

# Understanding the Research Data Lifecycle

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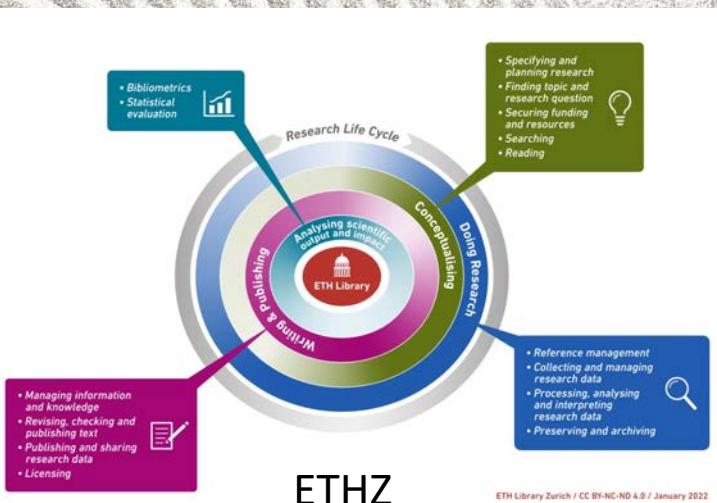


Image: <http://www.voicendata.com/wp-content/uploads/2015/07/Data-Management.png>

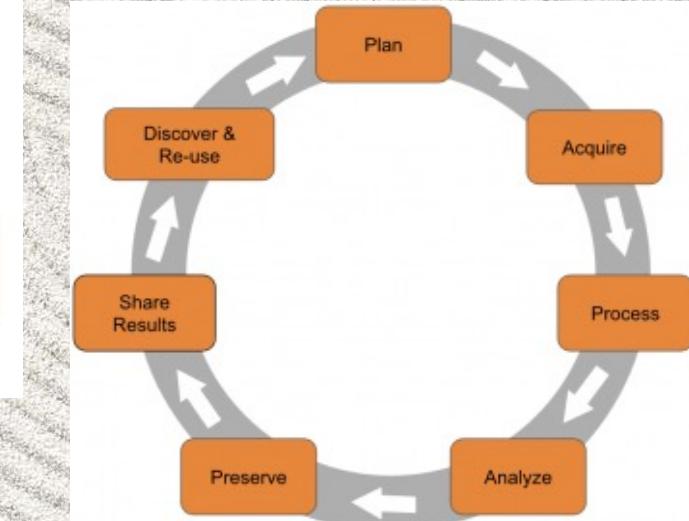
# Research Data Lifecycle



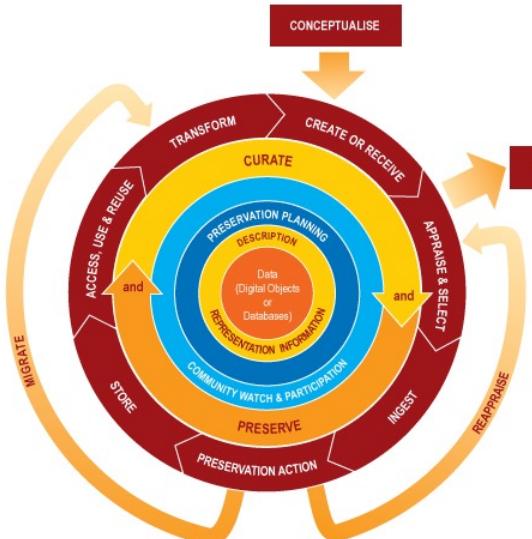
Harvard University



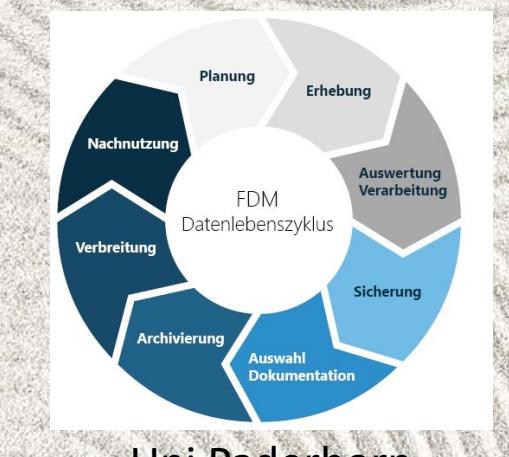
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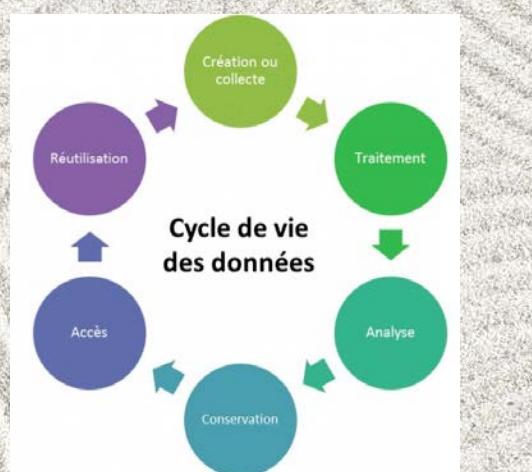
Princeton University



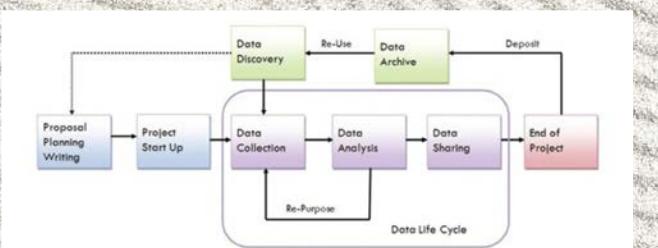
UK Digital Curation  
Centre



Uni Paderborn

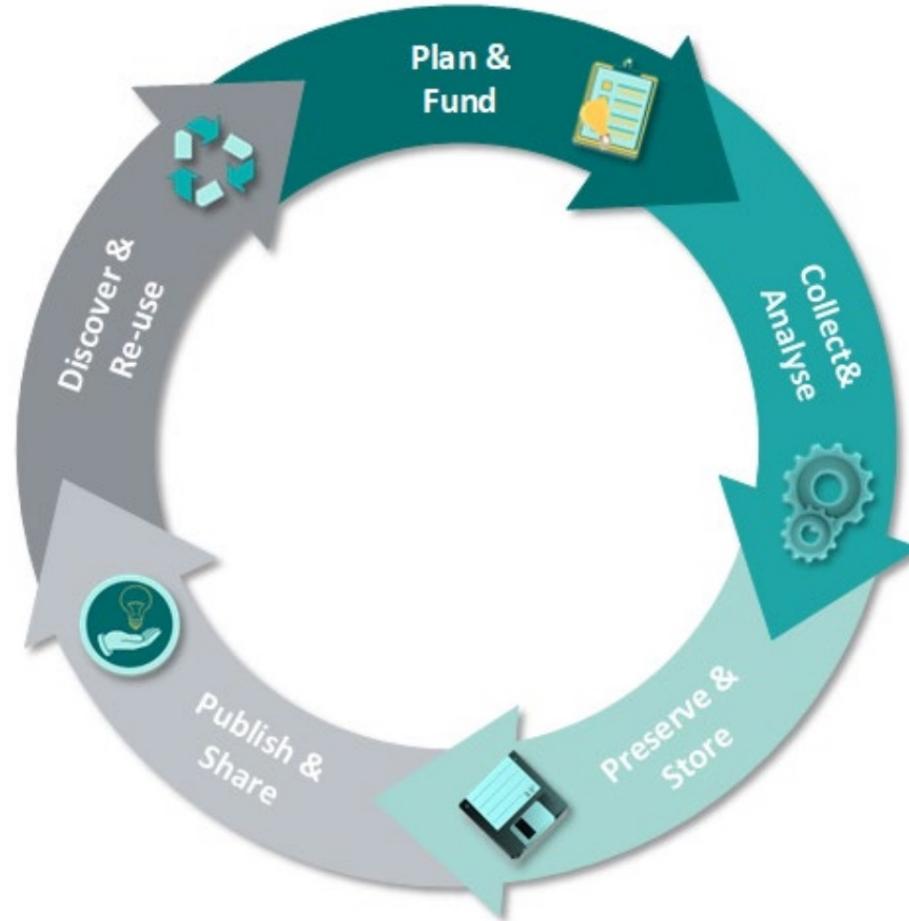


Institut Pasteur

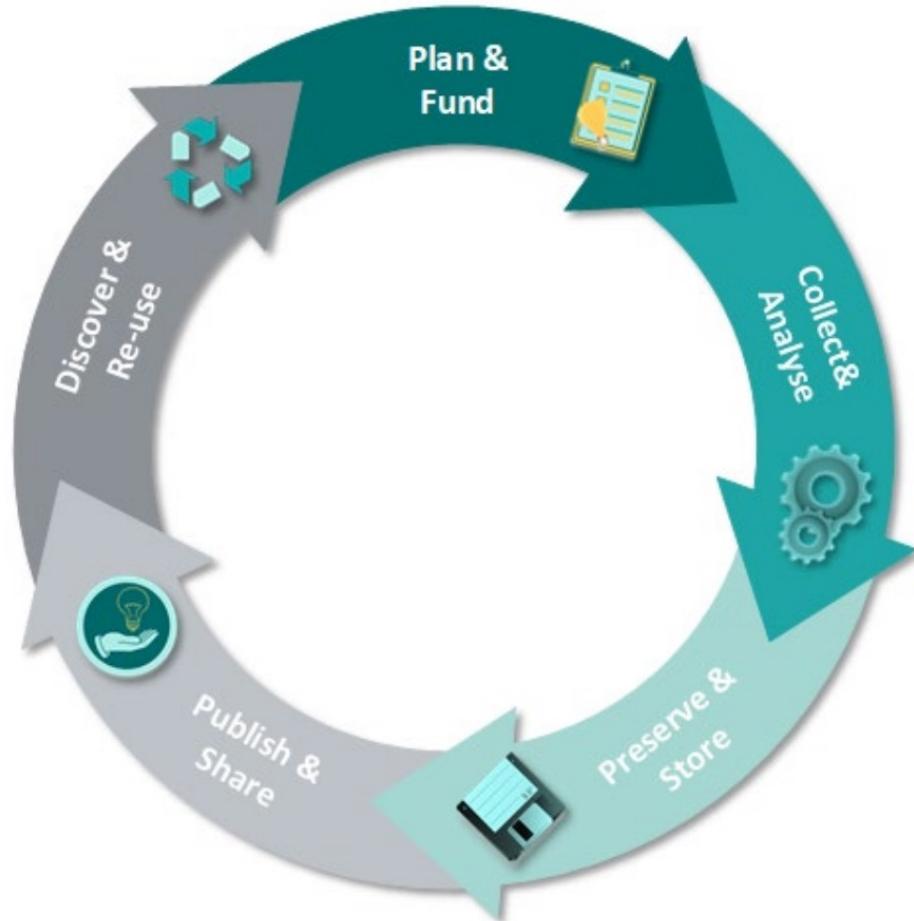


University of Oslo

# Research Data Lifecycle: Uni Basel Version



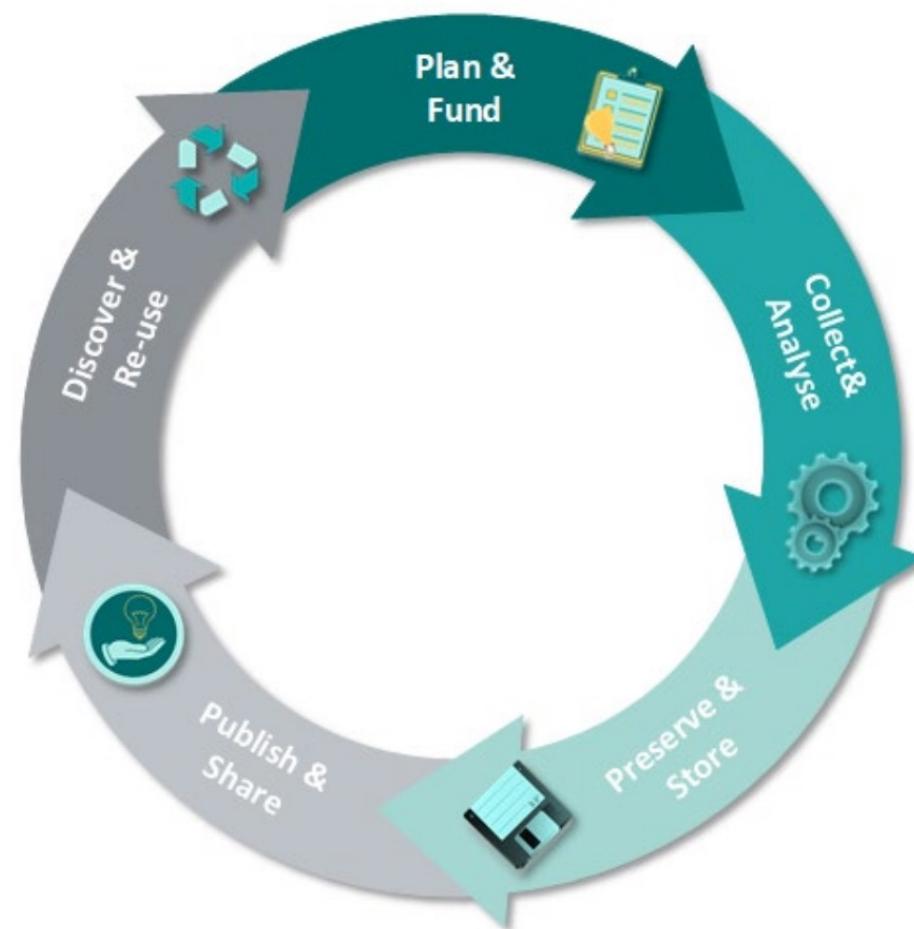
# First phase: Conceptualize and Receive



## Project Planning Phase

What data? How much data?  
What data is existing / can be re-used?  
How will data be collected, organized, managed?  
Ethical, legal, security concerns  
Data exchange, re-use?  
**-> Data Management Plans**  
**-> Part of planning a funded project**

# Phase 2: Create, Evaluate, Collaborate, Store



## Active Research

Experiments, observations, secondary materials collected

Data is being processed (transformed, collated, converted, validated, cleaned, etc.)

Metadata is created: Data sets are being described

Data is being analyzed and interpreted: Raw materials are turned into insights. Instruments, code, protocols need to be documented

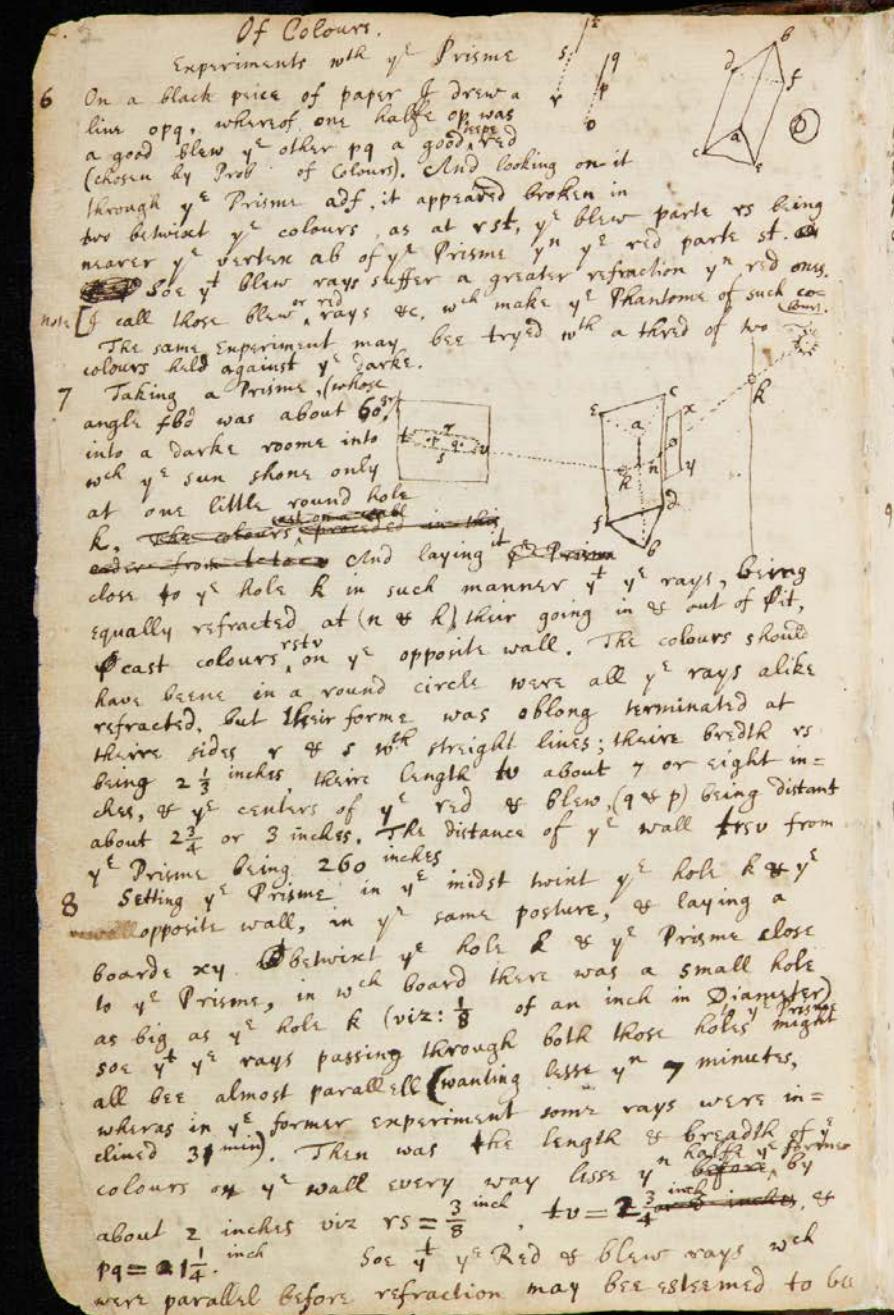
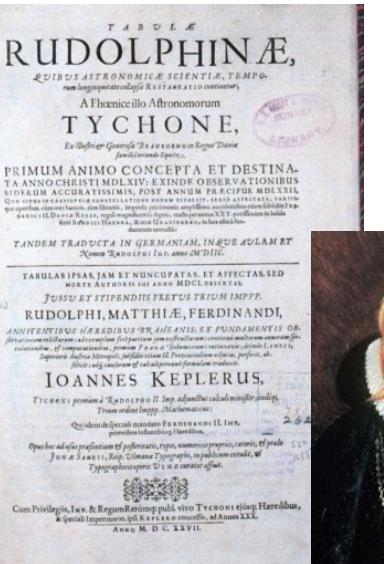
Scientific results are published or patented

Data is stored in active storage.

# Documenting your research is part and parcel of the scientific method

Document your work, your observations both expected and unexpected, and your data.

- Laboratory Notebooks
- Drawings, paintings
- Numeric data in tables, books



# The key to scientific record keeping:

Document your own research in a way that someone else with a comparable education/skill set as yours can understand it

Such as:

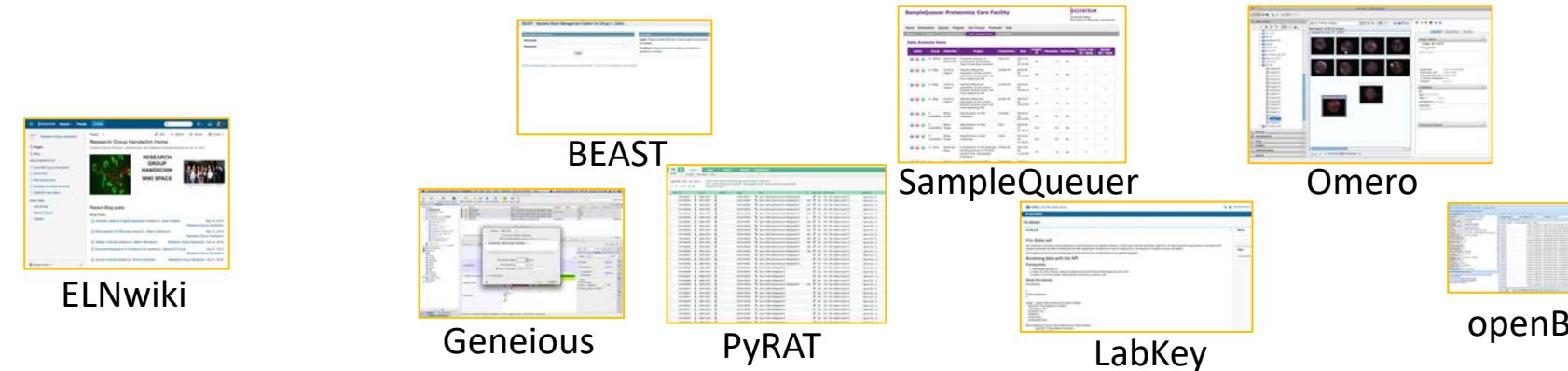
- A collaborator
- Your group leader
- Someone who picks up the project after you
- Someone in another group who wants to extend the research you published earlier
- ...
- Or you yourself, 7 years later, trying to write a publication

# Scientific Record Keeping Best Practices

Useful & good research records should include these details:

- **What** you did – experimental protocol
- **When** you did it - date
- **Why** you did it – objective
- **How** you did it – methods
- **Who** you are (the person creating the record)
- **What project(s)** this work was part of
- Who conceived of the study (if not yourself)
- Special materials & instruments utilized
- Source of materials & instruments
- Discussion of data – results – expected and unexpected
- Data handling and analyses
- Data interpretation by yourself (and others if pertinent)
- Next steps based on reported results

# Where to store your data: Data management systems



- **Information about experiments**
- *Who did what when and why?*
- *What was the outcome?*
- **Protocols**
- **Descriptions**
- **Gel pictures, ...**
- **Track samples and materials**
- Track provenance
- Traceability of processes
- Procedures and workflows
- Automation is relevant
- **Strains**
- **Vectors, ...**
- **Store and annotate large datasets**
- Provide programmatic access
- Make datasets searchable
- Analysis and visualization may be built-in
- **Microscopy images**
- **Sequencing data, ...**

# Where to store your data: Storing files safely

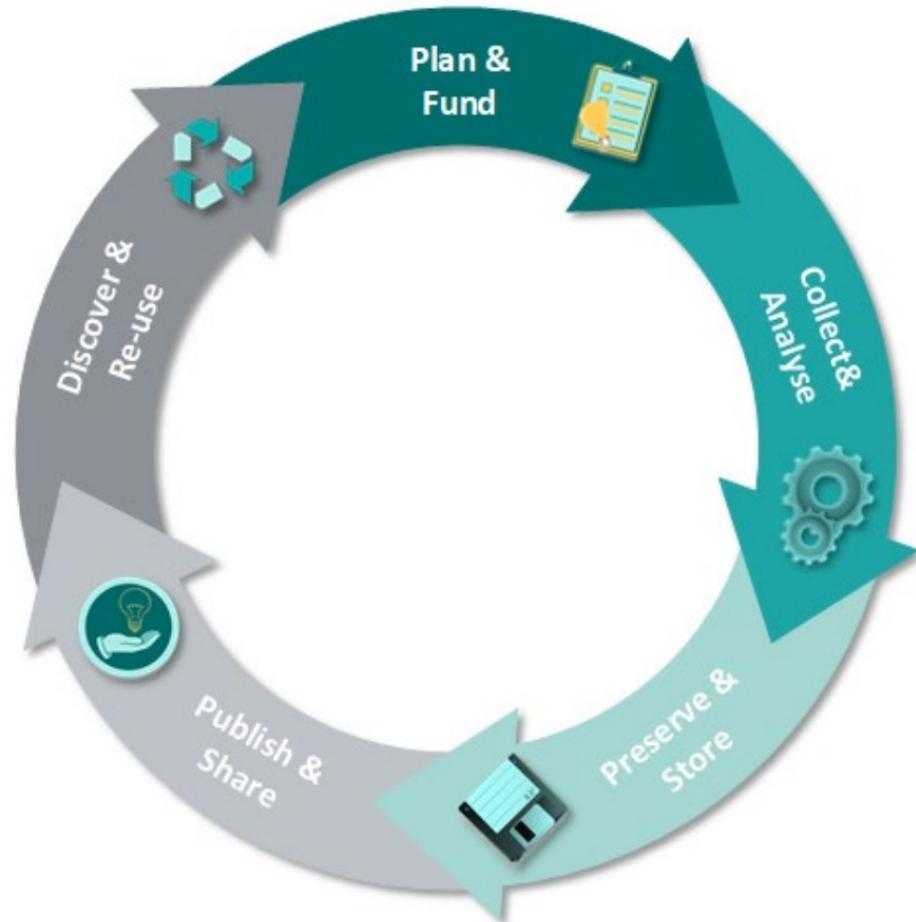


180TB of genomic data on USB disks.  
What could possibly go wrong...

When storing data in files,  
consider the following at the  
beginning of a project:

- Use centrally managed storage.
- What type, size is your data?  
Which storage system is best used?
- Use a hierarchical and logical folder structure
- Separate active and completed projects,  
**delete unused temporary files**
- **Version your files** using a convention or a SCM system (git)
- Use a **naming convention for your file names**
- Use well-known data formats  
(ideally non-proprietary, open-source formats)
- Describe your data with proper documentation and metadata.

# Phase 3: Conserve, Annotate, Curate, Archive



## Preserve & Store

Active storage vs. preservation for long term:

- Completion of metadata annotation
- Selection of data with long term value
- Convert files into the best file format
- Prepare for deposition
- Archive data to be retained but not published





Size of this preview: 800 × 536 pixels. Other resolutions: 320 × 214 pixels | 640 × 429 pixels | 1,024 × 686 pixels | 1,280 × 857 pixels | 7,000 × 4,687 pixels.

This file has annotations. Move the mouse pointer over the image to see them.

Original file (7,000 × 4,687 pixels, file size: 2.66 MB, MIME type: image/jpeg): ZoomViewer: flash/no flash



## Summary [ edit ]

Description	President Lyndon B. Johnson signs the 1964 Civil Rights Act as Martin Luther King, Jr., and others, look on
Date	2 July 1964
Source	<a href="http://photolab.lib.utexas.edu/detail.asp?id=18031">http://photolab.lib.utexas.edu/detail.asp?id=18031</a>
Author	Cecil Stoughton, White House Press Office (WHPO)
Permission (Reusing this file)	"This image is in the public domain and may be used free of charge without permissions or fees."

Technical Data:  
Type, File Format,  
Resolution ...

Content  
Description

Origin,  
Creator,  
License

# Discipline-specific metadata standards

<https://rd-alliance.github.io/metadata-directory/>



## Life Sciences

### ABCD (Access to Biological Collection Data) [Edit](#)

The Access to Biological Collections Data (ABCD) Schema is an evolving comprehensive standard for the access to and exchange of data about specimens and observations (a.k.a., primary biodiversity data). The ABCD Schema attempts to be comprehensive and highly structured, supporting data from a wide variety of databases. It is compatible with several existing data standards. Parallel structures exist so that either (or both) structured data and free-text can be accommodated.

Sponsored by Biodiversity Information Standards TDWG - the Taxonomic Databases Working Group, the current specification was last modified in 2007.

### Darwin Core [Edit](#)

A body of standards, including a glossary of terms (in other contexts these might be called properties, elements, fields, columns, attributes, or concepts) intended to facilitate the sharing of information about biological diversity by providing reference definitions, examples, and commentaries.

Sponsored by Biodiversity Information Standards TDWG; the current standard was last modified in October 2009.

### EML (Ecological Metadata Language) [Edit](#)

Ecological Metadata Language (EML) is a metadata specification particularly developed for the ecology discipline. It is based on prior work done by the Ecological Society of America and associated efforts (Michener et al., 1997, Ecological Applications).

Sponsored by ecomformatics.org, EML Version 2.2.0 was released in 2019.

### Genome Metadata [Edit](#)

Genome metadata on PATRIC consists of 61 different metadata fields, called attributes, which are organized into the following seven broad categories: Organism Info, Isolate Info, Host Info, Sequence Info, Phenotype Info, Project Info, and Others.

### ISA-Tab [Edit](#)

The Investigation/Study/Assay (ISA) tab-delimited (TAB) format is a general purpose framework with which to collect and communicate complex metadata (i.e. sample characteristics, technologies used, type of measurements made) from 'omics-based' experiments employing a combination of technologies.

Created by core developers from the University of Oxford, ISA-TAB v1.0 was released in November 2008.

### MIBBI (Minimum Information for Biological and Biomedical Investigations) [Edit](#)

A common portal to a group of nearly 40 checklists of Minimum Information for various biological disciplines. The MIBBI Foundry is developing a cross-analysis of these guidelines to create an interoperable, extensible community of standards.

The concept was initiated initially through the joint efforts of the Proteomics Standards Initiative, the Genomic Standards Consortium and the MGED FGSB Working Groups. The latest project to register with MIBBI is the MAFBle guidelines for reporting biofilm.

### Nexus [Edit](#)

Nexus is an international standard for the storage and exchange of neutron, x-ray, and muon experiment data. The structure of Nexus files is extremely flexible, allowing the storage of both simple data sets, such as a single data array and its axes, and highly associated metadata, such as measurements on a multi-component instrument or numerical simulations. Nexus is built on top of the container format HDF5, and adds domain-specific rules for organizing data within HDF5 files in addition to a dictionary of names.

### Observ-O [Edit](#)

Observ-O is founded on four basic concepts to represent any kind of observation: Targets, Features, Protocols (and their Applications), and Values. It is intended to lower the barrier for future data sharing and facilitate integrated search across panels and documentation, and software are available for free and open source (LODPLUG) at <http://www.observ-om.org>.

### ODAM Structural Metadata [Edit](#)

Open Data for Access and Mining (ODAM) Structural Metadata is a format describing how the metadata should be formatted and what should be included to ensure ODAM compliance for a data set. To comply with this format, two metadata files in TSV (text file). These two files describe the metadata of the dataset, which includes descriptions of measures and structural metadata like references between tables. The metadata lets non-expert users explore and visualize your data. By making data interpretable for humans and machines, it also encourages data dissemination according to FAIR principles. The structural metadata is specified in section 'Data collection and preparation' on the website.

### OME-XML (Open Microscopy Environment XML) [Edit](#)

OME-XML is a vendor-neutral file format for biological image data, with an emphasis on metadata supporting light microscopy. It can be used as a data file format in its own right, or as a way of encoding metadata within a TIFF or BigTIFF file (for which it is a specification).

The standard is maintained by the Open Microscopy Environment Consortium, and was last updated in June 2012.

### PDBx/mmCIF (Protein Data Bank Exchange Dictionary and the Macromolecular Crystallographic Information Framework) [Edit](#)

Protein Data Bank archive (PDB) is the single worldwide archival repository of information about the 3D structures of proteins, nucleic acids, and complex assemblies, managed by the Worldwide PDB (wwPDB). The PDB Exchange Dictionary (PDBx) is used for content deposition, annotation and archiving of PDB entries. PDBx incorporates the community standard metadata representation, the Macromolecular Crystallographic Information Framework (mmCIF), originally developed under the auspices of the International Union of Crystallography (IUC). PDBx has been extended by the wwPDB to include descriptions of other experimental methods that produce 3D macromolecular structure models such as Nuclear Magnetic Resonance Spectroscopy, 3D Electron Microscopy and Tomography.

### Protocol Data Element Definitions [Edit](#)

A draft set of data elements required by the National Institutes of Health (U.S.) for the submission of trial information to the ClinicalTrials.gov registry and results database.

### Repository-Developed Metadata Schemas [Edit](#)

Some repositories have decided that current standards do not fit their metadata needs, and so have created their own requirements.

### RO-Crate [Edit](#)

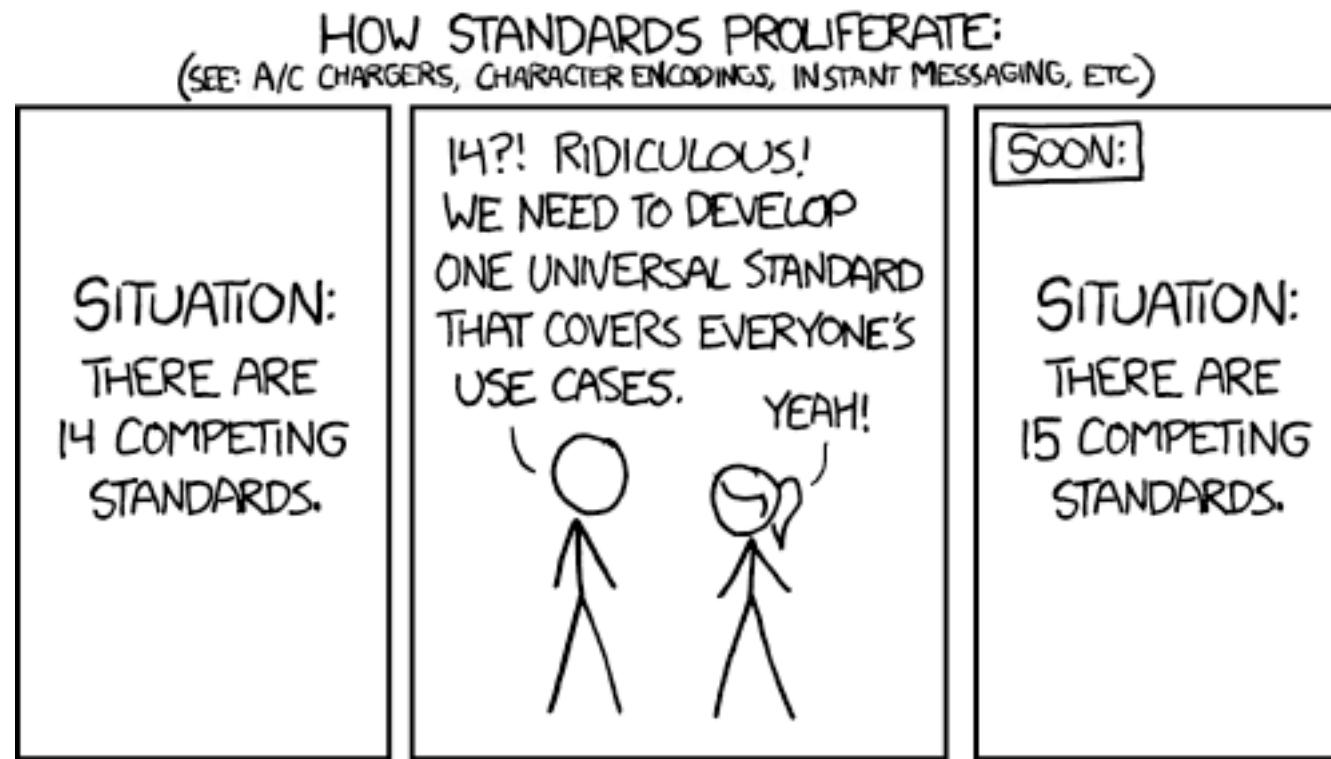
RO-Crate is a community effort to establish a lightweight approach to packaging research data with their metadata. It is based on schema.org annotations in JSON-LD, and aims to make best-practice in formal metadata description accessible and practical situations, from an individual researcher working with a folder of data, to large data-intensive computational research environments.

### UKEOF [Edit](#)

A metadata standard for describing environmental monitoring activities, programmes, networks and facilities published by the UK Environmental Observation Framework (UKEOF).

<https://fairsharing.org/>

# Why are there so many metadata standards?



The good news is:

**As research fields mature, standards will eventually become generally accepted, e.g PDB files, fasta files, pepXML, OME-TIFF, ...**

# README files

## README content:

- General information
- Data and file overview
- Sharing and access information
- Methodological information

## Best practices

- 1 README / 1 data folder (whenever possible)
- Name the README in accordance with described files
- Write the README as a plain text file (open format)
- Use the same format for multiple READMEs
- Follow your discipline's scientific taxonomic conventions
- Use standardized date formats.

Suggested: [W3C/ISO 8601 date standard](#),

YYYY-MM-DD or YYYY-MM-DD hh:mm:ss.

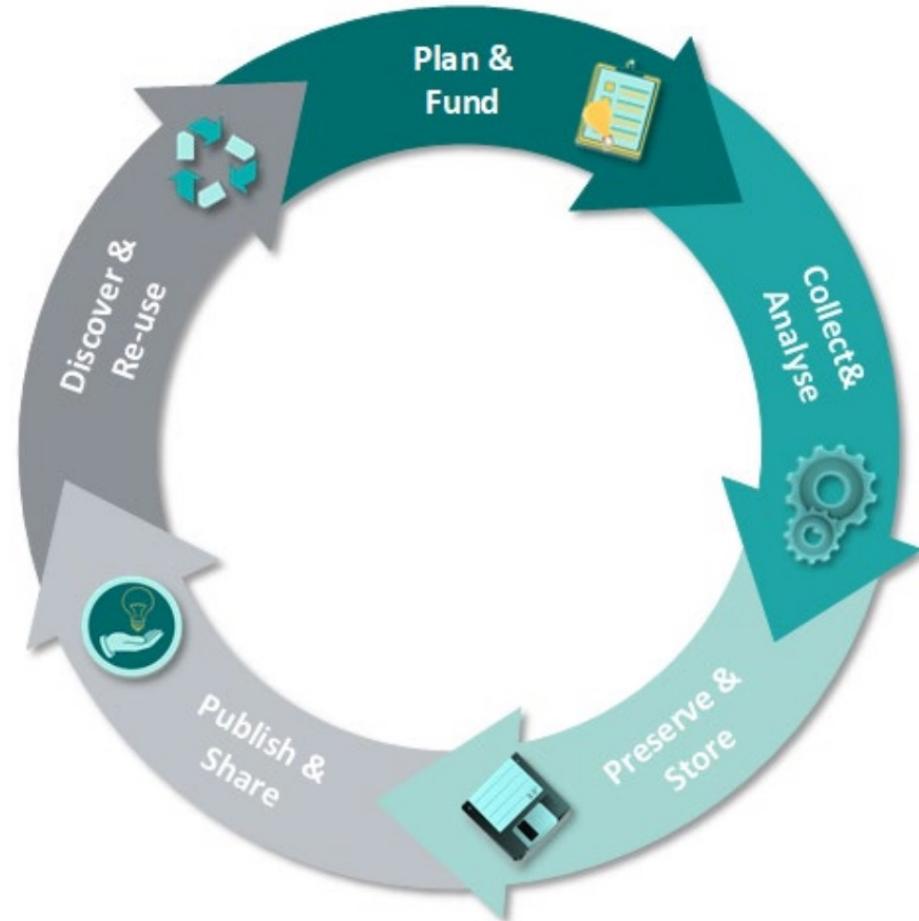
A README provides information about data file(s), granting better reusability.

## README Template

The screenshot shows a template for creating a README file. At the top, there is a logo for 'Georgia Tech Library'. Below the logo, the title 'README TEMPLATE' is centered. To the left, there is a sidebar with links to 'RESEARCH DATA MANAGEMENT', 'DATA MANAGEMENT PLAN' (with sub-links for 'Write a data management plan', 'Learn about data management requirements', and 'Request help with your plan'), 'ARCHIVE YOUR DATA' (with sub-links for 'Research data submission guidelines'), 'DATA MANAGEMENT TRAINING', and 'CONTACT US'. The main content area contains instructions for creating a README file, explaining its purpose and how it should be used. It also provides a list of numbered steps for what should be included in the file, such as file names, directory structure, and descriptions of parameters/variables.

<http://d7.library.gatech.edu/research-data/readme>

# Phase 4: Publish and Share



## Publish & Share

Publications based on data should include a data citation / reference how the data can be accessed

Selection of suitable repository and deposition

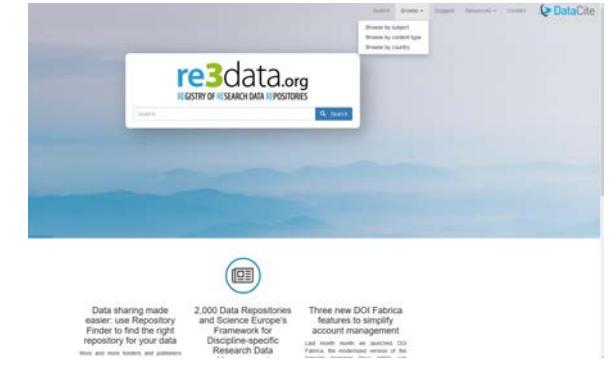
Assignment of Digital Object Identifiers

Decide on a data access model (“As open as possible, as restricted as needed”) and re-use conditions (e.g. CC-licenses)

## FAIR data

# How to find the best repository for your data:

- Repository **must support FAIR data** (Findable, Accessible, Interoperable, Reproducible)
- Choose **domain-specific** rather than general repositories
- Choose **common** rather than institutional repositories
- Choose what the **community** is using (foster re-use)
  - See example list of repositories and re3data.org



## SNSF uses this checklist to validate suitability:

- Datasets have globally unique and persistent identifiers (e.g. DOI)?
- Support intrinsic and submitter-defined metadata?
- Clarity under which license data will be available (e.g. CC0, CC BY, etc.)
- Citation information and metadata always publicly accessible (even for restricted datasets)
- Submission requests intrinsic metadata in a specific format (to ensure machine readability/interoperability)
- Repository has a long-term preservation plan for the archived data



# Zenodo: Everybody's general purpose repository

The screenshot shows the Zenodo homepage with a search bar, upload button, and communities section. Below is a list of recent uploads from the 'Research Data University of Basel' community. Each upload entry includes a thumbnail, title, authors, and a 'View' button.

- Recent uploads:**
  - Raw data to: "Persistent RNA virus infection is short-lived at the single cell level but leaves transcriptomic footprints"
  - Data\_Table S2\_Impact on Bile Acid Concentrations by Alveolar Echinococcosis and Treatment with Albendazole in Mice
  - Data\_Table S1\_Impact on Bile Acid Concentrations by Alveolar Echinococcosis and Treatment with Albendazole in Mice
  - Data\_Figure S5\_Impact on Bile Acid Concentrations by Alveolar Echinococcosis and

- Hosted by the CERN
- Free of charges
- Max 50GB/dataset (unlimited datasets)
- Automated DOI assignment
  
- OpenAIRE integration (EC reporting)
- GitHub integration
- ORCID integration
  
- All file formats accepted
- Usage statistics interface
- OAI-PMH protocol (content harvesting)
- 18 petabytes disk cluster
- Each file has 2 replicas on different servers
- 2 independent MD5 checksums per file
- Metadata 12-hourly backup cycle
- ...

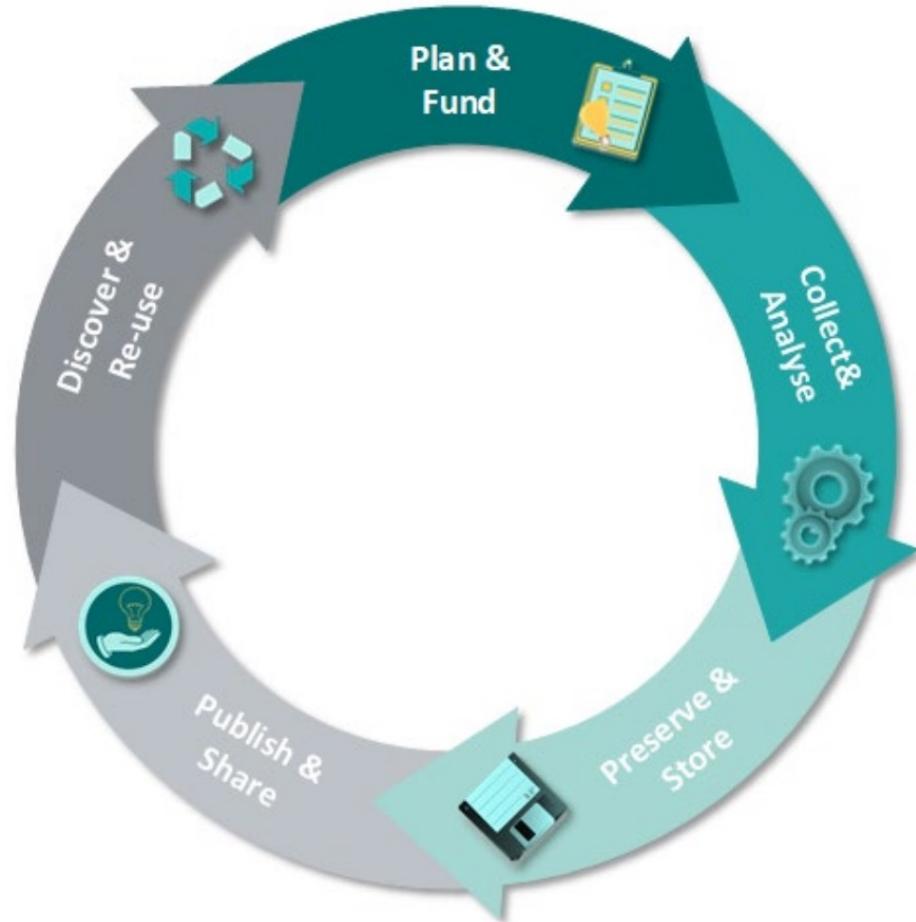
[sandbox.zenodo.org](https://sandbox.zenodo.org)

# EBI BioStudies: a new general repository for biological studies and their data

The screenshot shows the BioStudies website interface. At the top, there's a navigation bar with links to EMBL-EBI home, Services, Research, Training, About us, and EMBL-EBI. Below the header is a search bar with placeholder text "Search BioStudies" and an example "hyperplasia, PMC516016". To the right of the search bar are "Feedback" and "Login" buttons. The main content area has a blue header with the BioStudies logo. Below it, there are three filter sections: "Collection", "Released", and "Link Type". The "Collection" section lists various data sources like ArrayExpress, BioImages, BioImages-EMPIAR, EU-ToxRisk, Eurocan Platform, Europe PMC, HeCaToS, JCB, None, and SourceData. The "Released" section shows a list of years from 2014 to 2023 with corresponding counts. The "Link Type" section lists Array Design, DOI, ENA, FNA, GEO, and GATC. In the center, search results are displayed in a grid format. One result is highlighted: "Source Data of Manuscript EMBOJ-2022-112953" (E-MTAB-11704) dated 13 February 2023, which includes a link to "Tnseq to identify recipient factor to modulate conjugative plasmid transfer". Other results include "SNP analysis of J2530" (E-MTAB-12475), "Whole genome sequencing of caspofungin adaptors derived from homozygous deletion strains of PKC and calcineurin pathways in Candida albicans" (E-MTAB-10539), and "A surface morphometrics toolkit to quantify organellar membrane ultrastructure using cryo-electron tomography" (EMPIAR-11370). A footer at the bottom of the page contains a "Feedback" button.

- **Hosted by EBI**
- **Integration with other EBI databases**
- **Integration with PMC**
- **Link to data or contain the data itself**
- Recommended ELIXIR Deposition Database
- Deposition can be manual or automated
- Accompanies structured EBI databases
- Aims to accept all «other» data generated from a biological study as well as a study description
- ...

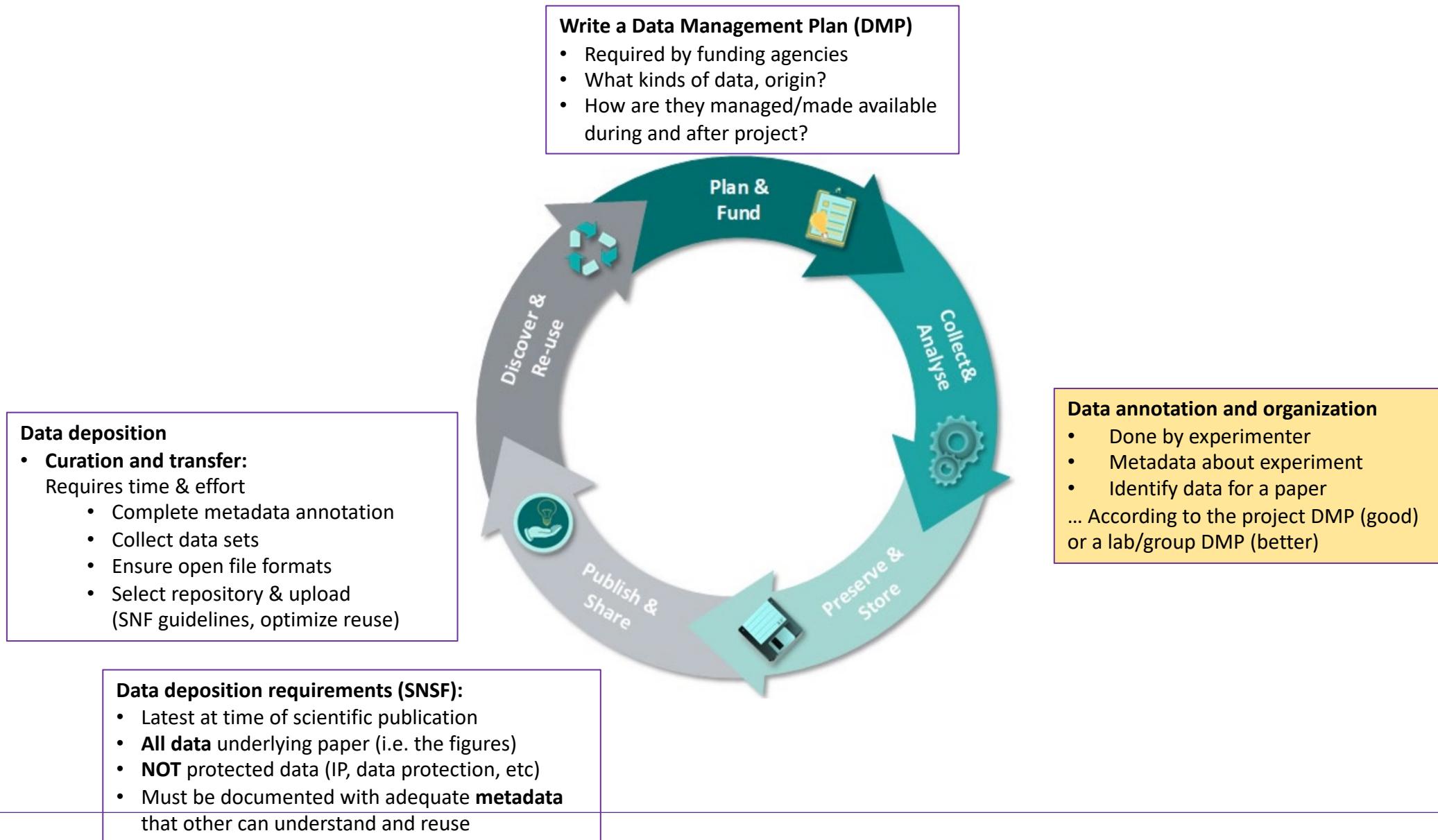
# Phase 5: Discover, Re-use, Learn and Teach



## Discover & Re-use

- Prepare new studies
- Reexamine existing data
- Mining data sets for new insights
- Use data sets for method development
- Training and Teaching with existing data sets
- Respect the usage license

# Quality checkpoints in the research data life cycle



# DATA CURATION:

# DOES IT SPARK JOY?

NETFLIX | NEW YEAR'S DAY

TIDYING UP  
WITH MARIE KONDO



# Thank you.

# Dr. Michael Podvinec

## Biozentrum Research IT



Image: <http://www.voicendata.com/wp-content/uploads/2015/07/Data-Management.png>

# Forgotten Data: A film by Simone Pengue

Prix Média Newcomer 2021 of the Swiss Academy of Sciences



# About this material/acknowledgements

This material is licensed under the CC-BY-SA license

- Exception:
  - Adobe Stock Images under license to University of Basel (Slide 2)

Adobe Stock Images licensed to University of Basel

This presentation is standing on the shoulders of others:

- University of Basel Research Data Network
- Aude Bax de Keating, EPFL Library (at the time)
- Others (references provided wherever possible)

# «Data available on request»

Current Biology

Volume 24, Issue 1, 6 January 2014, Pages 94-97

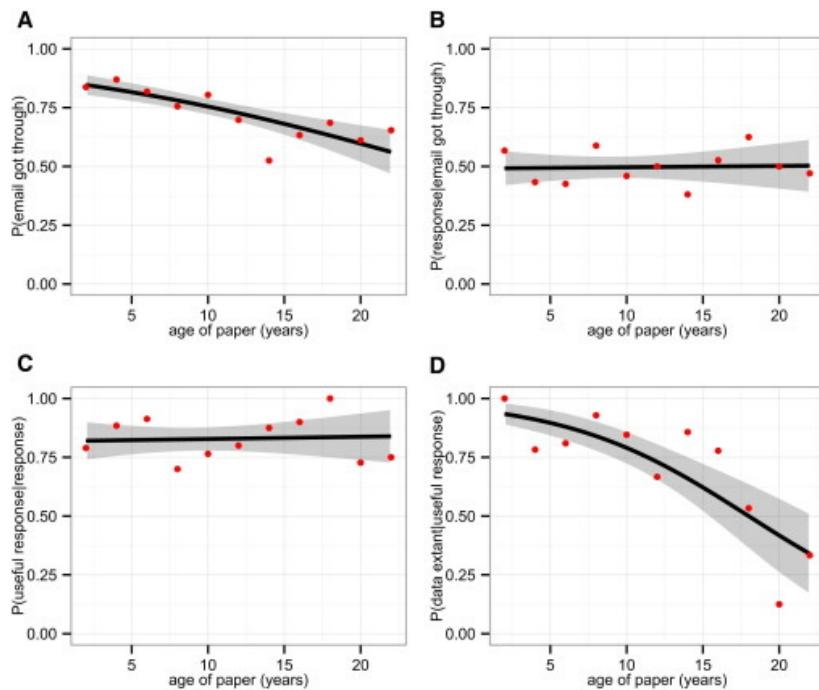


Report

## The Availability of Research Data Declines Rapidly with Article Age

Timothy H. Vines<sup>1, 2, 3, 4</sup>, Arianne Y.K. Albert<sup>3</sup>, Rose L. Andrew<sup>1</sup>, Florence Débarre<sup>1, 4</sup>, Dan G. Bock<sup>1</sup>, Michelle T. Franklin<sup>1, 5</sup>, Kimberly J. Gilbert<sup>1</sup>, Jean-Sébastien Moore<sup>1, 6</sup>, Sébastien Renaud<sup>1</sup>, Diana J. Rennison<sup>1</sup>

Show more ▾



“Policies ensuring that research data are available on public archives are increasingly being implemented(...). These policies are predicated on the idea that authors are poor stewards of their data, particularly over the long term [7], and indeed many studies have found that authors are often unable or unwilling to share their data [8, 9, 10, 11].

However, there are no systematic estimates of how the availability of research data changes with time since publication. We therefore requested data sets from a relatively homogenous set of 516 articles published between 2 and 22 years ago, and found that availability of the data was strongly affected by article age.

For papers where the authors gave the status of their data, the odds of a data set being extant fell by 17% per year. In addition, the odds that we could find a working e-mail address for the first, last, or corresponding author fell by 7% per year.

Our results reinforce the notion that, in the long term, research data cannot be reliably preserved by individual researchers, and further demonstrate the urgent need for policies mandating data sharing via public archives.”

# What is a Data Management Plan?

Required by all major funders (SNSF, Horizon Europe, NIH) - with minor variations

## Example: SNSF Data Management Plan questions

- 1.1 What data will you collect, observe, generate or re-use?
- 1.2 How will the data be collected, observed or generated?
- 1.3 What documentation and metadata will you provide with the data?
- 2.1 How will ethical issues be addressed and handled?
- 2.2 How will data access and security be managed?
- 2.3 How will you handle copyright and Intellectual Property Rights issues?
- 3.1 How will your data be stored and backed-up during the research?
- 3.2 What is your data preservation plan?
- 4.1 How and where will the data be shared?
- 4.2 Are there any necessary limitations to protect sensitive data?
- 4.3 I will choose digital repositories that are conform to the FAIR Data Principles.
- 4.4 I will choose digital repositories maintained by a non-profit organisation.

## Help:

- Websites:
  - [researchdata.unibas.ch](http://researchdata.unibas.ch)
  - <https://wiki.biozentrum.unibas.ch/x/X8KBAw>
- Uni Basel Research Data Network / Data Stewards

## A living document:

The direction of research is not completely predictable.

The data management challenges are expected to evolve during a project.

Likewise, the DMP can evolve. It has to be finalized at the conclusion of the project

# Data Management Requirements from Funders

## SNSF:

**DMPs required since 2017, format unchanged so far.**

Data policy: All data pertaining to publications within the project must be made available in recognized repositories meeting FAIR principles  
(see SNSF IR 11.8.1)

**DMP (for now) not part of the scientific evaluation**, but must conclusively describe data collected, generated and observed at the project end time. (SNSF IR 11.8.3, sep. guidelines).

**New: Must only be submitted upon successful scientific evaluation**

2025-2028 Priorities: Open research data, infrastructure and services + Skills one of four pillars.

**Note:** SNSF allows for CHF10'000 for data curation efforts

## EU FPs: FAIR DMPs

Horizon 2020 2014-2017: ORD Pilot (Opt-in) – 2017-2020: All projects

Horizon Europe: Open Data Policy (“as open as possible, as closed as necessary”)

## NIH:

2023 policy: All grants must provide DMS plan. Compliance can be reviewed during reporting intervals and provisions to terminate the agreement.  
Established repositories strongly encouraged

# Why do I need to keep old data and how long?

## Data retention serves you and others:

- Continue working internally with the same data / internal re-use
- Ensure reproducibility of research results
- Keep the data for legal requirements
  - To defend yourself against accusations of plagiarism, fraud, unethical conduct
  - Intellectual Property: Prove priority of invention
- Re-use within the scientific community
  - SNSF requires Data Management Plans with grant proposals
  - SNSF expects funded researchers to generate FAIR data
  - Publishers require publication of data underlying a publication

## How long do I have to keep my data?

- Uni Basel: 5 years minimum
- SNSF: 10 years
- -> 10 years is a good general rule

## Where do I keep/publish my data?

- Active storage?
- Archival systems / “Deep Storage”?
- Data repositories? -> Allow reproducibility and re-use by applying the FAIR principles

# A simple guide to repositories for common life science data types

Data type	Pre-publishing/Internal	Post-publishing/External	
Imaging data	OMERO	IDR Bioimage Archive* EMPIAR ??? (size-dependent)	<b>More:</b> See <a href="http://re3data.org">re3data.org</a> for other relevant research data repositories. The SNSF uses re3data as a reference when deciding on the FAIRness of a repository.
Proteomics/Mass Spectrometry	SampleQueue PUMA storage	PRIDE & others (during publication)	<ul style="list-style-type: none"> <li>re3data shows that no persistent identifier is used by these databases. This seems strange – and likely will be updated.</li> </ul>
Genomics	sciCORE HPC storage LabKey PUMA storage	NCBI and EBI sequence databases	** Zenodo accepts any research data sets up to 50GB per submission and is funded by the EU through OpenAIRE. There is a « <b>Research Data University of Basel</b> » community you can associate your research with.
“Everything else”: Western Blots qPCR Sample information Electrophysiology Video microscopy (...)	PUMA storage JUMBO storage Annotations in ELN LabKey Sample Manager(?) (...)	EBI BioStudies* Zenodo** Dryad***	*** Dryad is comparable to Zenodo (but from the US). It has a financial model for submissions > 50GB

# Naming your file

## File Naming Caveats

Please keep in mind that not all characters are allowed in file names and name length restrictions on file and path level also do apply.

We recommend to keep the path length **below 260 characters and only use safe characters**. Keep the file name at **32 characters or less**.

## Develop a file naming convention:

### Info tracked & the convention used

The researchers wanted to track several things about the tiles:

1. **Study site.** Indicated by the name, ex. FR3, FR7, FR9.
2. **Depth of the water.** Indicated by S (shallow), M (middle), or D (deep).
3. **Date.** Indicated by YYMMDD.
4. **Tile number.** Indicated on the tile.
5. **Tile treatment.** Indicated by C (caged) or U (uncaged).
6. **Number assigned to photo by camera.**
7. **Whether the post-removal photo was of the entire tile or a tile section.** Indicated by W (whole area), A (upper right), B (lower right), C (lower left), or D (upper left).

Example: FR3S.140623.129C.2653.W.JPG

This was image 2653 of whole, uncovered tile 129 from study site 3 in shallow water, taken on June 23, 2014.

### Safe characters:

A B C D E F G H I J K L M N O P a b c d e f g h i j k l m n o p 0 1 2 3 4 5 6 7  
8 9 - \_

### Don't use:

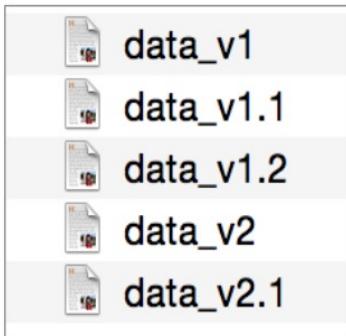
< > : " / \ | ? \*

Another example: <https://www.unibas.ch/de/Universitaet/Administration-Services/Generalsekretariat/Archive-Sammlungen/Universitaetsarchiv1.html>

# Versioning your file

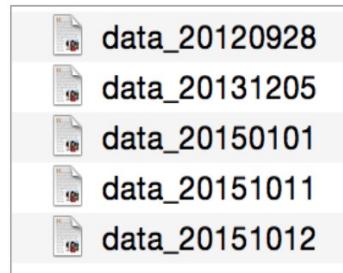
Try to:

**Sort**



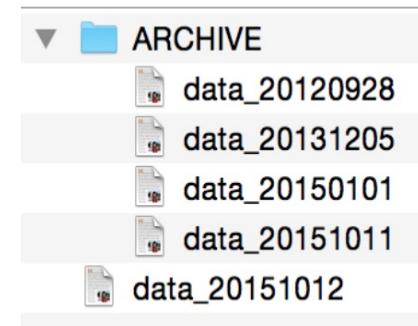
Major changes: ordinal numbers  
Minor changes: decimal numbers

**Distinguish**



Dates distinguish between the different versions.

**Separate**



Archive older files in a separate folder.

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Version control software:



# The FAIR Data Principles

