## **Open Science**

**Publicly Available Resources** 

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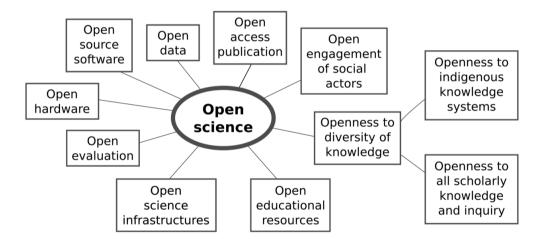


#### **Outline**

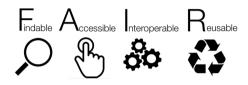
- What is Open Science?
- FAIR Data and Software
- Challenges sharing and reusing data
- Standardisation and Ontologies
- Publicly available resources

# What is Open Science Impact, Contribution, Trust

- Make scientific research accessible to all levels of society:
  - Publications
  - Samples
  - Methods
  - Software
  - Data
- Advantages:
  - Reproducibility and replicability
  - Societal **responsibility** publicly funded, publicly available
  - Multi-purpose of research outputs
- Disadvantages: concerns of data misuse



#### **FAIR Data and Software**



- Findable and Accessible
  - Add enough metadata data about your data

Minimum Information for Biological and Biomedical Investigations

• Deposit your data in **public repositories** or make them available in **databases** 

Zenodo Figshare Pride Metabolights GEO GitHub

- Interoperable:
  - Use standard and open formats
  - Provide all data needed to reproduce your analysis
- Reusable:
  - Describe your data well, e.g., good metadata but also

Provide README files describing the data Use descriptive column headers for the data tables

Attach a license

### **Challenges Sharing and Reusing**

The marshmallow test — delayed gratification

- Open does not mean FAIR
- Requires an effort
- Metadata becomes the most important data
- In many cases there are no standards or multiple ones
- Most of the data out there not FAIR







https://imgflip.com/memegenerator

https://en.wikipedia.org/wiki/Stanford\_marshmallow\_experiment

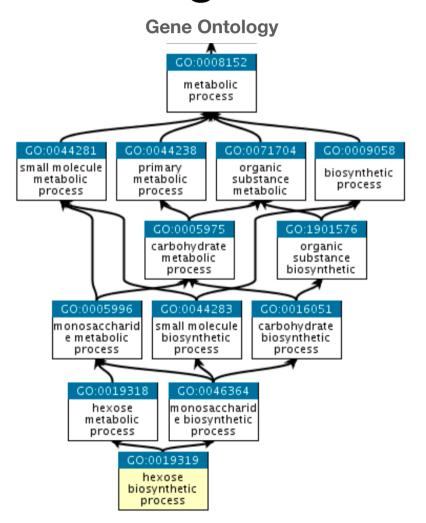
### Standardisation and Ontologies

- Data standardisation requires defining terminologies and vocabularies that:
  - Assign unique identifiers to entities/concepts such as proteins, genes, diseases
  - Describe those entities/concepts and provide meaning
  - Relate those concepts to other terms
  - Classify those entities/concepts into categories
- Solution —> Ontologies
- Ontology:

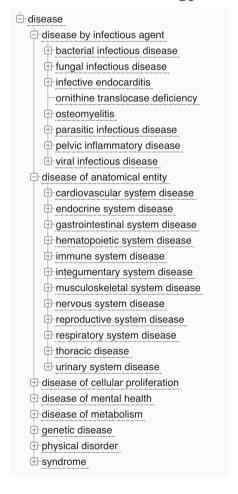
formal way of representing knowledge in which concepts are described both by their meaning and their relationship to each other

A collection of terms and their definitions for a specific domain

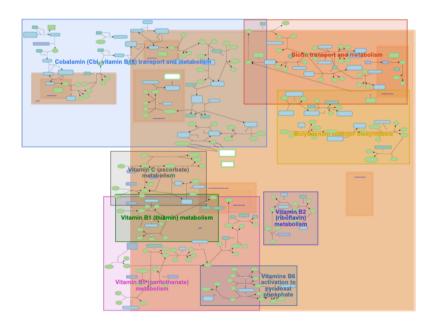
#### **Ontologies**



#### **Disease Ontology**



#### **REACTOME Pathways**



https://www.ebi.ac.uk/ols/ontologies

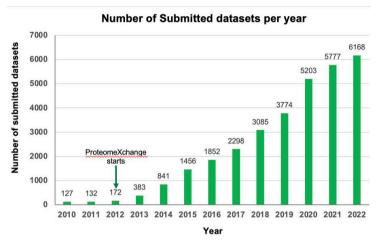
https://reactome.org/

http://geneontology.org/



- Do not reinvent the wheel
- Extend the life and purpose of publicly available data
- Build in-silico hypotheses before jumping into experiments (cheaper, higher success rate)
- Download Use Test Transform Upload
- Growing number of resources and datasets available





#### **Examples of Microbes Resources**

ALEdb 1.0: a database of mutations from adaptive laboratory evolution experimentation <a href="https://aledb.org/">https://aledb.org/</a>

MiMeDB: the Human Microbial Metabolome Database <a href="https://mimedb.org/">https://mimedb.org/</a>

Web of microbes (WoM): a curated microbial exometabolomics database for linking chemistry and microbes <a href="https://metatlas.nersc.gov/wom/project-begin.view">https://metatlas.nersc.gov/wom/project-begin.view</a>

mBodyMap: a curated database for microbes across human body and their associations with health and diseases <a href="https://mbodymap.microbiome.cloud/">https://mbodymap.microbiome.cloud/</a>

MicroPhenoDB Associates Metagenomic Data with Pathogenic Microbes, Microbial Core Genes, and Human Disease Phenotypes <a href="http://www.liwzlab.cn/microphenodb">http://www.liwzlab.cn/microphenodb</a>

BacDive in 2022: the knowledge base for standardized bacterial and archaeal data https://bacdive.dsmz.de/

MASI: microbiota—active substance interactions database <a href="http://www.aiddlab.com/MASI/">http://www.aiddlab.com/MASI/</a>

iModulonDB: a knowledgebase of microbial transcriptional regulation derived from machine learning <a href="https://imodulondb.org/index.html">https://imodulondb.org/index.html</a>

MIBiG 3.0: a community-driven effort to annotate experimentally validated biosynthetic gene clusters <a href="https://mibig.secondarymetabolites.org/">https://mibig.secondarymetabolites.org/</a>