



# **GGBN – Strategies for standardized exchange of genetic resources on a global scale**

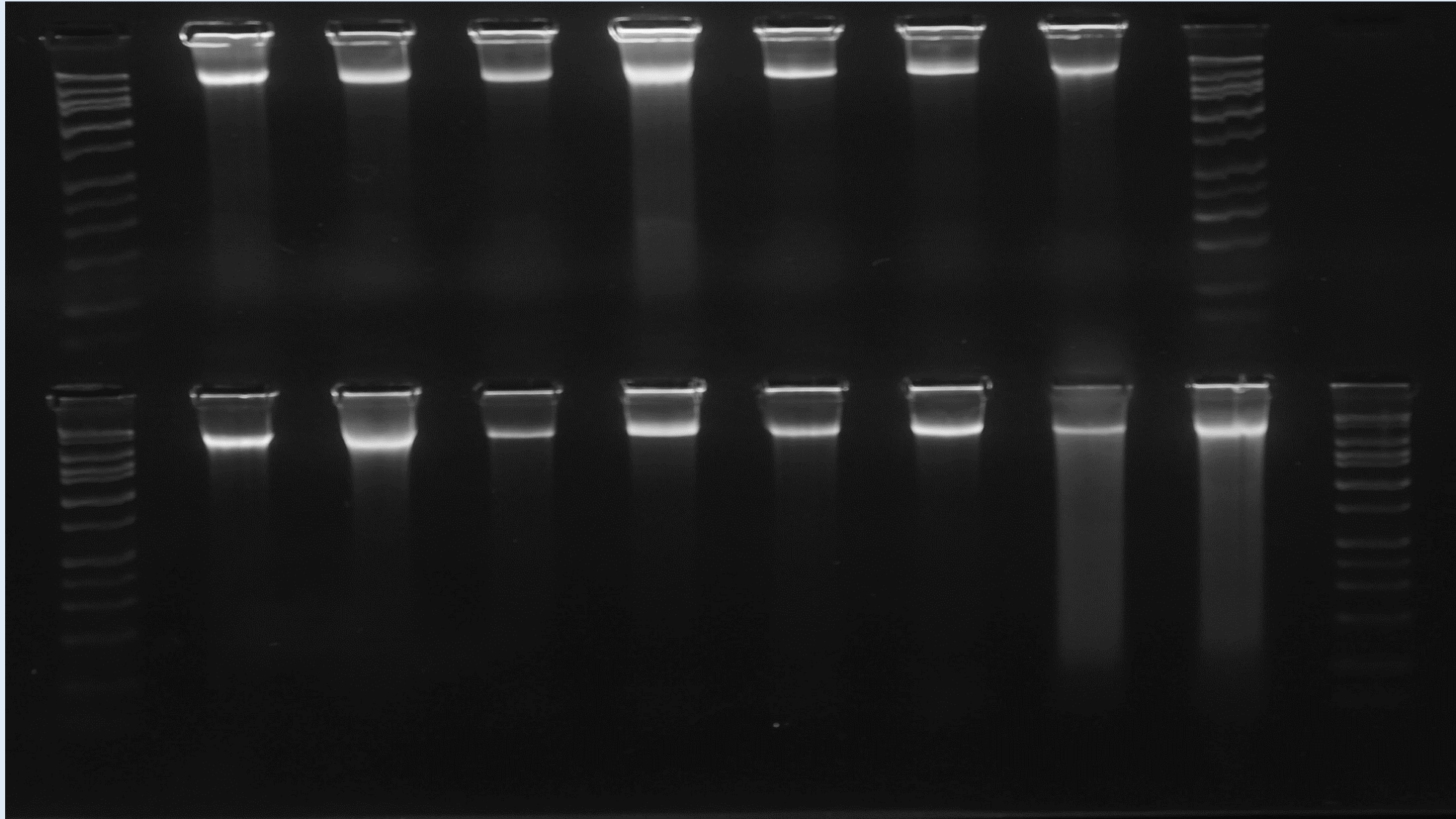
Jonathan Coddington  
Smithsonian Institution

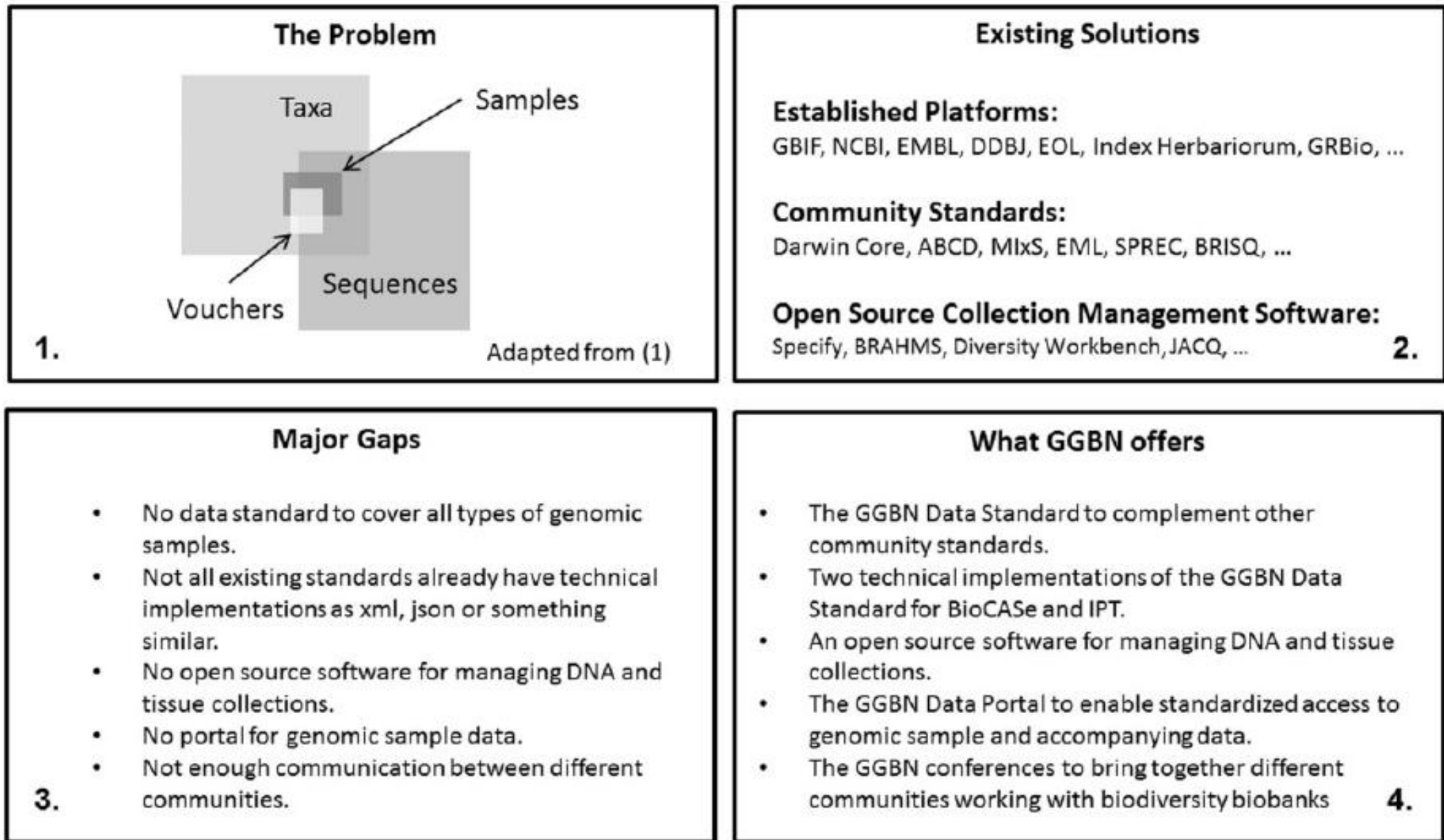


# Global Genome Biodiversity Network

- **Network for non-human biobanks** (e.g. DNA, tissue), founded in 2011
- Based on DNA Bank Network, 2007
- General secretariat: Smithsonian Institution
- Technical secretariat: Botanic Garden and Botanical Museum Berlin
- <http://www.ggbn.org>

# High Molecular Weight DNA





**Figure 1.** Bridging the gaps. Schematic representation of (1) Low percentage of available sequence data in public repositories with proper information where the voucher and/or sample is deposited. (2) Existing tools and platforms for standardized management and access to biodiversity data. (3) Major gaps identified by GGBN and (4) what GGBN has developed to fill these gaps.

# GGBN Data Standard



GGBN Wiki



## GGBN Data Standard v1

Search 

 Log in

Page

Discussion

View

View source

History

What links here

Related changes

Special pages

Printable version

Permanent link

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](https://doi.org/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.

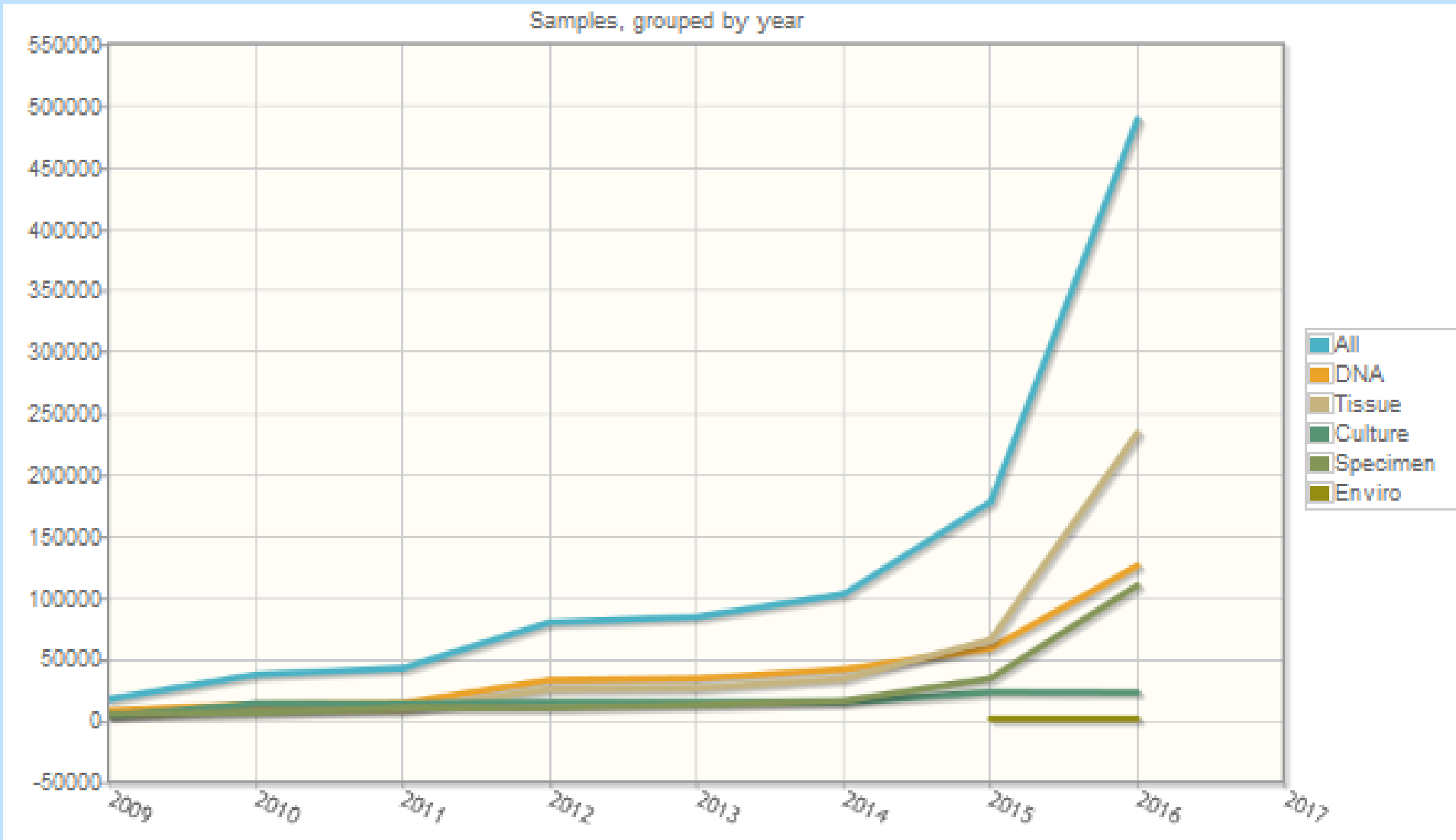
A new stable version of the standard will be released earliest at the end of 2019.

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 [MixS](#). 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 [MixS](#) and

# 50 members, 22 countries: 518,000 samples, 2200 families, 11,000 genera, 32,000 species



# GGBN Growth: 17 Biobanks to date



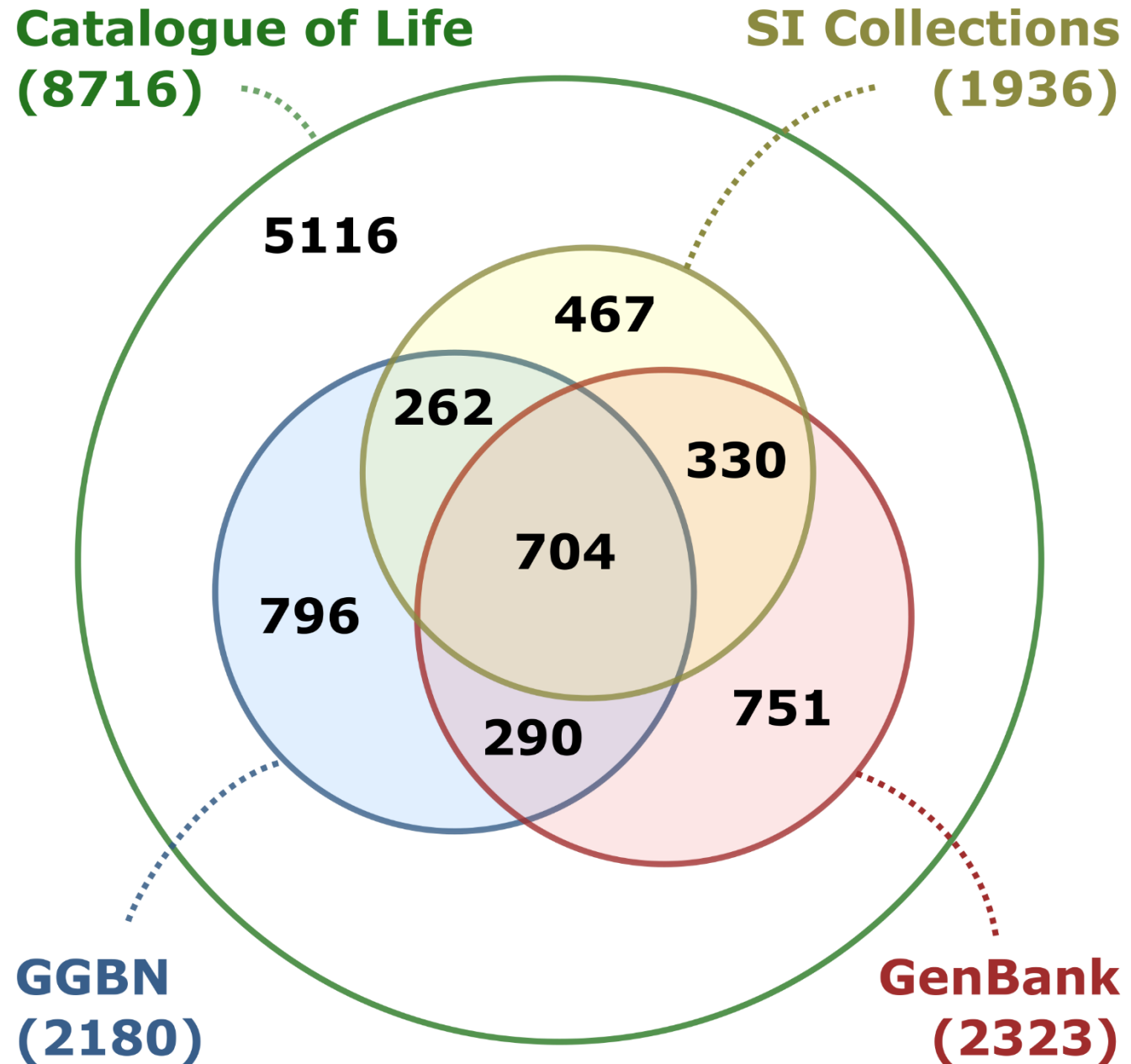
# GGBN Distribution of Families (2329)

N Families	N_BioBanks	Percent
1704	N_BioBanks=1	0.73
442	N_BioBanks=2	0.19
121	N_BioBanks=3	0.05
38	N_BioBanks=4	0.02
20	N_BioBanks=5	0.01
4	N_BioBanks=6	0.00



# Gap Analysis Families

## Families



# Thanks!



GLOBAL  
GENOME  
INITIATIVE

Global Genome  
Biodiversity  
Network (GGBN)



AMERICAN MUSEUM OF NATURAL HISTORY



museum für  
naturkunde  
berlin



SENCKENBERG  
world of biodiversity

