# Data Management for Transparent Research

Serena Bonaretti https://sbonaretti.github.io/

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## Data in life sciences

- Scope, size, and complexity of datasets has increased dramatically over the last 10-20 years
- Traditionally, the knowledge of how to manage this data is often limited to "information managers"
- In response to new funding requirements, researchers are asked to be part of data management

Griffin et al. 2019



GAGTO

SCCTCGTATTGGGAGTCCTCGGCG AGTAGCATTCGCCTCGCATCGGGA

GCAGTAGCATCAGCCTCGCA GCAGTAGCATCAGCCTCGCA expression Interactions functional domains biological non-coding available type function other metadata product structural using glycan terms thaliana sequences damage Piam other metadata product structural using glycan terms thaliana sequences damage Piam genetic acid Drug organism Via features DNA based sequencing PDB chromatin genome names microarray plant chemical factor RNA CDS resource site maps sites regions target RNAs domain about 60 all interaction molecular nematode KEGG annotation database modification structure disease predicted pathways collection protein including Ontology annotations

### Why do we need to manage data?

- We want to be able to reuse data produced by ourselves and by other researchers
- "We are losing data at a rapid rate, with up to 80% unavailable after 20 years" (<u>Griffin et al. 2019</u>)



### Data life cycle in research



## Collecting



- Repositories are indexed in registries of repositories (e.g. <u>re3data</u>, <u>FAIRsharing</u>, etc.)
- Data need to have accurate metadata
  - E.g. Use of vocabulary from ontologies (= set of categories that define objects and the relationships among them)

# Integrating



- Main issue: Standardization
  - Data can have different formats
- Possible solutions:
  - Linked Data and Semantic Web
  - Format converters

## Processing and Analyzing



- Processing and Analyzing are crucial for computational reproducibility and assessment of quality
  - Use of electronic notebooks with dependences
  - Virtual machines containing the whole computational environment (e.g. <u>Docker</u>)
- In the lab: Electronic laboratory notebooks
  - E.g. LabTrove, <u>BlogMyData</u>, <u>Benchling</u>, etc.

# Publishing



- Journals require more and more to share both raw data and derived data
  - Issues: Ethical constraints and commerciallyrelevant data
- New journals that publish papers describing datasets
  - Authors can get credit for labor-intensive and expensive data collections
  - E.g. <u>Data descriptor in scientific data</u>, <u>Data note</u> <u>in gigascience</u>, etc.

# Storing



- Local data storage during collection, integration, processing and analysis
  - E.g. <u>Dropbox</u>, <u>Google Drive</u>, local servers

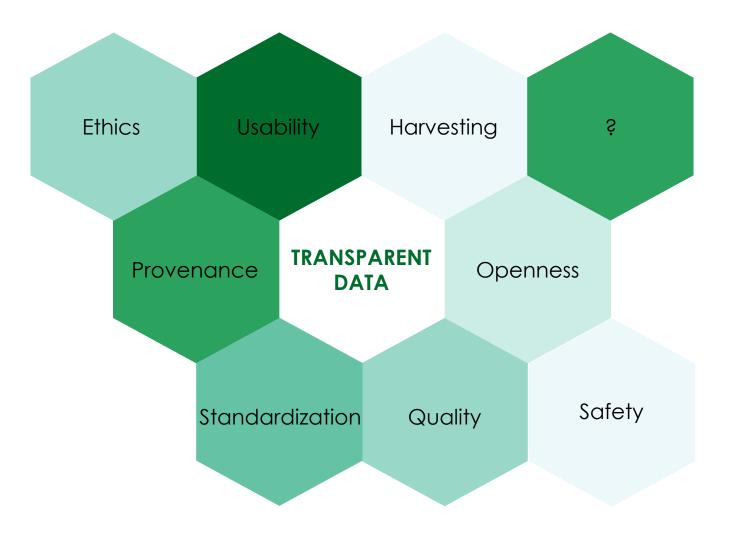
- Data repositories for sharing
  - E.g. <u>Zenodo</u>, <u>Figshare</u>, domain-specific repositories

# Sharing and Finding

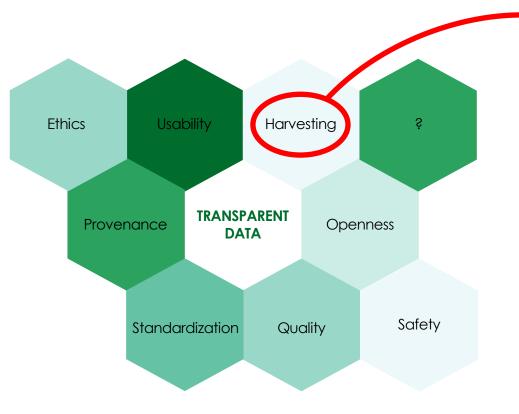


- Characteristics of open datasets:
  - Data are hosted in a public repository
  - Data are accompanied by descriptive metadata
  - Data have unique identifiers (e.g DOI)
- Issues:
  - Data quality
  - Ethical constraints

### Topics in data management



# The FAIR guiding principles



Findable Accessible Interoperable Reusable

#### To be Findable:

- F1. (meta)data are assigned a globally unique and persistent identifier
- F2. data are described with rich metadata (defined by R1 below)
- F3. metadata clearly and explicitly include the identifier of the data it describes
- F4. (meta)data are registered or indexed in a searchable resource

#### To be Accessible:

- A1. (meta)data are retrievable by their identifier using a standardized communications protocol
- A1.1 the protocol is open, free, and universally implementable
- A1.2 the protocol allows for an authentication and authorization procedure, where necessary
- A2. metadata are accessible, even when the data are no longer available

#### To be Interoperable:

- 11. (meta)data use a formal, accessible, shared, and broadly applicable language for knowledge representation.
- I2. (meta)data use vocabularies that follow FAIR principles I3. (meta)data include qualified references to other (meta)data

#### To be Reusable:

- R1. meta(data) are richly described with a plurality of accurate and relevant attributes
- R1.1. (meta)data are released with a clear and accessible data usage license
- R1.2. (meta)data are associated with detailed provenance
- R1.3. (meta)data meet domain-relevant community standards

#### Wilkinson et al. 2016

### Academic discussion about FAIR

#### • What FAIR is

- It "refers to a set of principles, focused on ensuring that research objects are reusable" (<u>Mons et al. 2017</u>)
- "They deliberately do not specify technical requirements" but "They describe characteristics and aspirations for systems and services" (<u>Mons et al. 2017</u>)
- My interpretation: They are principles to organize metadata in such a way that machines can harvest data for us

#### • What FAIR is **not**

- It is not a standard, it is not specific to life sciences, it is not equal to semantic web (Mons et al. 2017)
- It does not imply open data (<u>Mons et al. 2017</u>)
- My interpretation: It is limited to data harvesting, i.e. it does not provide guidelines for format standardization, data quality, etc.