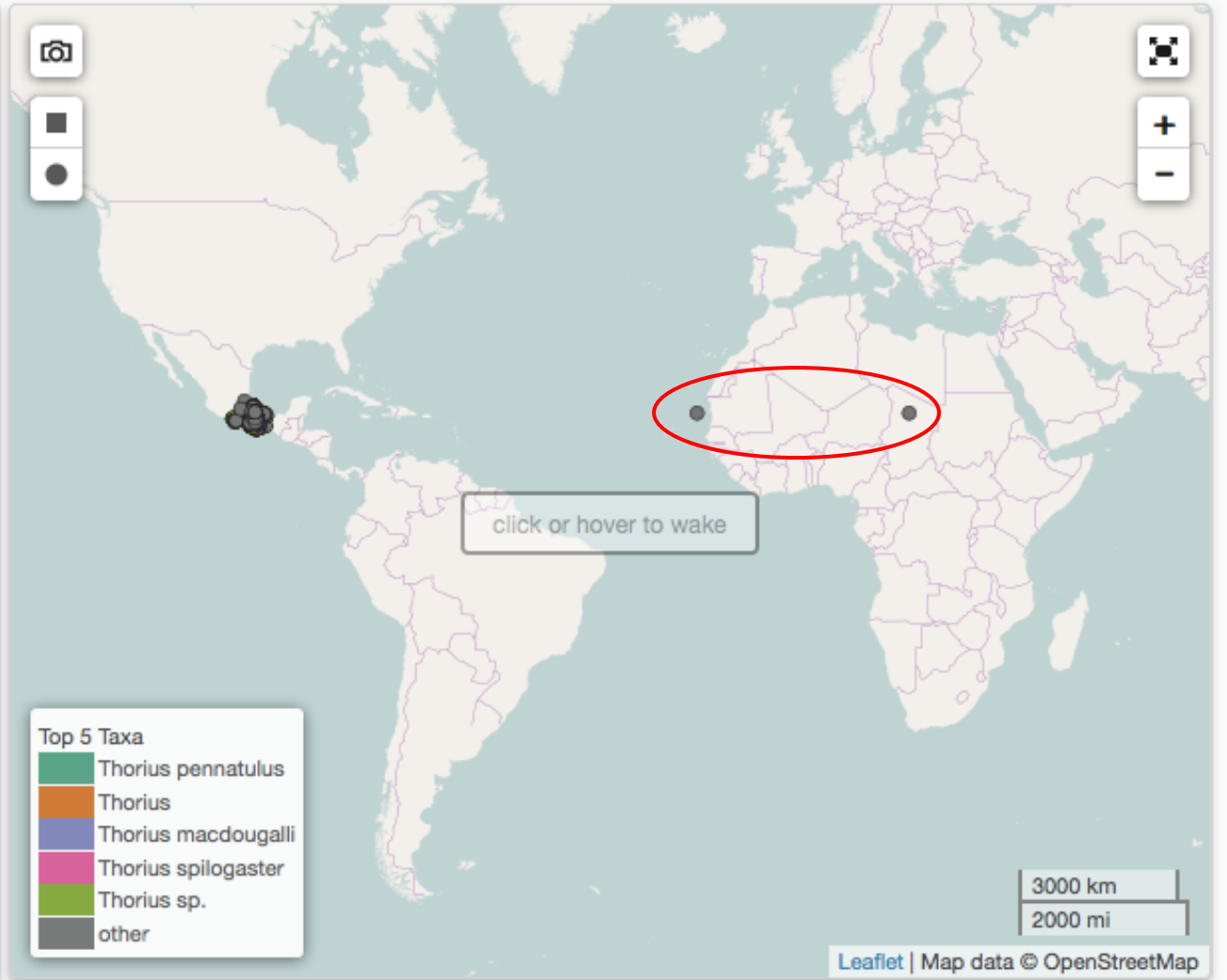
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Scientific Name [Add EOL Synonyms](#) ✕
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Date Collected Start: End: ✕
 Present Missing

Country ✕



Search Records

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thorius

Must have media

Must have map point

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Add a field [Clear](#)

Scientific Name

dwc:scientificName

[Add EOL Synonyms](#)

Present

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Date Collected

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End:

yyyy-mm-dd

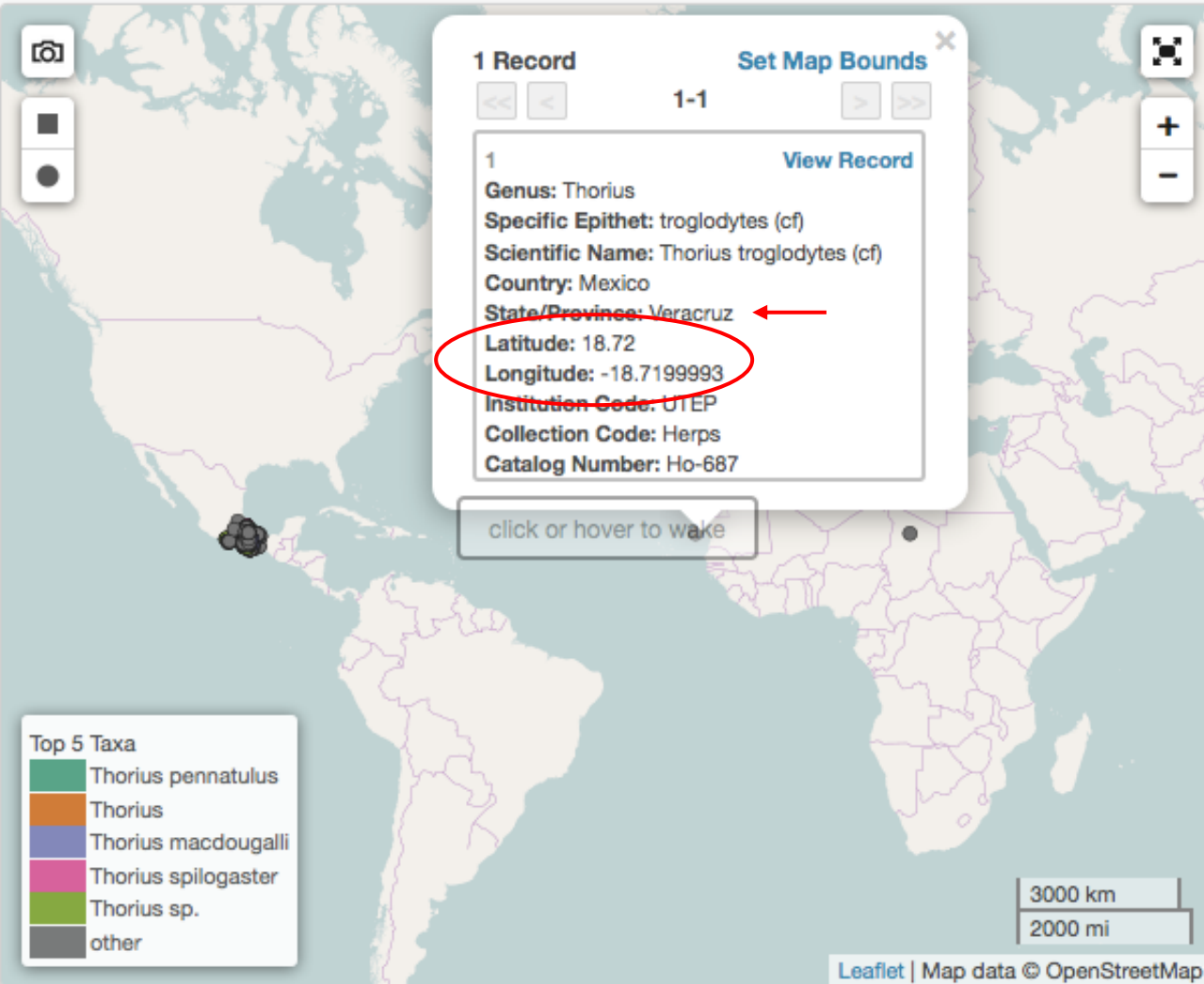
yyyy-mm-dd

Present

Missing

Country

dwc:country





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Biodiversity Heritage Library

*Stats as of February 2018

Proceedings of the Academy of Natural Sciences of Philadelphia.

A review of the species of Plethodontidae and Desmognathidae

v.21 (1869)

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Scientific Names on this Page

Page 111 (Text)

Plethodontidae 

Spelerpes

Thorius 

Indexed by  Global Names

entirely that of the Plethodontidae. This was the less to have been anticipated, as the general characters of the only genus are those of the genus Spelerpes. The history of the metamorphosis is as yet unknown. The only known genus is Mexican.

THORIUS Cope.

Parietal and palatine bones rudimental, represented by cartilage and membrane. Posterior nares therefore not separated from orbit; sphenoidal patches of teeth entirely united; tongue boletoid, free in front. Toes distinct, rudimental, 4—5.

The tarsal bones consist of astragalus, calcaneum, a scaphoid and three minute cuneiform bones. The metatarsals and phalanges are fully ossified, as are the corresponding elements of the fore limbs.

This genus is highly interesting, as indicating the lowest grade of ossific deposit found among the tailed Batrachians, accompanied by characters of full development in other respects. Thus, while the cranium is but imperfectly ossified, and less developed than in a comparatively early larval stage of Amblystoma, the tongue, vertebral column, and extremities have advanced far beyond its larval condition, which is permanent in the latter genus, and the branchial apparatus disappears while the individuals are but little more than half their adult size.

It is represented as yet by but one species, from Mexico, of terrestrial habits.

THORIUS PENNATULUS Cope.

American Naturalist, 1869, 222.

This is a small species, with smooth skin, very weak limbs, and stout tail. The head is scarcely wider than the neck; it is not flattened, the loreal region is rather elevated and distinct, and the muzzle slightly prominent. The upper lip is sometimes truncate, with infranarial angle prominent, sometimes regularly rounded. The nostril is larger than any known salamander, its diameter equalling half that of the pupil.

The vomerine teeth are situated on a transverse, elevated crest, which is a little behind between the inner nares, and though curved backwards, is but little interrupted medially. Each half contains four teeth, perhaps five when complete. The sphenoidal series is large, pyriform, the anterior extremity narrowed and prolonged to opposite the middle of the orbits.

There are thirteen costal folds; three and one-half of their interspaces are by the extended hind limb, from its origin. The toes are very

Biology of the of minute species (*Thorius*) from

Gabriela Parra-Olea¹,
Jessica A. Maisano⁴ D

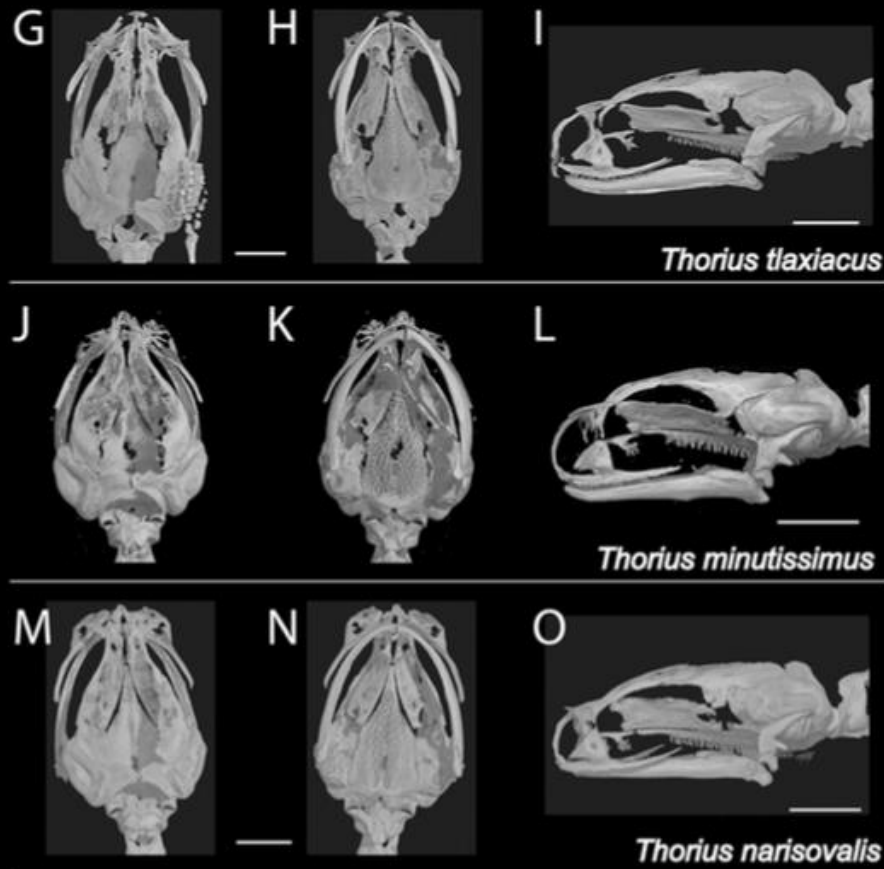


Figure 5 X-ray micro-computed tomography (μ CT) scans of adult skulls. (A–C) *Thorius pinicola*, MCZ A-136429, paratype, male; (D–F) *T. longicaudus*, MCZ A-137819, holotype, female; (G–I) *T. tlaxiacus*, MVZ 183447, paratype, male; (J–L) *T. minutissimus*, IBH 23011, female; and (M–O) *T. narisovalis*, MVZ 162257, female. Each skull is shown in dorsal (left), ventral (middle) and left lateral views. The skeleton of the right hand is visible in G. Total length of each skull is only 3–4 mm; scale bar, 1 mm.

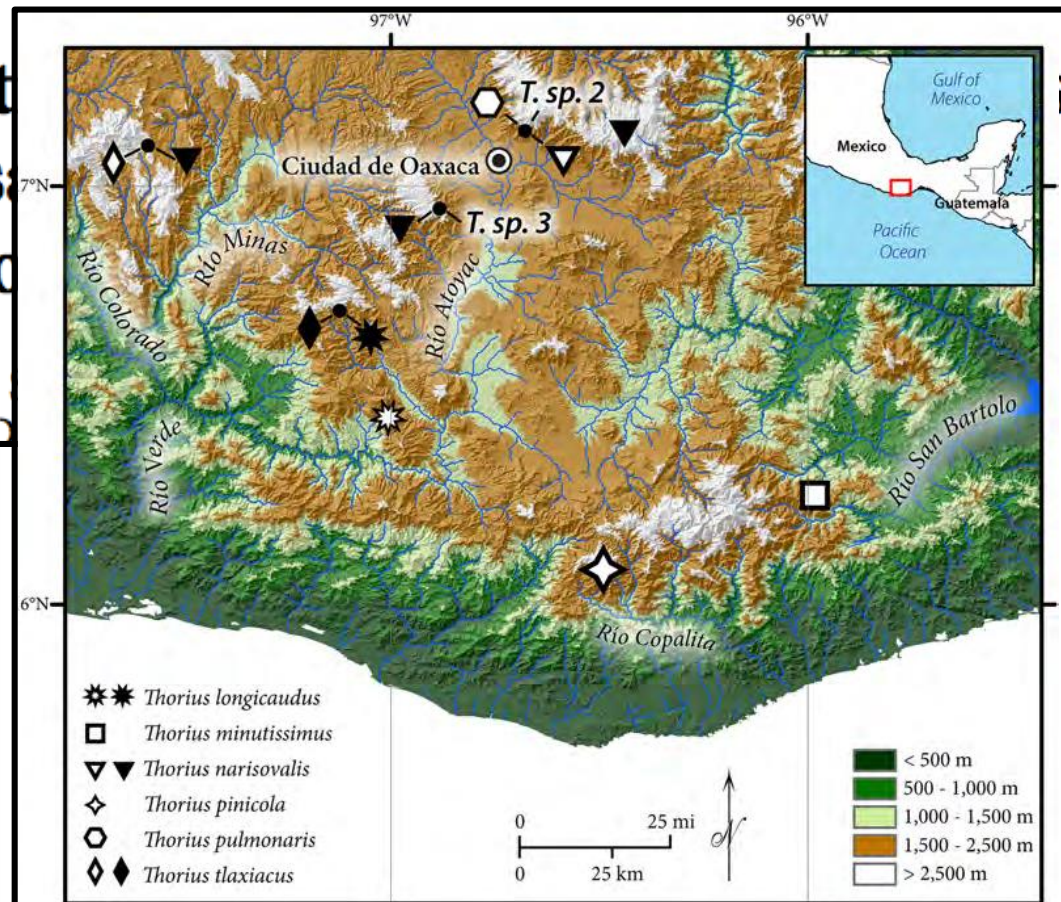


Figure 1 Geographic distribution of *Thorius* in western and southern Oaxaca, Mexico. Type localities of six named species are denoted by open symbols; additional localities are denoted by closed symbols. Known localities of *T. sp. 2* and *T. sp. 3*, two unnamed Oaxacan species, are also shown (Rovito *et al.*, 2013). Small closed circles denote four localities where two or three species are sympatric or nearly sympatric (from left to right): Heroica Ciudad de Tlaxiaco, San Vicente Lachixio, Zaachila and Cerro San Felipe.

g geographically disjunct populations. Adult *T. sp. nov.*, and *T. tlaxiacus* sp. nov. are larger than *T. minutissimus* sp. nov. rather than oval nostrils; *T. pinicola* and *T. narisovalis* sp. nov. differ from each other and from other named species by differences in adult body size, external morphology and internal anatomy (allozymes) and differences in DNA sequences.

What happened, and when?



ADBC →



ORNIS2 PORTAL LAUNCHES
The **ORNIS2 portal** was launched.



HERPNET2 PORTAL LAUNCHES
The **HerpNET2 portal** was launched.



FISHNET2 LAUNCHES
The **FishNet2 portal** was launched.



ORNIS FUNDED
ORNIS was funded by the National Science Foundation (NSF), DBI-0345448, 2004-2010.



FISHNET2 FUNDED
FishNet2 was funded in 2004 by the National Science Foundation (NSF), DBI-0415600, 2004-2010.



HERPNET FUNDED
HerpNET was funded by the National Science Foundation (NSF), DBI-0132303, 2002-2008



MANIS FUNDED
MaNIS was funded by the National Science Foundation (NSF), DBI-0108161, 2001-2006.



FISHNET FUNDED
FishNet was funded by the National Science Foundation (NSF), DEB-9910159, 1998-2001.

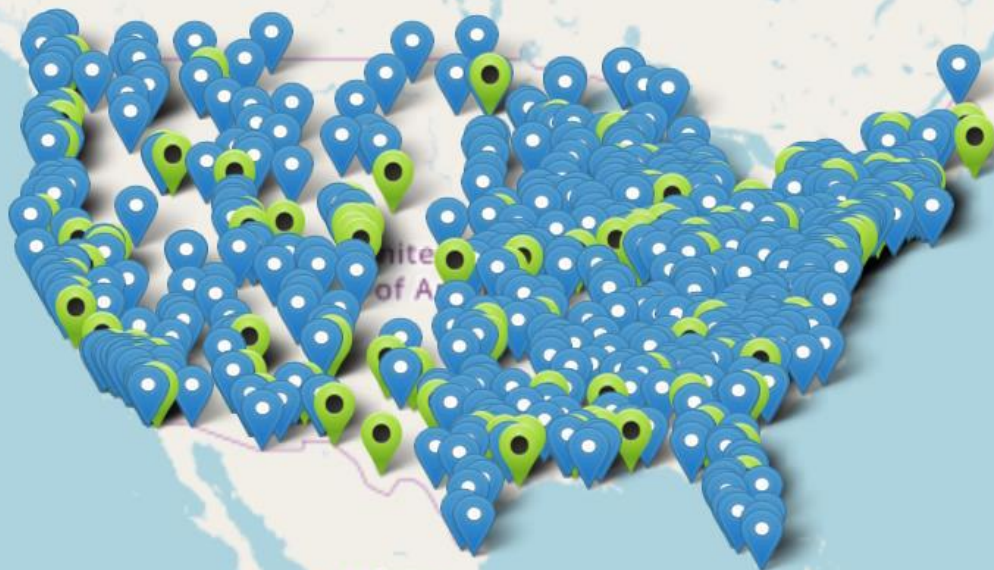
Advancing Digitization of Biodiversity Collections (ADBDC)

10 years @ \$100 million

April 2018:

- iDigBio is working with 708 collections in 396 institutions
- 20 TCNs and 17 PENs in 50 states
- 1 coordinating center (“Hub”): iDigBio

iDigBio Portal has
109M records for
327M specimens
with 23M
associated media
records



Blue = NSF-funded, green = other.

Why?

HARNESSING THE POWER of DIGITAL DATA

SCIENCE

EXECUTIVE OFFICE OF THE PRESIDENT
OFFICE OF SCIENCE AND TECHNOLOGY POLICY
WASHINGTON, D.C. 20502

Science Agencies

March 20, 2014

Scientific collections provide an essential base for developing scientific evidence and are an important resource for scientific research, education, and resource management.... Policies and procedures for maintaining, preserving, and developing Federal scientific collections while also increasing access to those collections for appropriate use are, therefore, central to their value.

Report of
to the Committee

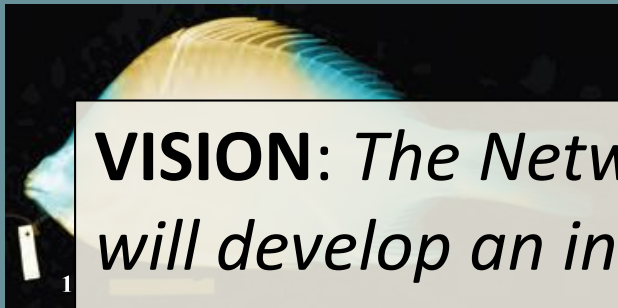
collections



National Science Foundation

NIBA Strategic Plan (2010)

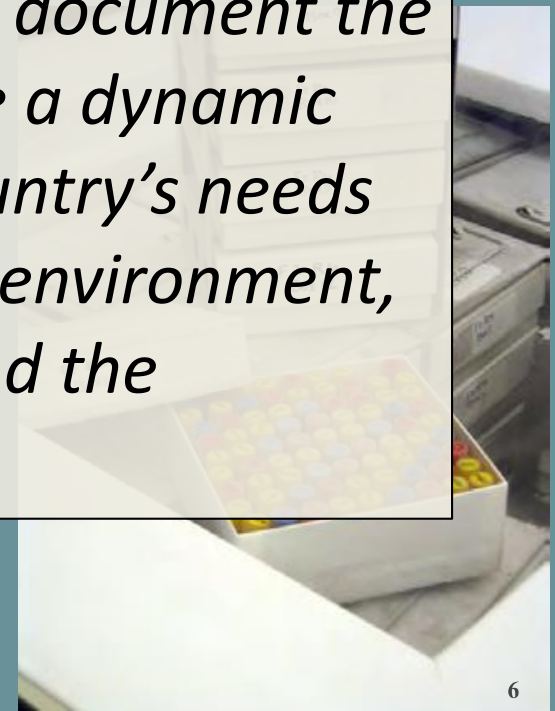
The Network Integrated Biocollections Alliance



VISION: *The Network Integrated Biocollections Alliance will develop an inclusive, vibrant, partnership of U.S. biological collections that collectively*

VISION: *The Network Integrated Biocollections Alliance will develop an inclusive, vibrant, partnership of U.S. biological collections that collectively will document the nation's biodiversity resources and create a dynamic electronic resource that will serve the country's needs in answering critical questions about the environment, human health, biosecurity, commerce, and the biological sciences.*

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IMPLEMENTATION PLAN FOR THE NETWORK INTEGRATED BIOCOLLECTIONS ALLIANCE

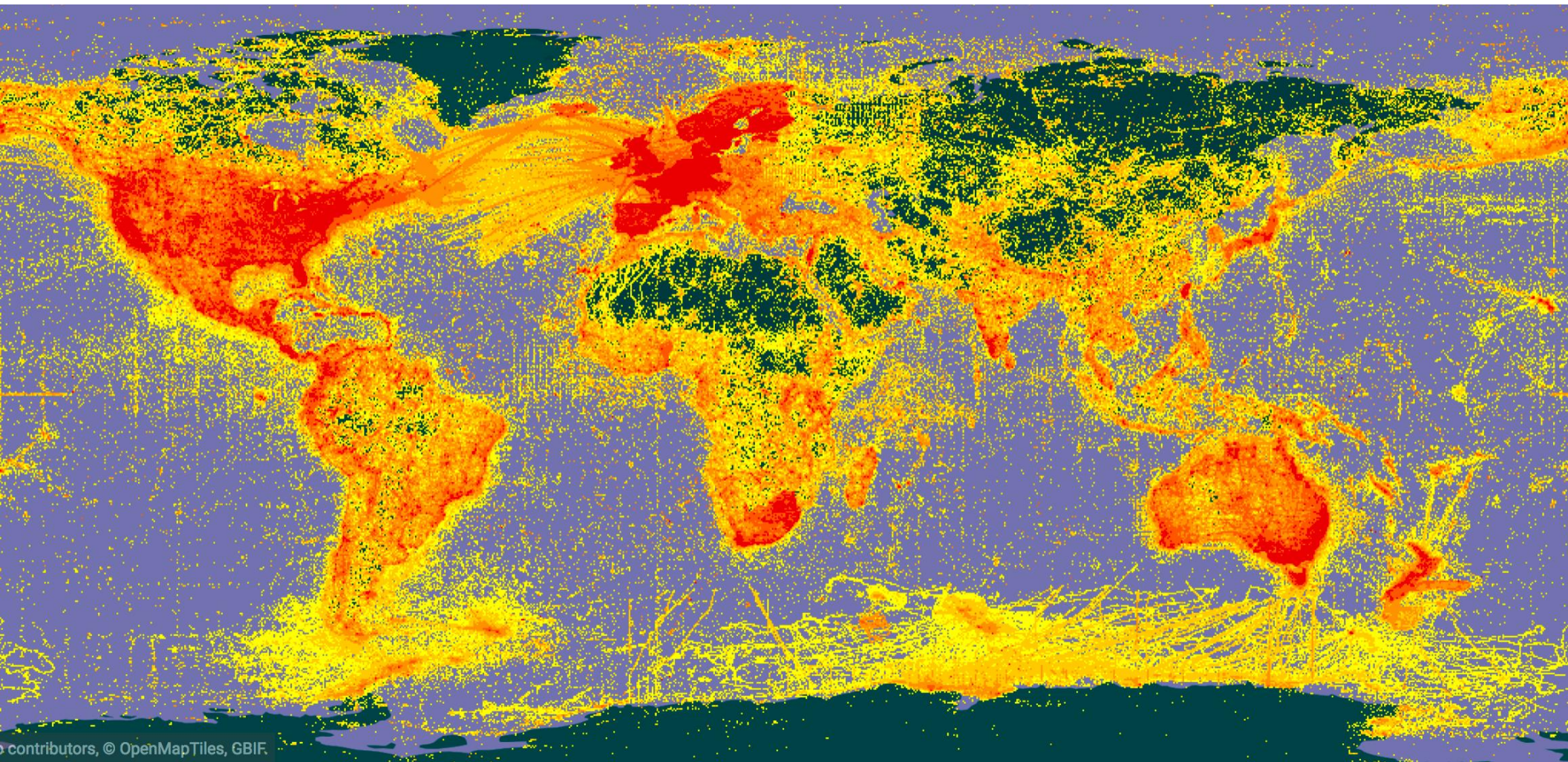


NIBA Implementation Plan (2013)

Digitization is a Global Effort

- Atlas of Living Australia
- CONABIO (Mexico)
- Canadensys (Canada)
- India Biodiversity Portal
- Swedish LifeWatch
- PORBIOTA (Azores, Portugal)
- Edaphobase (Germany)
- Ocean Biographic Information System

Global Biodiversity Information Facility



June 1st 2018: 984,441,947 occurrence records

Data Error is a Serious Problem

CellPress

Current Biology 25, R1057–R1069, November 16, 2015

Current Biology
Magazine

Correspondence Widespread mistaken identity in tropical plant collections

specimens had the wrong name prior to a recent taxonomic study. A similar pattern of wrongly named specimens is also shown for Dipterocarps and *Ipomoea* (morning glory). We also examine the number of available plant specimens worldwide. Our data demonstrate that, while the world's collections have more than doubled

(Supplemental Figure S1A). Finally, we documented the increase in the number of tropical herbaria and the accumulation of specimens for several taxa and geographical regions (Supplemental Figure S1B–C1–X).

Figure 1B charts the complete determination history of all names for all specimens of *Aframomum*. Before

“while the world’s [plant] collections have more than doubled since 1970, more than 50% of tropical specimens ... are likely to be incorrectly named. This finding has serious implications for the uncritical use of specimen data from natural history collections.”

analyses [1]. Thus, digitisation of natural history collections is providing unprecedented information to facilitate the study of the natural

over time relative to the current name, as determined in the recent monograph [2]. Additionally, we measured the inconsistency of

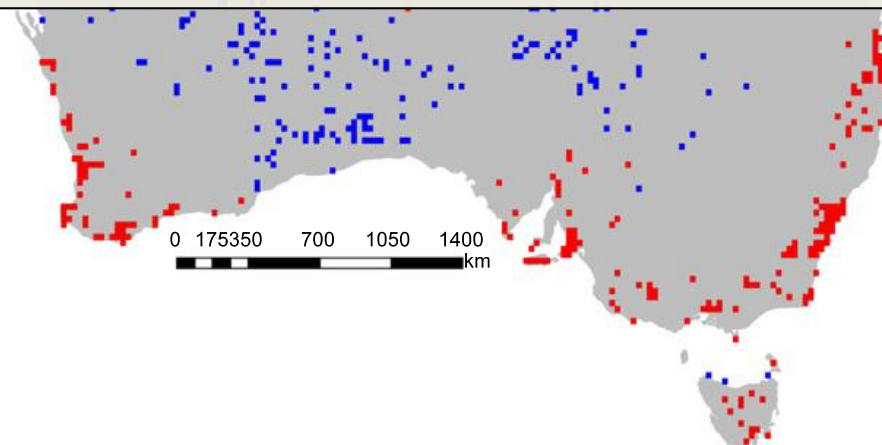
specimens of Dipterocarpaceae from nine herbaria, we identified 9,222 collections, each represented by at least two duplicate specimens

Widespread sampling biases in herbaria revealed from large-scale digitization

- “distance to herbaria explained 45% of the variance in collecting effort in AU, 29% in SA and 12.3% in NE, with a higher density of specimens closer to herbaria.”
- There was a significant trend toward collection on weekends (Saturdays and Sundays) in NE ... and midweek in SA and AU.

New Phytologist (2017)
doi: 10.1111/nph.14855

Key words: collector bias, geographic bias, herbarium, regional flora, sampling bias, temporal bias, trait bias.



Most Basic Digitization Remains To be Done

- 3–4 billion specimens are housed in natural history collections.
- Core data are online for < 50%.
- Many fewer specimens are georeferenced, imaged.
- Linkages among data types are typically limited and inconsistent.
- Special collections are largely untouched.

Building the future of Europe's Natural Science Collections



Our mission

To unify European natural sciences collections, effectively transforming a scattered and fragmented access model to a central facility with an integrated data-driven pan-European research infrastructure.

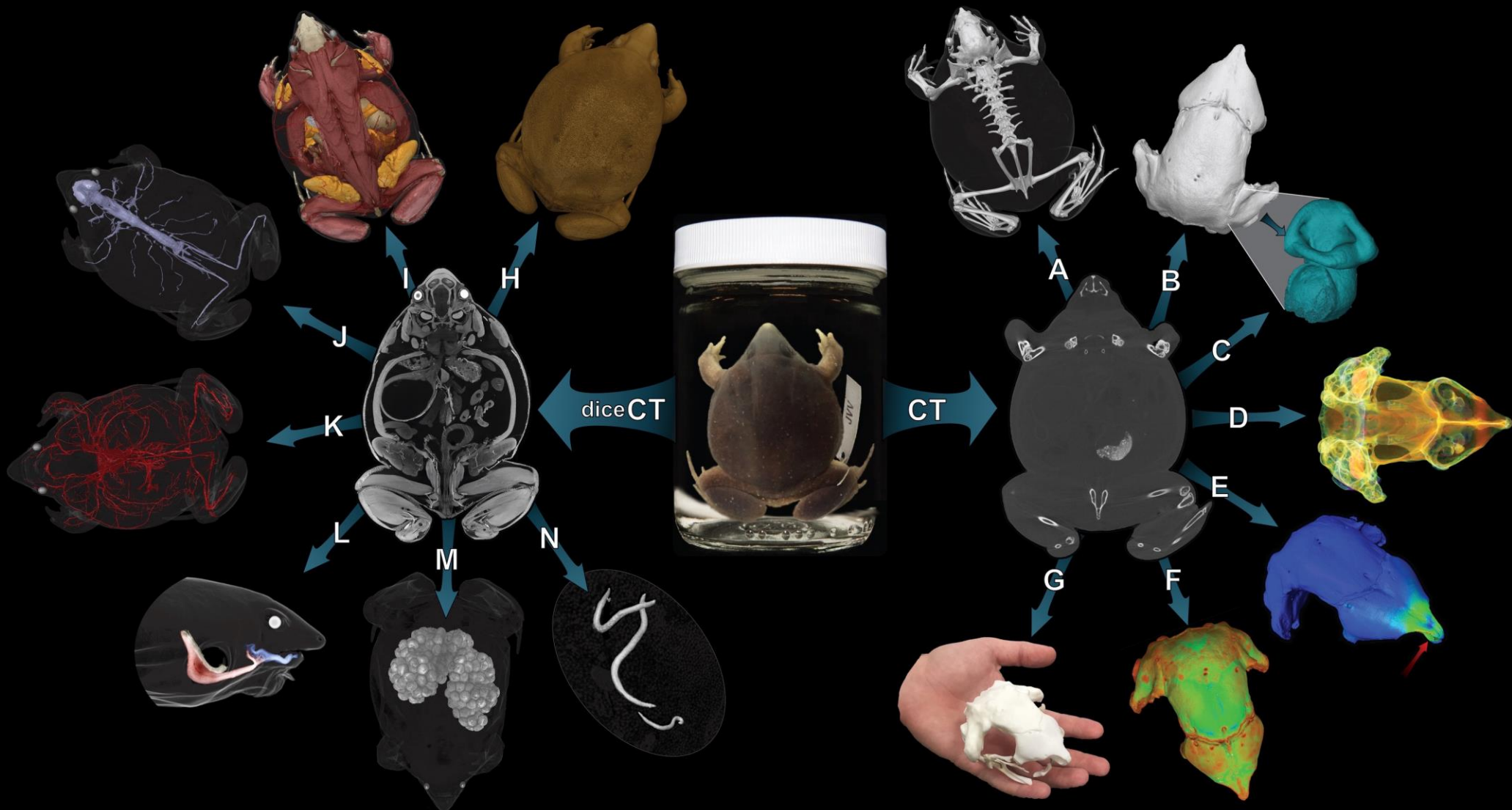
- 1.5 billion specimens held in European facilities
- 80% of global biodiversity described held by European collections
- 100 collaborative projects
- 5,000 scientists
- 15,000 visiting scientists annually
- 3,000 scientific publications annually
- 10 million public visitors annually
- 25 million web visitors annually

Distributed System of Scientific Collections (DISSCO)

- 114 European museums in 21 countries
- 1.5 billion specimens!
- 2018-2022: Preparatory phase (Innovation and Consolidation programmes), e.g., *European Open Science Cloud Summit*, 11 June 2018
- 2019-2024: Construction phase/programme
- 2024-2025: Deployment phase

oVert TCN

Microcomputed Tomography (μ CT)

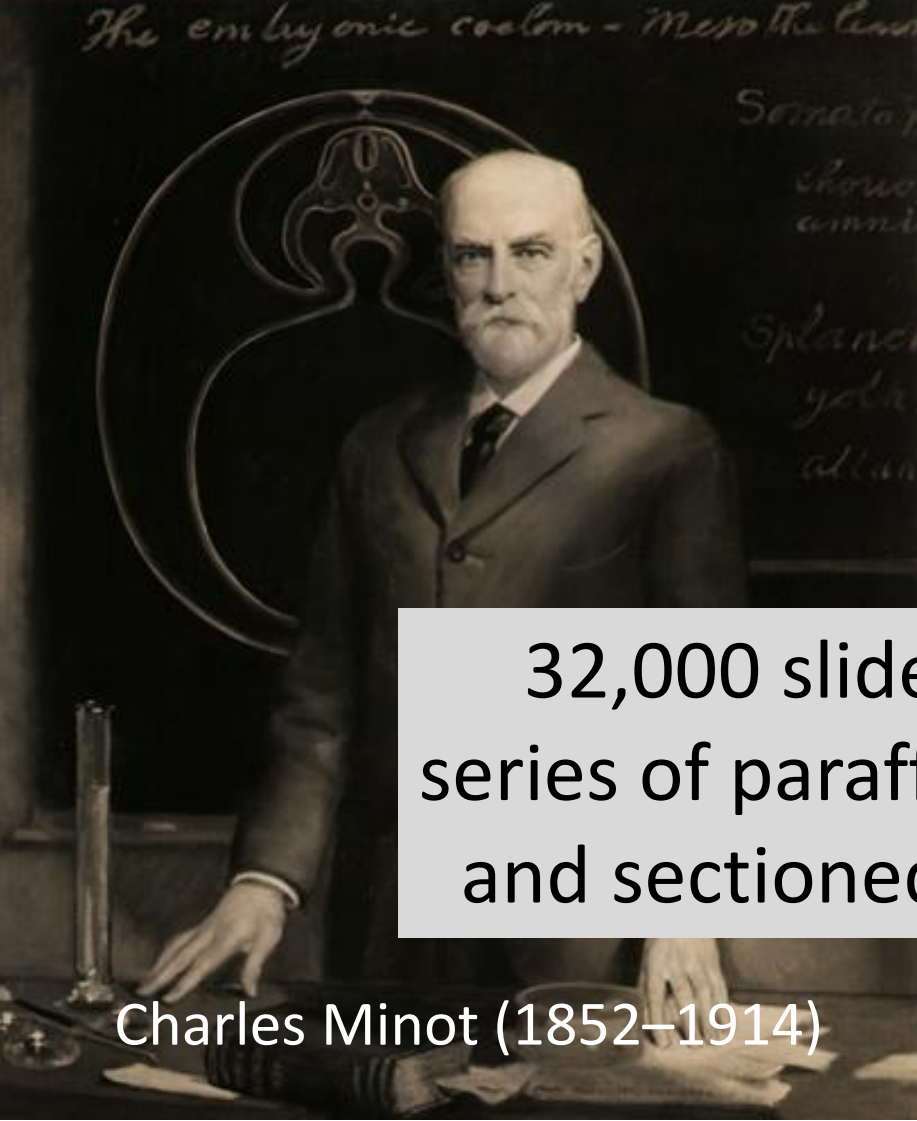


Scanning Microscope Slides

NO. 1999 SL. P
Amia calva 25.6 mm.
Frontal 10 μ
Cochineal, Alum
haematoxylum & Eosin
178-187
HARVARD
EMBRYOLOGY
A



- FMNH: ca. 400,000 slides, including 10,000 slides of type specimens.
- MCZ: > 32,000 slides of the Harvard Embryological Collection (Charles Minot).
- HUH: > 15,000 slides depicting wood anatomy, fungi, diatoms, insect pests, etc.
- NMNH: > 2,300 types/paratypes on slides and > 10,000 histological slides just for IZ.

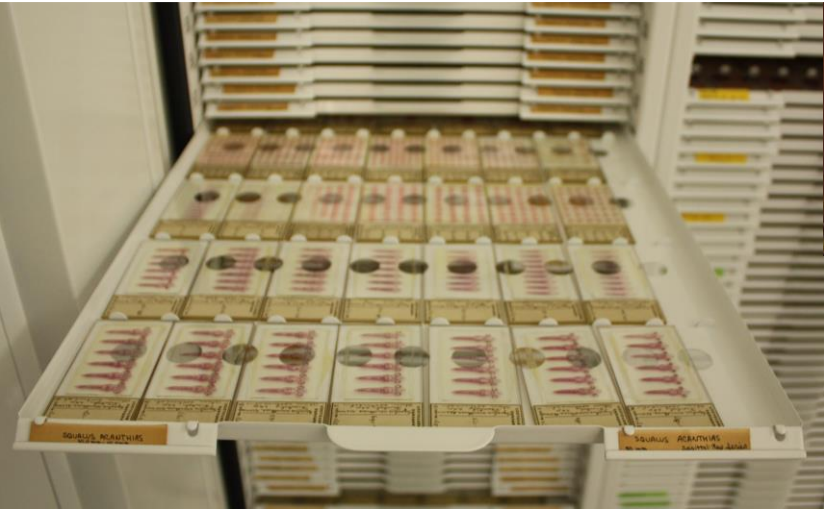


Charles Minot (1852–1914)



Harvard Embryological Collection

32,000 slides comprising 2,535 series of paraffin-embedded, stained and sectioned vertebrate embryos





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Research

Cite this article: Sanger TJ, Gredler ML, Cohn MJ. 2015 Resurrecting embryos of the tuatara, *Sphenodon punctatus*, to resolve vertebrate phallus evolution. *Biol. Lett.* **11**: 20150694. <http://dx.doi.org/10.1098/rsbl.2015.0694>

Received: 12 August 2015

Accepted: 30 September 2015

Subject Areas:

developmental biology, evolution

Evolutionary developmental biology

Resurrecting embryos of the tuatara, *Sphenodon punctatus*, to resolve vertebrate phallus evolution

Thomas J. Sanger¹, Marissa L. Gredler² and Martin J. Cohn^{1,2,3}

¹Department of Molecular Genetics and Microbiology, ²Department of Biology, and ³Howard Hughes Medical Institute, University of Florida, PO Box 103610, Gainesville, FL 32610, USA

The breadth of anatomical and functional diversity among amniote external genitalia has led to uncertainty about the evolutionary origins of the phallus. In several lineages, including the tuatara, *Sphenodon punctatus*, adults lack an intromittent phallus, raising the possibility that the amniote ancestor lacked external

BIOLOGICAL REVIEWS

Cambridge
Philosophical Society

Biol. Rev. (2015), pp. 000–000.
doi: 10.1111/brv.12187

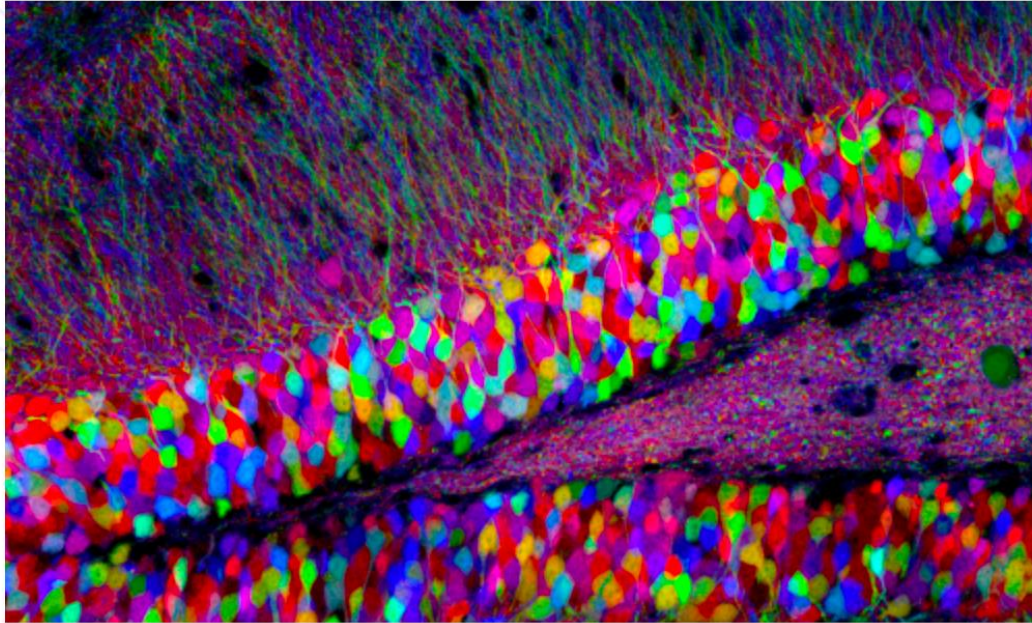
Fishing for jaws in early vertebrate evolution: a new hypothesis of mandibular confinement

Tetsuto Miyashita*

Department of Biological Sciences, University of Alberta, Edmonton, Alberta T6G 2E9, Canada



1 2 3 4 5 6



How the brain is wired

UPCOMING EVENTS

THURSDAY SEMINAR SERIES

Thursday Seminar Series

Gordon Berman (Emory)

Thu 18 Jan 4:00pm - Northwest Building Room 243

Assistant Professor

CBS SEMINAR

Developmental origin of neural circuits in *Drosophila*

Christopher Doe (University of Oregon)

Tue 6 Feb noon - Northwest Building

SlideAtlas

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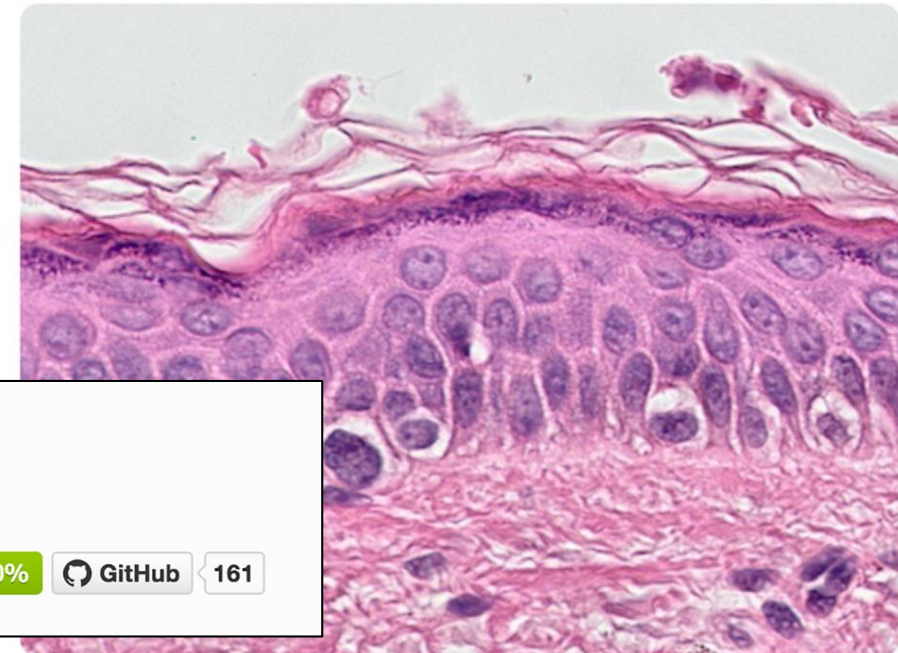
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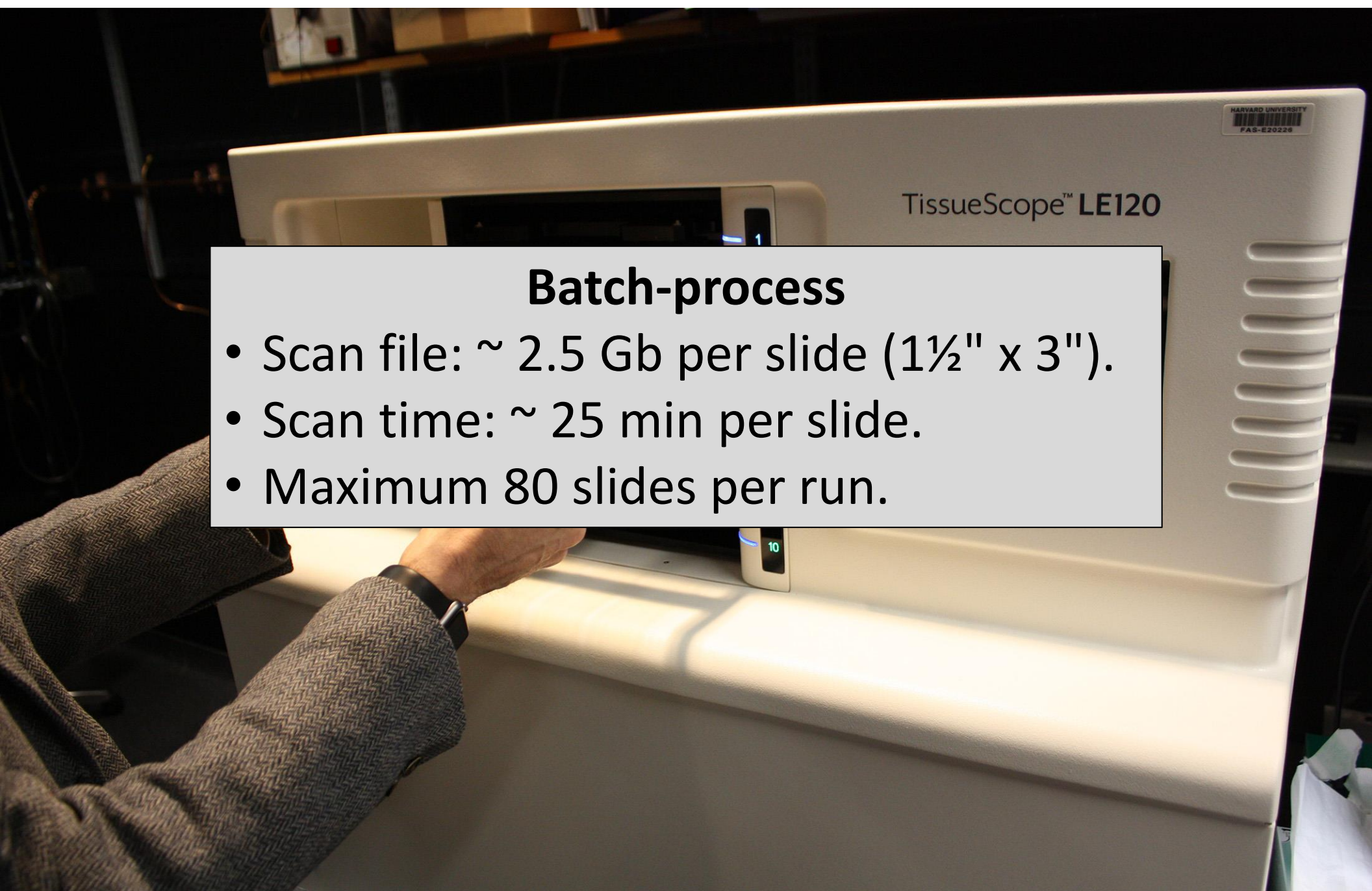
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coverage 90%

GitHub

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- Scan time: ~ 25 min per slide.
- Maximum 80 slides per run.

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MCZ / MCZ SC-1995 / HEC-1606 (free-tailed bat)


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Media Details
HEC-1606 slide J

Ultimate Goal: Linking Research to Collections*

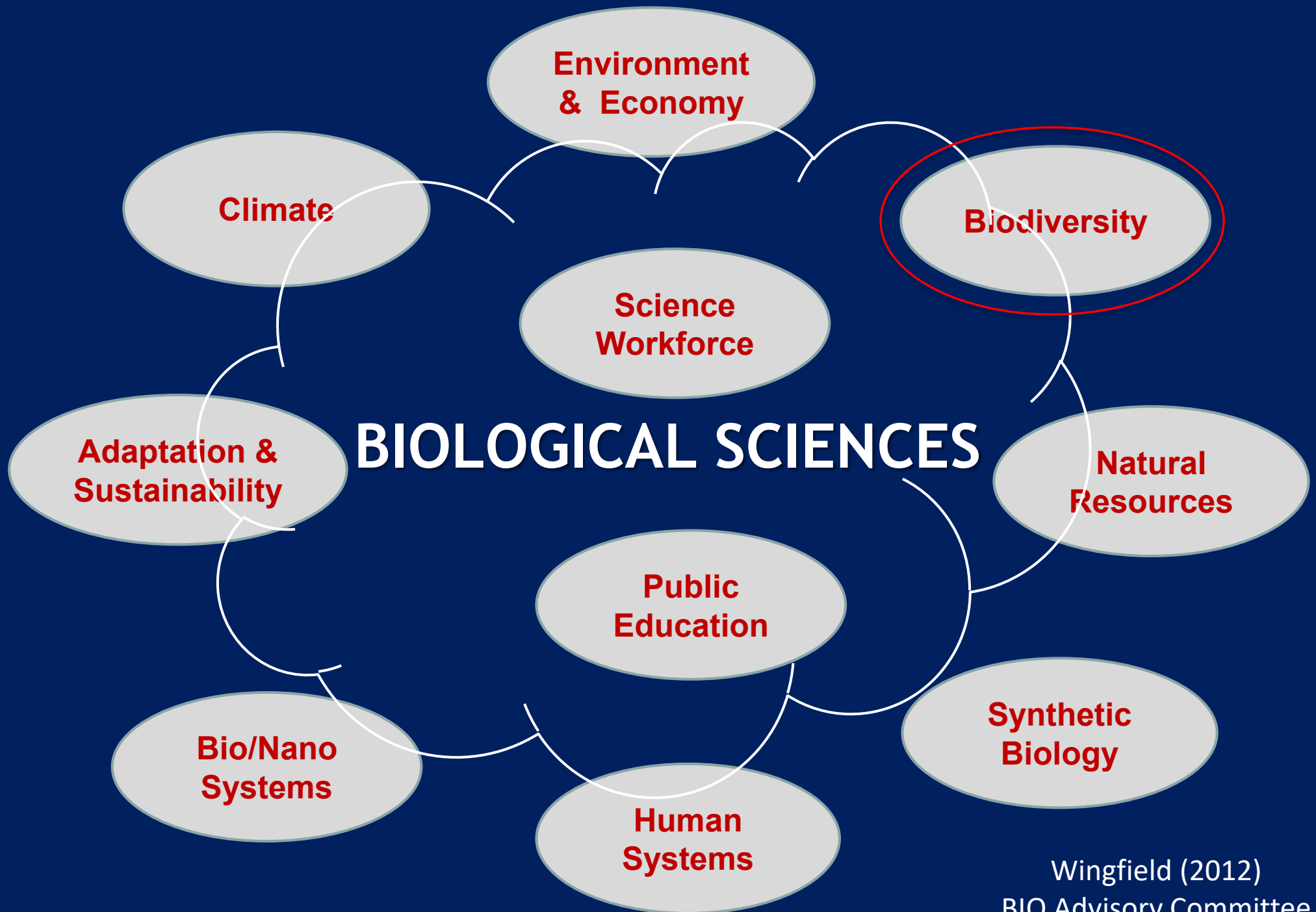
The screenshot shows the NSF website interface. At the top, there is a navigation bar with links: Home, Funding, Awards, Discoveries, News, Publications, Statistics, About, and FastLane. Below this is the NSF logo and the text "National Science Foundation Directorate for Biological Sciences (BIO)". A search bar is visible on the right. A secondary navigation bar includes: BIO Home, BIO Funding, BIO Awards, BIO Discoveries, BIO News, and About BIO. On the left side, there is a sidebar for "Molecular and Cellular Biosciences (MCB)" with various sub-links. The main content area features a "Press Release 12-090" titled "Cellular Secrets of Plant Fatty Acid Production Understood". The sub-headline reads "Chalcone-isomerase protein holds much promise of economic benefit". Below the text is an image of a hand holding a blue protein structure over a field of yellow flowers, with a red fuel can and a basket of fruit nearby. The NSF logo is in the top right corner of the image. Below the image, the text states "Basic discovery is key to rapid economic development." and provides a link for "Credit and Larger Version".

* Wingfield (2012)
BIO Advisory Committee



Protein \Rightarrow alignment \Rightarrow phylogenetics \Rightarrow sequence \Rightarrow GenBank \Rightarrow **vouchered specimen** \Rightarrow museum

Challenges Facing Humanity in the Next 50 Years



“e-research infrastructure for biodiversity knowledge generation”

PHILOSOPHICAL
TRANSACTIONS B

Biodiversity analysis in the digital era

rstb.royalsocietypublishing.org

Opinion

Cite this article as

Moritz C.

digital era

20150337

<http://dx.doi.org/10.1098/rstb.2015.0337>

“integration of various taxon-level data types (genome, morphology, distribution and species interactions) within a phylogenetic and environmental framework.... Living in this future will require the adoption of new ways of integrating scientific knowledge into societal decision making.”

CT 2601,

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Accepted: 2 February 2016

One contribution of 16 to a theme issue

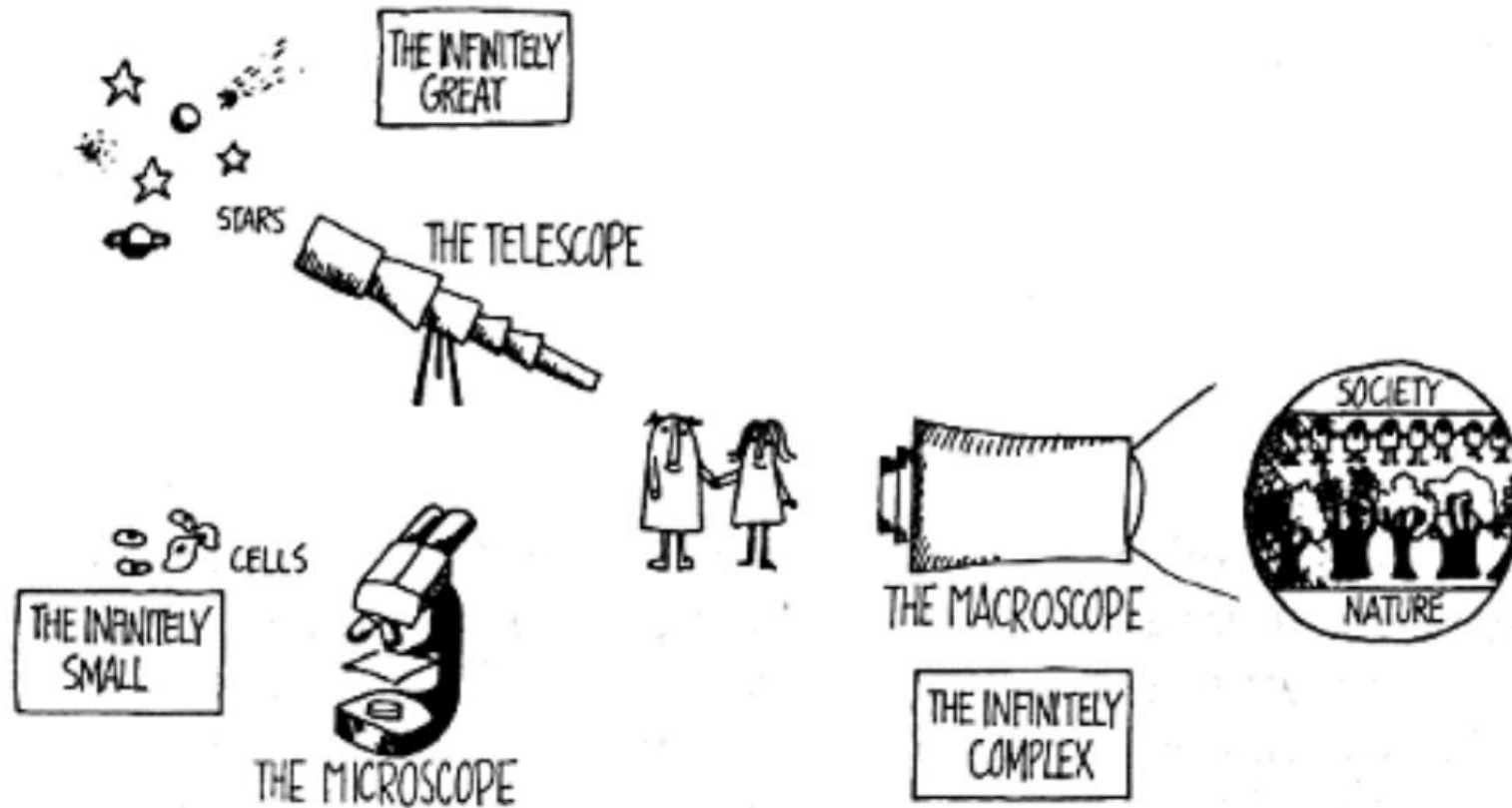
‘From DNA barcodes to biomes’.

Subject Areas:

data scaling problem in ecology, this integrative framework will provide richer information and fast learning to enable a deeper understanding of biodiversity evolution and dynamics in a rapidly changing world. The Atlas of Living Australia is used as one example of the advantages of progressing towards this future. Living in this future will require the adoption of new ways of integrating scientific knowledge into societal decision making.

This article is part of the themed issue ‘From DNA barcodes to biomes’.

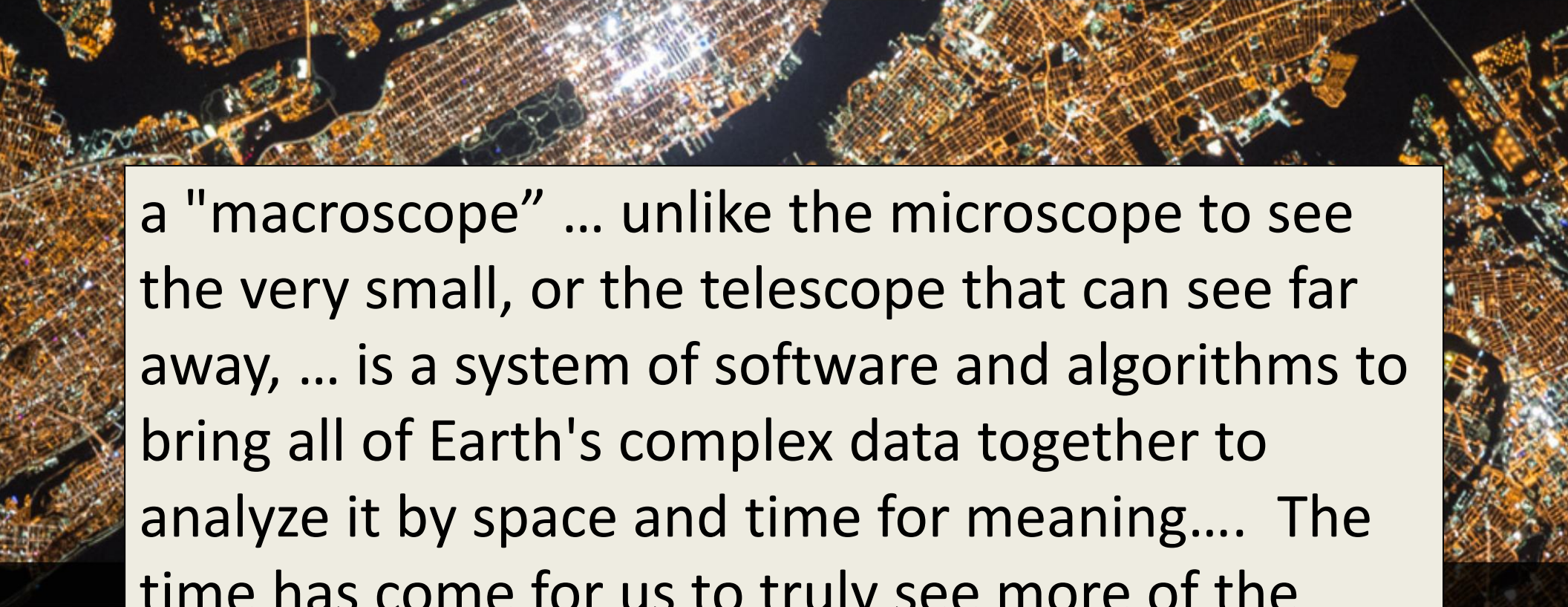
The Macroscope



The macroscope can be considered the symbol of a new way of seeing, understanding, and acting.

Joël de Rosnay (1975)

Macroscopes will help us understand Earth's complexity in infinite detail



a "macroscope" ... unlike the microscope to see the very small, or the telescope that can see far away, ... is a system of software and algorithms to bring all of Earth's complex data together to analyze it by space and time for meaning.... The time has come for us to truly see more of the world thanks to being able to digitize and collect new sources of data from millions of connected objects....

In

org
dat

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We call this a "macroscope" – but unlike the microscope to see the very small, or the telescope that can see far away, it is a system of software and algorithms to bring all of Earth's complex data together to analyze it by space and time for meaning.

FLORISTICS

An integrated assessment of the vascular plant species of the Americas

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wonders don't cease soon.”

—Alexander von Humboldt (1)

Fernández de Oviedo's chronicles (2) from
1526 contain the first European accounts

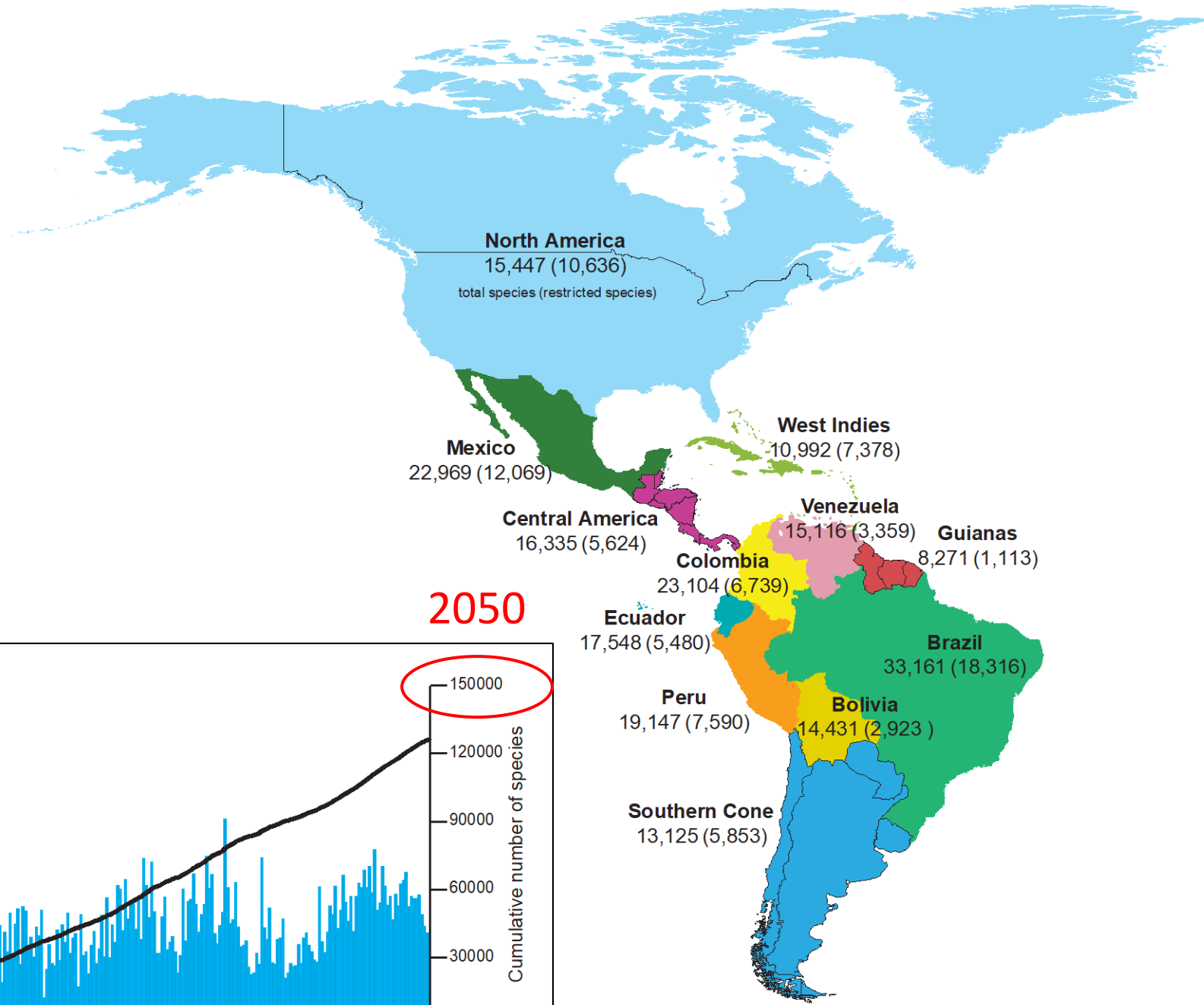
We present an integrated assessment of all known native species of vascular plants in the Americas. Twelve regional and national checklists ... were merged into a single list. Our publicly searchable checklist includes 124,993 species..., which correspond to 33% of the 383,671 vascular plant species known worldwide.

expert knowledge of plant specialists. The task of preparing authoritative checklists often proved more difficult than anticipated, in view of the large amounts of data to be synthesized within them, and some of the projects took a decade or more to complete. Checklists are extensively used

larger than the 77,100 species recorded for Africa (23), which has an area twice its size. Although China has a surface area of similar size to the United States or Canada, its mostly temperate flora consisting of 30,426 species (24) is 97% larger than the flora of the United States and Canada combined.

Within the Americas, Brazil has the most diverse flora, with 33,161 species, followed by Colombia (23,104) and Mexico (22,969) (Fig. 1 and

hidaceae, eae with fig. S2A); s (figs. S2 laceae is Andean of those bia 15%, s also the 13%) and t diverse he most- te South- d Mexico in most America.razil (8%) (8%), the enezuela se family is Rubiaceae (7%), which is well represented in all tropical regions; second in diversity are Orchidaceae and Asteraceae (table S1). Fifty-two families are endemic or near endemic to the Americas, including the nearly endemic Berberidopsidaceae,



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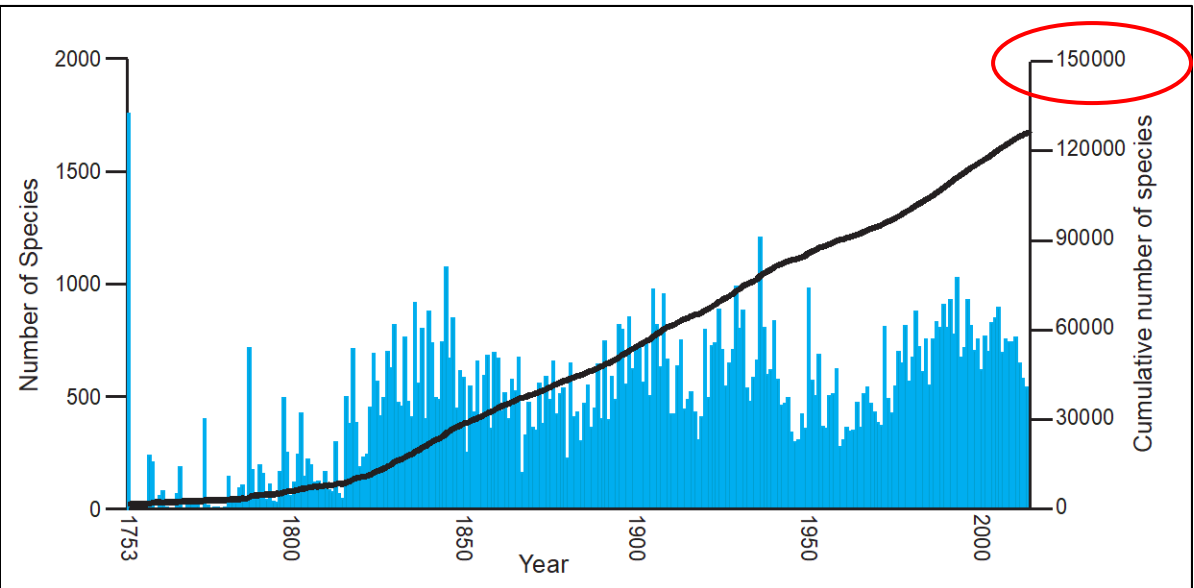


Fig. 4. Species described per year. The number of plant species (basinonyms) described per year from 1753 to 2015 for the Americas list (blue bars), and the cumulative number of accepted species (black line).

The biomass distribution on Earth

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A census of the biomass on Earth is key for understanding the structure and dynamics of the biosphere. However, a global, quantitative view of how the biomass of different taxa compare with one another is still lacking. Here, we assemble the overall biomass composition of the biosphere, establishing a census of the

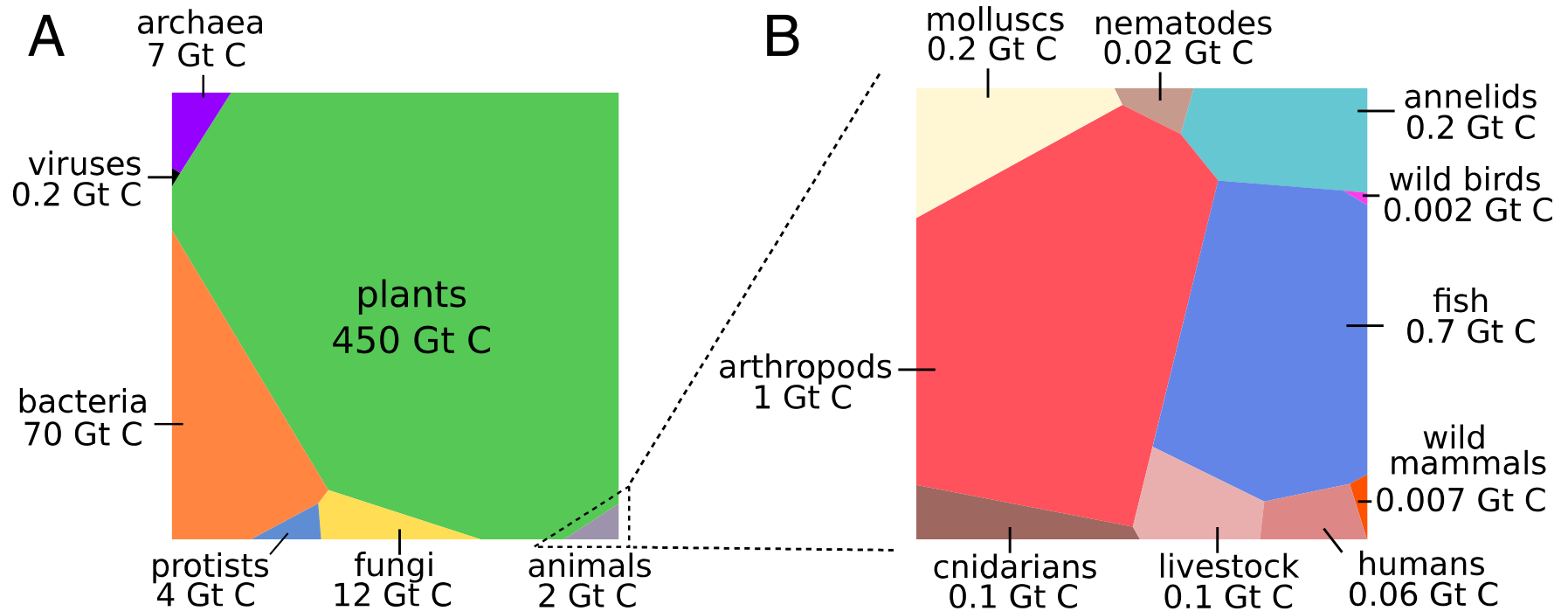
In the past decade, several major technological and scientific advances have facilitated an improved quantitative account of the biomass on Earth. Next-generation sequencing has enabled a more detailed and cultivation-independent view of the composition of natural communities based on the relative abundance of

We assemble a census of the biomass of all kingdoms of life. This analysis provides a holistic view of the composition of the biosphere and allows us to observe broad patterns over taxonomic categories, geographic locations, and trophic modes.

One of the most fundamental efforts in biology is to describe the composition of the living world. Centuries of research have yielded an increasingly detailed picture of the species that inhabit our planet and their respective roles in global ecosystems. In describing a complex system like the biosphere, it is critical to quantify the abundance of individual components of the system (i.e., species, broader taxonomic groups). A quantitative description of the distribution of biomass is essential for taking stock of biosequestered carbon (1) and modeling global biogeochemical cycles (2), as well as for understanding the historical effects and future impacts of human activities.

Table 1, we report our best estimates for the biomass of each taxon analyzed. We use biomass as a measure of abundance, which allows us to compare taxa whose members are of very different sizes. Biomass is also a useful metric for quantifying stocks of elements sequestered in living organisms. We report biomass using the mass of carbon, as this measure is independent of water content and has been used extensively in the literature (6, 16, 17). Alternative measures for biomass, such as dry weight, are discussed in *Materials and Methods*. For ease of discussion, we report biomass in gigatons of carbon, with 1 Gt C = 10^{15} g of carbon. We supply additional estimates for the number of individuals of different taxa in *SI Appendix, Table S1*.

Global biomass of different forms of life (A) and different animal taxa (B)*



* “We estimate that the contribution of reptiles and amphibians to the total animal biomass is negligible....”

Building—and Sustaining—the Macroscope Will Require Targeting Diverse User Communities Beyond Natural History & Systematics

- Conservation and land-use planning
- Climate change and invasive species
- Medicine (infectious disease, drug discovery, etc.)
- Agriculture and forestry
- Physiology, developmental biology, ecology, etc.
- Biologically inspired engineering
- K-12 education