



# **Global Genome Biodiversity Network – Infrastructure for Genomic Research**

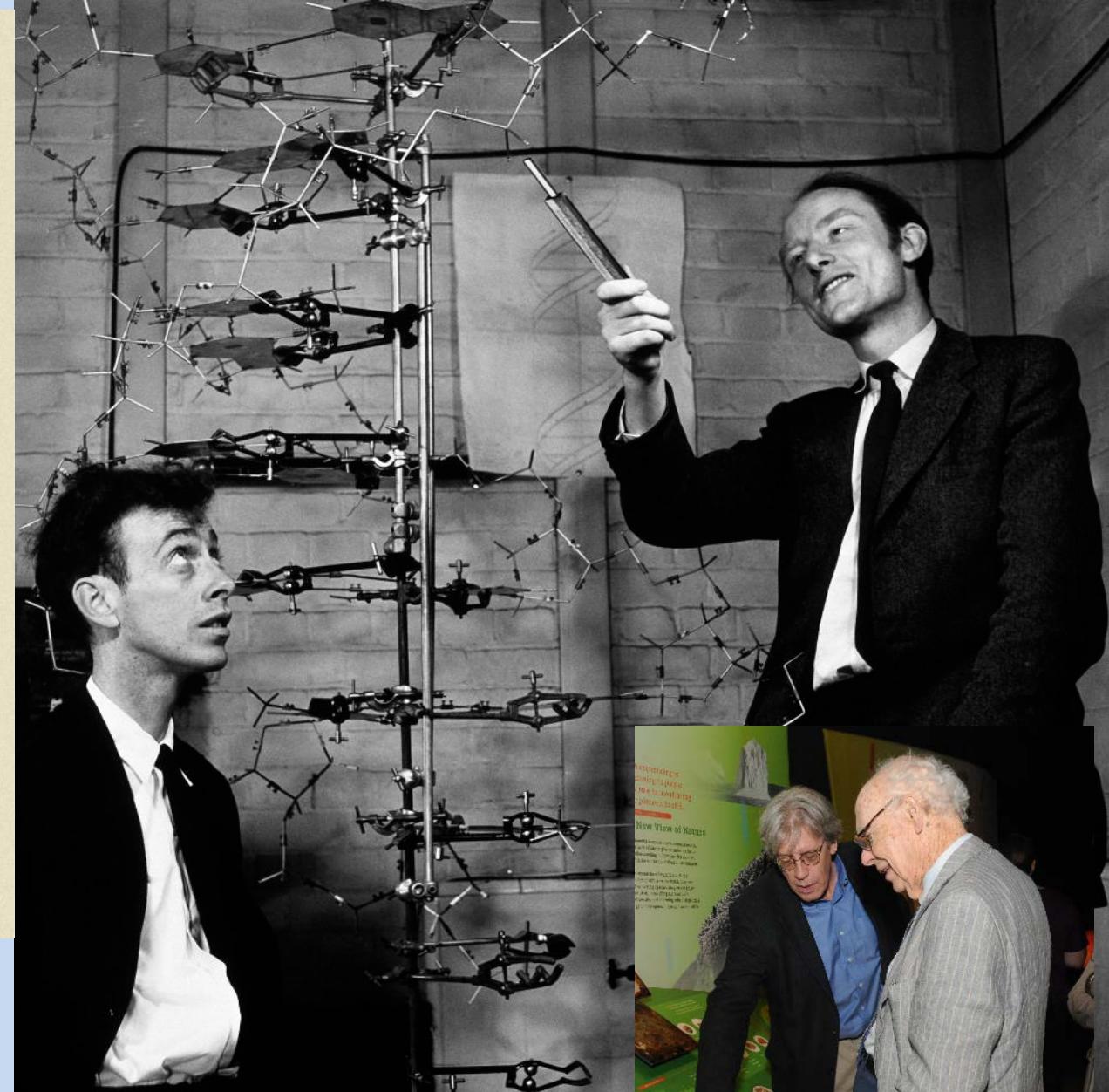
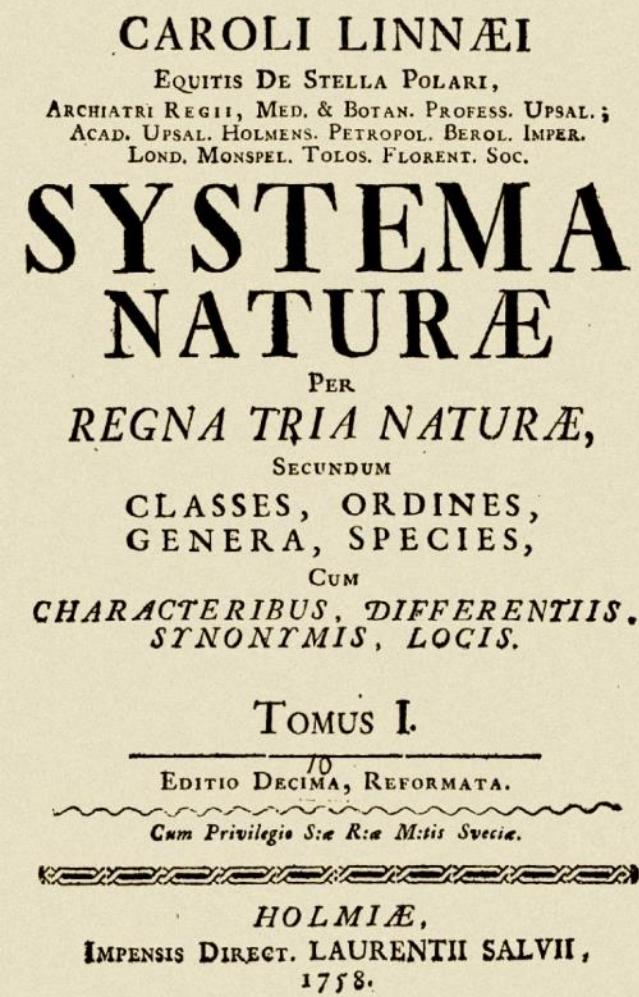
**Jonathan Coddington**

**Executive Committee GGBN**

**National Museum of Natural History,  
Smithsonian Institution**



**Smithsonian  
Institution**



1953

1758



2016



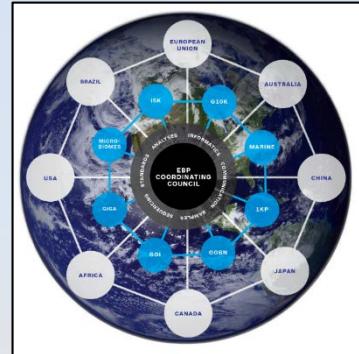
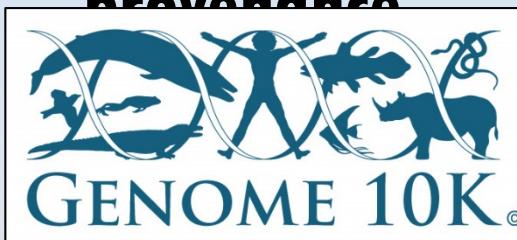
# The Challenge: Sample Access

## — Needs:

- High-quality
- Good metadata
- Legal

## — Sample info dispersed

- Hard to find
- Heterogeneous
- Ambiguous quality & provenance





**Sequences: OK! Specimens: OK! Tissues Not  
OK!**

## Taxa (CoL, GBIF)

Unsequenced

Dark Taxa

Vouchers

“Tissues”

**GenBank  
Sequences**

**GGBN**



# Existing Solutions

- **Established Platforms:**
  - GBIF, NCBI, EMBL, DDBJ, EOL...
- **Standard Data Models:**
  - Darwin Core, ABCD, MiSx...
- **Open Source Software:**



**NOTHING  
COMPARABLE  
FOR TISSUES,  
DNA, etc**



# History and Framework

- **Founded in 2011**
- **Botanic Garden and Botanical Museum Berlin (Technical)**
- **Smithsonian (General)**
- **MOU, Terms of Reference**
- **Core, Associate, Observer members**
- **Museums, Gardens, Zoos, Biorepositories**





# Benefits

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- **Quality standards, best practice, transparency**
  - Data Model & Standardized vocabulary
  - Reciprocal links with other aggregators (GBIF, GenBank)
  - Technical support
  - ABS Guidance
- **Visibility, impact of genetic collections**
- **Biobanking document library**
- **Funding opportunities**
- **Networking**







## GGBN Wiki

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### GGBN Data Standard v1

This page describes the GGBN Data Standard version 1 (v1), released on 04. October 2016.

**Citation:** Droege et al. (2016): The Global Genome Biodiversity Network (GGBN) Standard Specification. DATABASE. doi: 10.1093/database/baw125

**Note:** Ongoing discussions and updates of the GGBN Data Standard will be documented at [http://terms.tdwg.org/GGBN\\_Data\\_Standard](http://terms.tdwg.org/GGBN_Data_Standard)

**The GGBN Data Standard is intended to be used with ABCD or Darwin Core and is not a stand-alone solution!**  
Implementations are available for both ABCD and DarwinCore-Archive. For ABCD two implementations have been made. One for single species samples ABCDGGBN and one for environmental samples ABCDGGBN-Enviro. Please check out the TDWG wiki for updates.

A new stable version of the standard will be released earliest at the end of 2019.

The GGBN Data Standard is a set of terms and controlled vocabularies designed to represent sample facts. It does not cover e.g., scientific name, geography, or physiological facts. This allows combining the GGBN Data Standard with other complementary standards, such as DwC, ABCD or MLxS. Potentially, the standard can be used not only for non-human genomic samples but also human samples. It builds upon existing standards commonly used within the communities extending them with the capability to exchange data on tissue, environmental and DNA sample as well as sequences. The GGBN Data Standard incorporates all molecular terms of MLxS and

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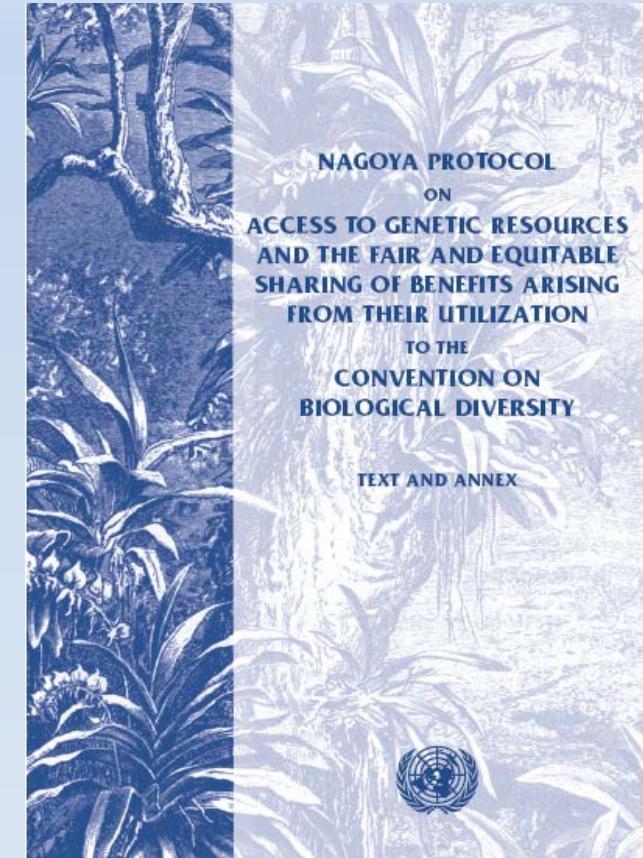
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[http://org/ggbn/Main\\_Page](http://org/ggbn/Main_Page)



# ABS Best Practices

- **Access and Benefit Sharing (ABS) Compliance**
  - **Guidance documentation**
    - **Code of Conduct**
      - **Statement of Use of Genomic Material**
    - **Best Practice**
  - **Example Material Transfer Agreements**
  - **GGBN Data Standard permit vocabulary**



\* Consortium of European Taxonomic Facilities

\*\* Society for the Preservation of Natural History Collections



# Best Practices

## Recommendation for Biorepositories

- Collaboration with ISBER
- Submitted April 2017
- May 2017, in review

\* International Society for Biological and Environmental Repositories

ISBER 4<sup>th</sup> edition





# Data Portal

GGBN Global Genome Biodiversity Network

518362 Records

Samples	
DNA	127828
Tissues	256995
Envrios	74
Repositories	17
Vouchers	
Cultures	21589
eVouchers	2
Specimens	105853
Unknown	6021
Collections	16
Taxa	
Families	1933
Genera	9652
Species	28158
Total	518362

**News**

- 16. August 2016 - GGBN 2016 Conference Presentations now Available
- 08. August 2016 - Entomological Collections Network Joins GGBN
- 02. August 2016 - GGBN - now on Twitter and Facebook
- 28. July 2016 - Korea National Research Resource Center Joins GGBN
- 20. June 2016 - GGBN - a botanical perspective paper published in Annals of Botany
- 18. June 2016 - Museum für Naturkunde (MfN) makes its genomic collections available via GGBN
- 16. June 2016 - Zoology Museum (QCAZ) at Catholic University of Ecuador makes its genomic collections available via GGBN
- 14. June 2016 - Natural History Museum at the University of Oslo (NHMO) makes its genomic collections available via GGBN
- 06. June 2016 - GBIF & GGBN survey on DNA and biodiversity
- 06. June 2016 - Institute of Vertebrate Biology, Academy of Sciences of the Czech Republic makes its genomic collections available via GGBN
- 31. May 2016 - Natural History Museum of Denmark makes its genomic collections available via GGBN
- 25. May 2016 - GGBN Conference: Full Program Now Available

**Explore GGBN**

**Repositories**

**Search**

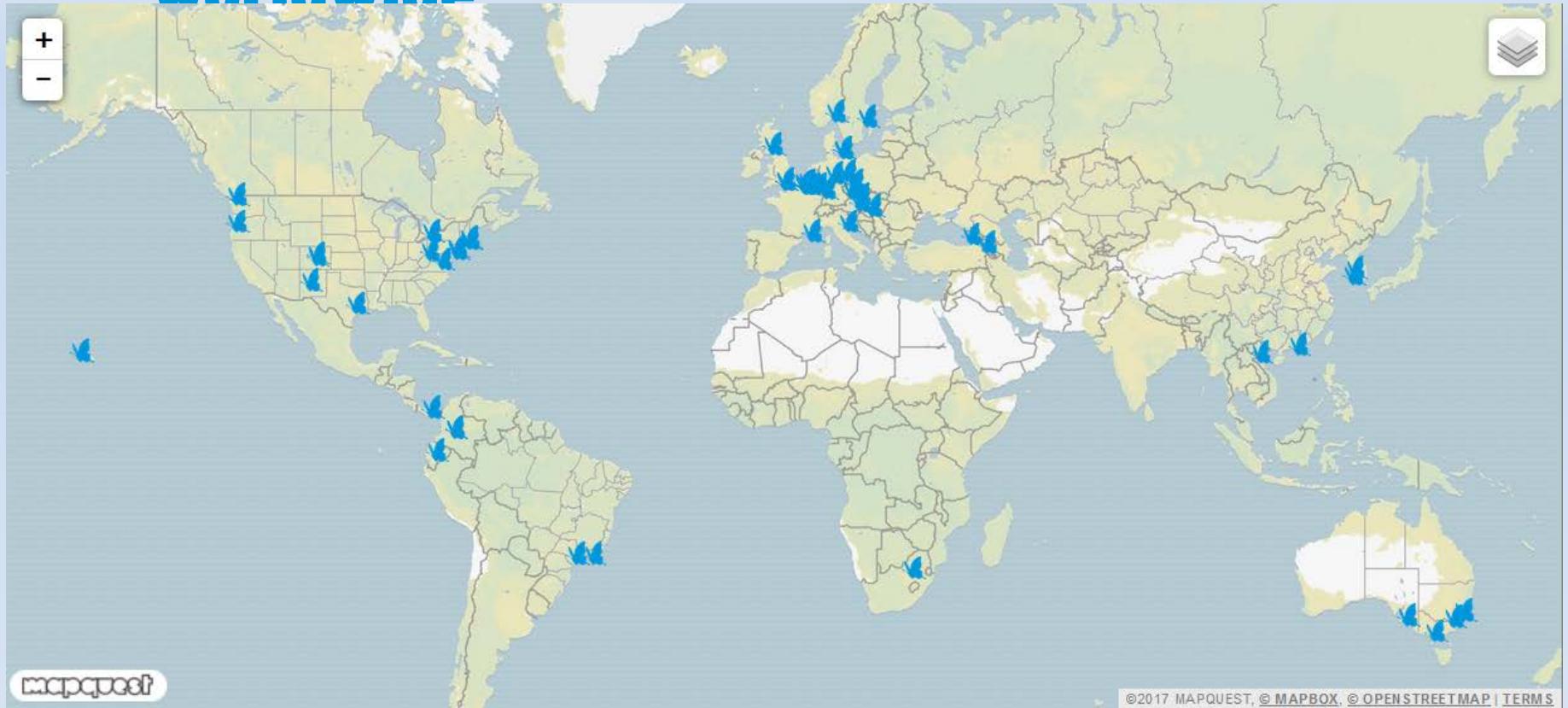
**Documents**

**About**

Cookies Information



# Today: 59 GGBN members, 20 countries worldwide



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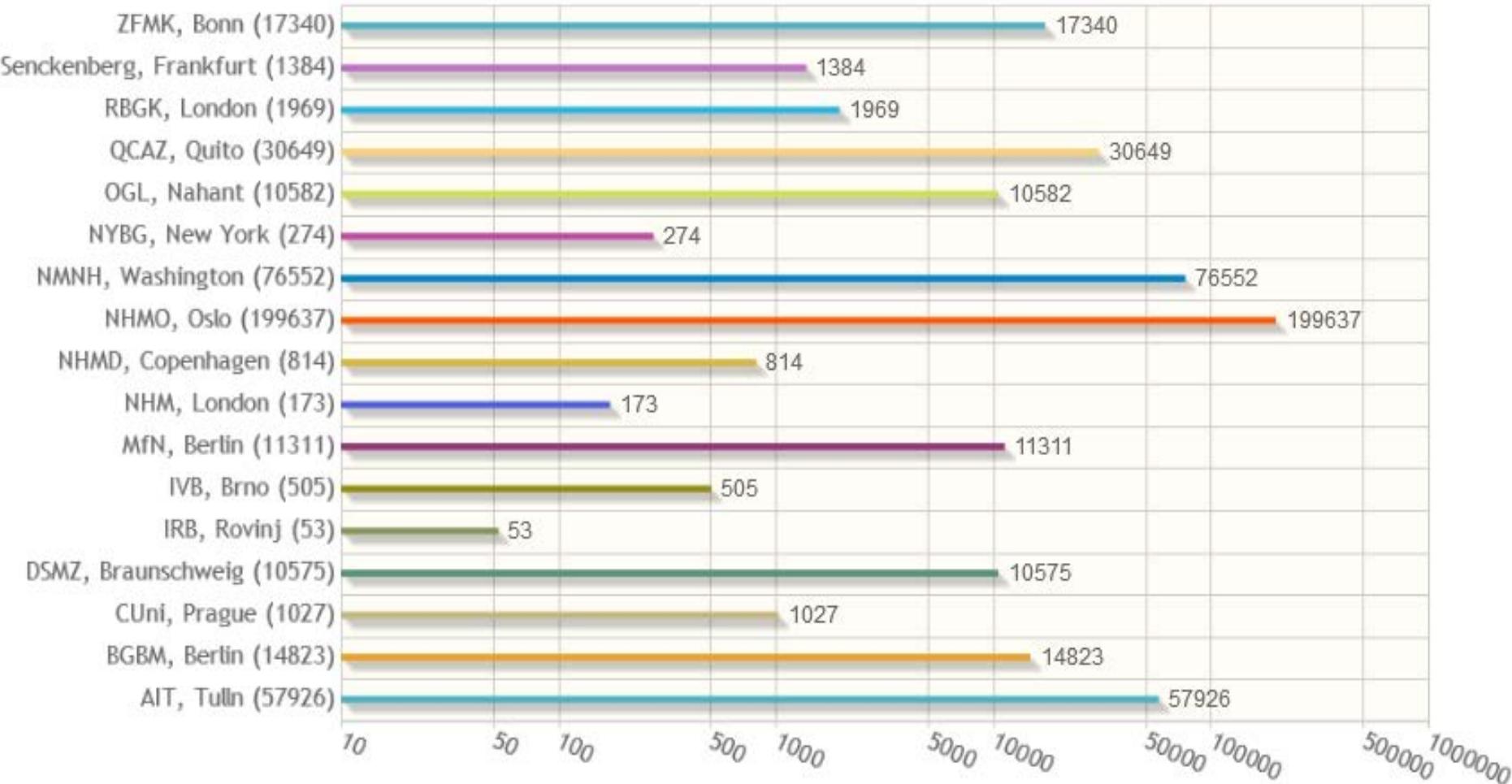




# Repositories (17 push data)

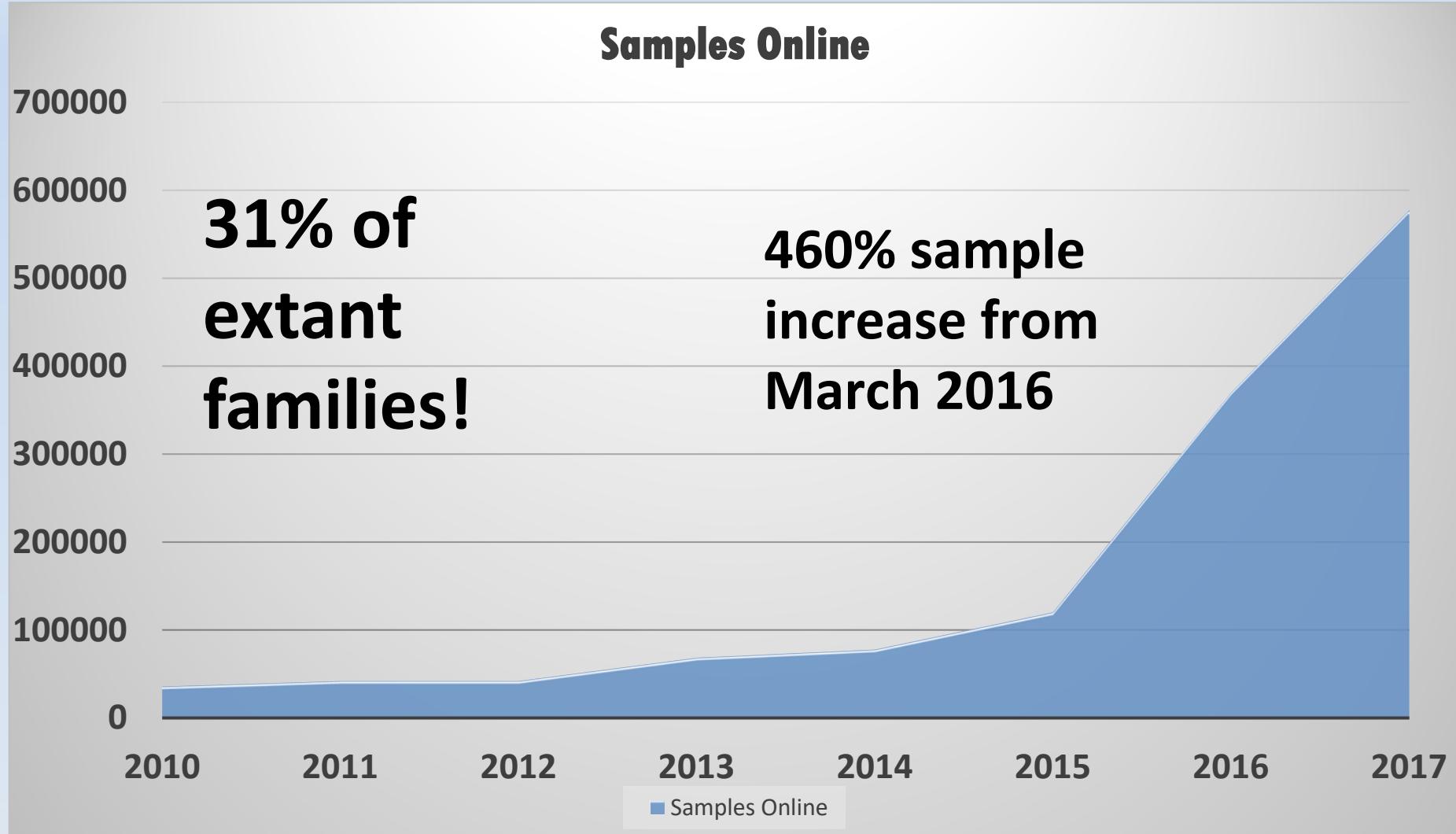
Global statistics **Repositories** No. samples No. taxa

Number of DNA, tissue or environmental sample records (including multiple samples from the same specimen).  
Display with logarithmic scale.





# GGBN Online Sample Growth





# Eukaryotic Biodiversity

- **Domains:** **3 (Eubacteria, Archaea, Eukarya)**
- **Kingdoms:** **5 (animal, plant, fungi, chromista and protozoa)**
- **Phyla:** **61 (35 animal; 10 plant; 2 fungi; 14 chromists+protozoa)**
- **Classes:** **266**
- **Orders:** **1253**
- **Families:** **8,716-9,330**
- **Genera:** **140,000-200,000**
- Sources: *Global Genome Initiative Knowledge Portal (Jonathan Coddington); Catalog of Life (Luisa Abucay & Yuliya Roskov)*
- **Species:** **1.9 million known (est. 10-15 M)**

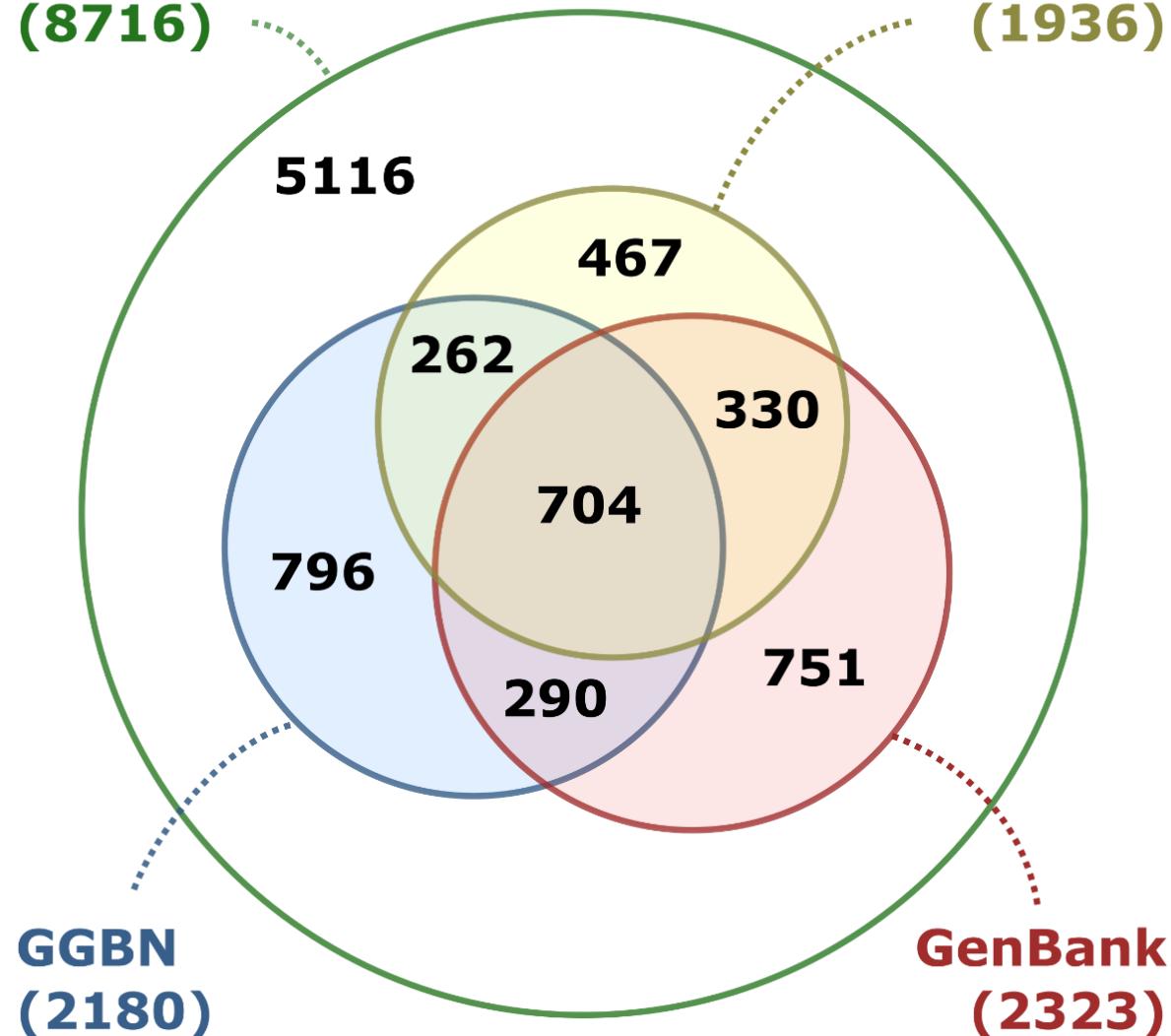


# Gap Analysis

## Families

Catalogue of Life  
(8716)

SI Collections  
(1936)





# Families mostly Singletons! (2180)

N Families	N_BioBanks	Percent
1,591	N_BioBanks=1	0.73
414	N_BioBanks=2	0.19
109	N_BioBanks=3	0.05
43	N_BioBanks=4	0.02
21	N_BioBanks=5	0.01
2	N_BioBanks=6	0.00



# Discoverability, Best Practices, Scale & Integration

- Increase discoverability
  - Biorepositories (new/existing)
  - Aggregators (e.g. iDigBio, VertNet):
    - GGBN data standard
    - Standard usage statistics





# Thank you



<http://www.ggbn.org>

[http://lists.fu-berlin.de/listinfo/GGBN-  
info/](http://lists.fu-berlin.de/listinfo/GGBN-info/)

<https://twitter.com/GGBNOutreach>

<https://www.facebook.com/ggbnoutreach>

**GGBN Executive Committee**

**GGBN Members**

**GGBN Collaborators**

**GGBN Task Forces**

**DFG**

**SYNTHESYS**

**ISBER & ESBB**

**NSF**

**GGI**

