



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

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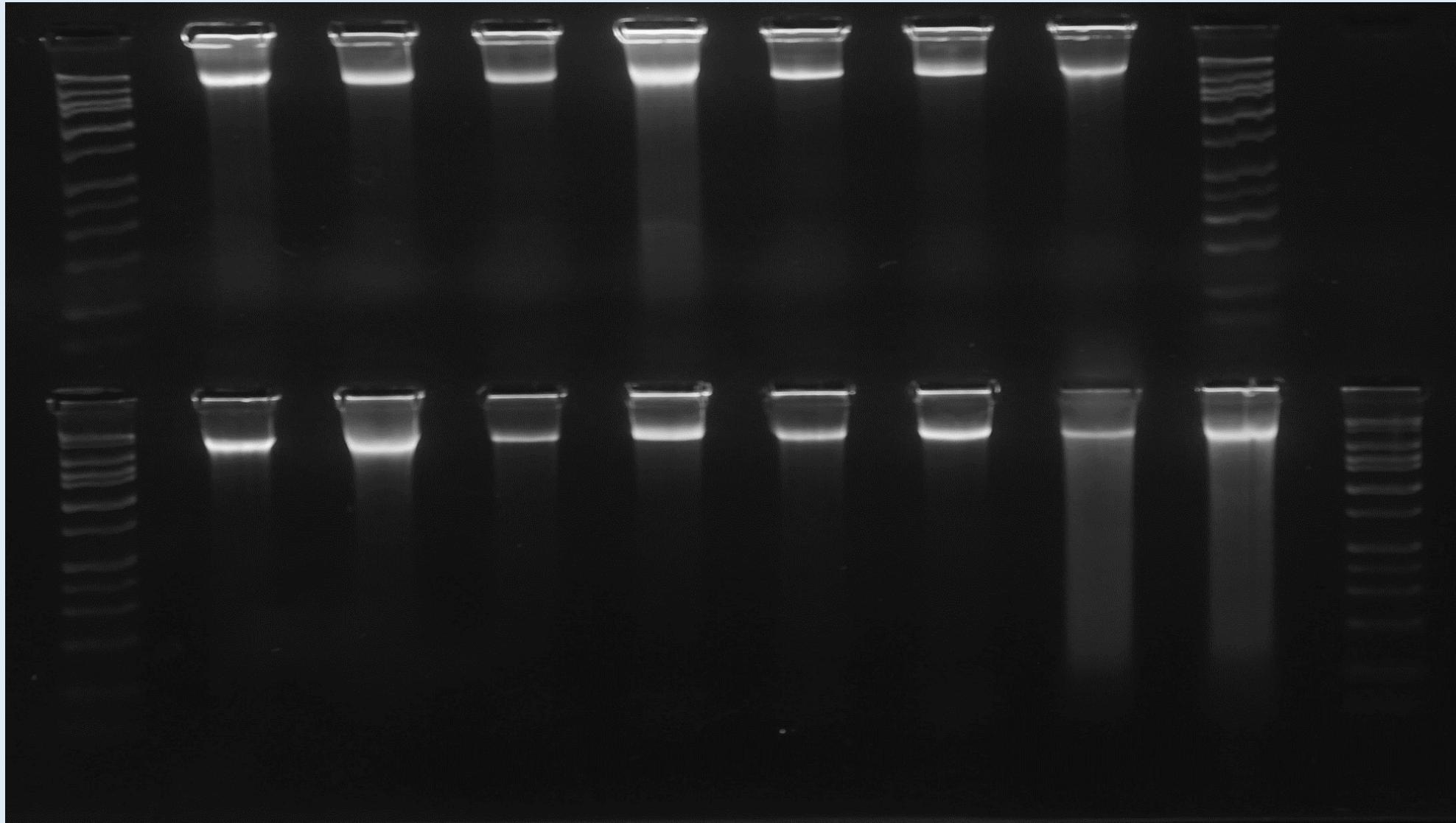
Smithsonian Institution



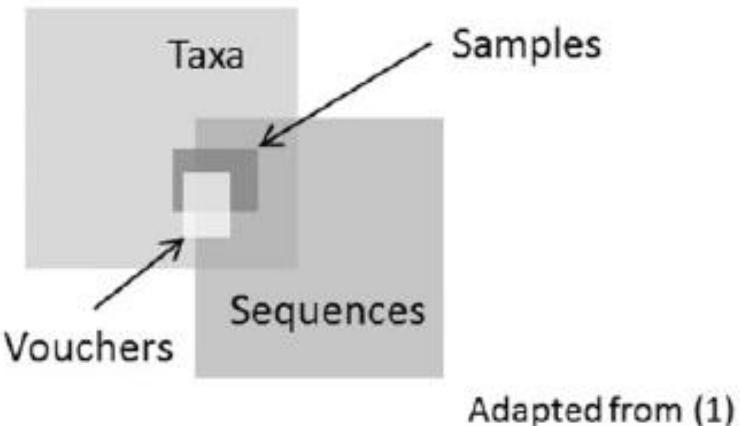
Global Genome Biodiversity Network

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

High Molecular Weight DNA



The Problem



Existing Solutions

Established Platforms:

GBIF, NCBI, EMBL, DDBJ, EOL, Index Herbariorum, GRBio, ...

Community Standards:

Darwin Core, ABCD, MiS, EML, SPREC, BRISQ, ...

Open Source Collection Management Software:

Specify, BRAHMS, Diversity Workbench, JACQ, ...

2.

Major Gaps

- No data standard to cover all types of genomic samples.
 - Not all existing standards already have technical implementations as xml, json or something similar.
 - No open source software for managing DNA and tissue collections.
 - No portal for genomic sample data.
 - Not enough communication between different communities.
- 3.

What GGBN offers

- The GGBN Data Standard to complement other community standards.
 - Two technical implementations of the GGBN Data Standard for BioCASE and IPT.
 - An open source software for managing DNA and tissue collections.
 - The GGBN Data Portal to enable standardized access to genomic sample and accompanying data.
 - The GGBN conferences to bring together different communities working with biodiversity biobanks
- 4.

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

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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

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](#).

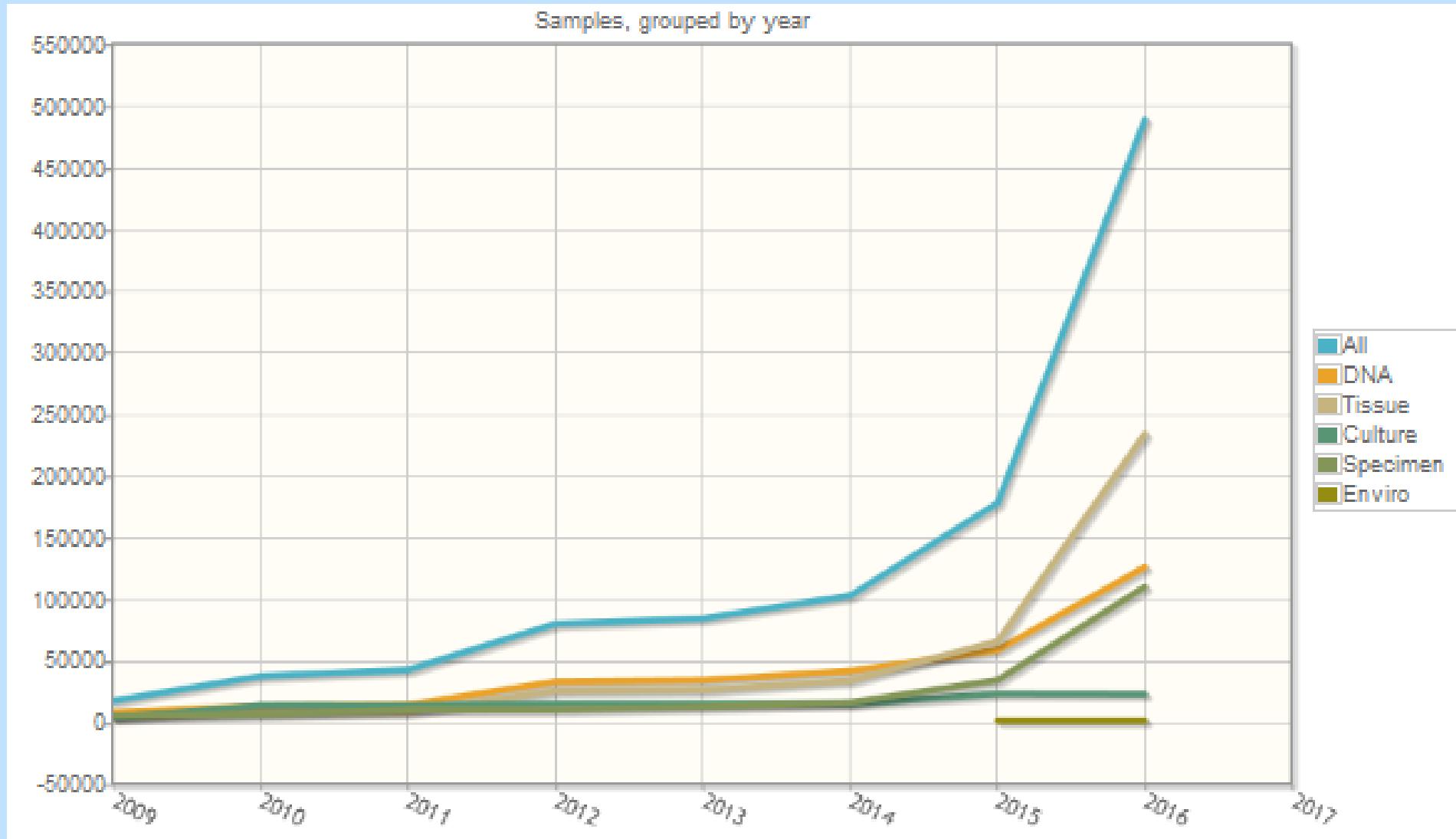
A new stable version of the standard will be released earliest at the end of 2019.

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

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



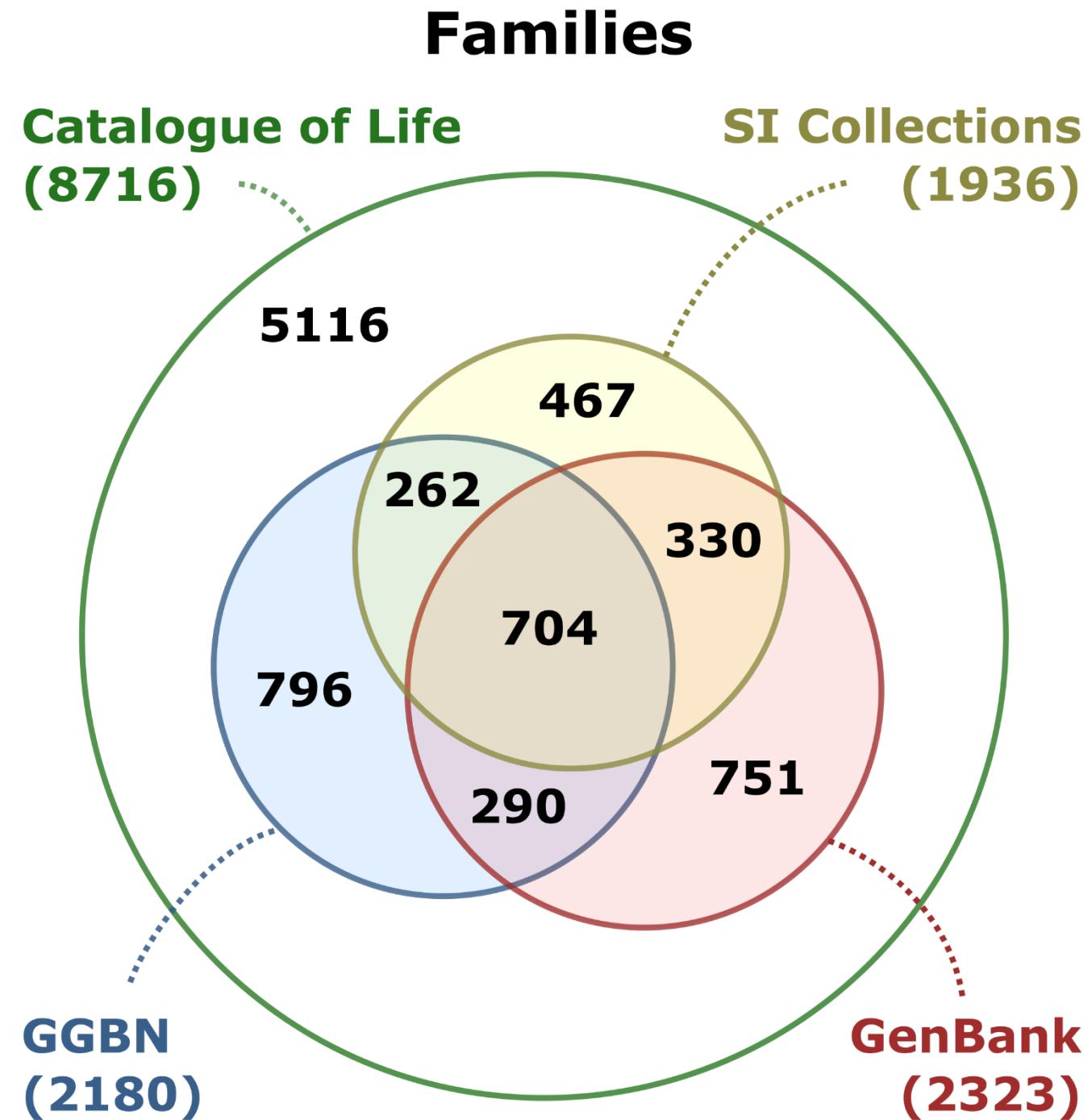
GGBN Growth: 17 Biobanks to date



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



Thanks!



GLOBAL
GENOME
INITIATIVE

Global Genome
Biodiversity
Network (GGBN)



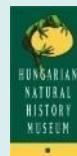
AMERICAN MUSEUM & NATURAL HISTORY



SENCKENBERG
world of biodiversity



museum für
naturkunde
berlin



Smithsonian
National Museum of Natural History



NCBI
National Center for
Biotechnology Information



Integrated Taxonomic
Information System



THE NEW YORK BOTANICAL GARDEN

Barcode of Life
Identifying Species with DNA Barcoding



Smithsonian
National Zoological Park

CONSORTIUM FOR THE
BARCODE OF LIFE



Smithsonian Tropical Research Institute
Center for Tropical Forest Science