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# DATA PUBLICATION 101



## **OUTLINE**

- What is data publishing?
- Why data sharing?
- Data sharing policy
- Data sharing using repositories
  - What is a data repository?
  - Examples
  - How repositories lead to FAIR and/or open data (metadata, persistent identifiers, standards)
  - How to select a data repository?
  - How to use a data repository
- Data re-use, data citation, licenses
- Biblio data register (refer to demos)



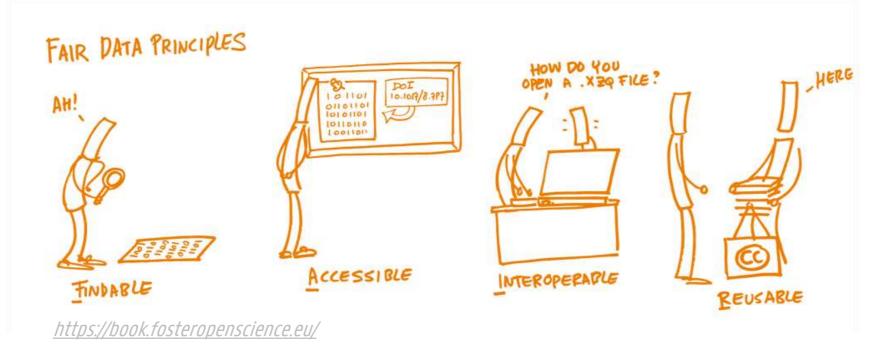
#### **PUBLISHING DATA**

- Sharing data from
  - → completed (parts of) research projects
  - → outside your project or research team.
- Not
  - exchanging data with collaborators while your research is active.
  - Share data with your promotor



### RATIONALE FOR DEVELOPING FAIR PRINCIPLES

- Urgent need to improve the infrastructure supporting the reuse of research data
- Specific emphasis on enhancing the ability of machines to automatically find and use the data, in addition to supporting its reuse by individuals
- The FAIR principles are a guideline to enhance the data reusability





# WHY SHARE DATA?



#### THE BENEFITS

#### FOR RESEARCHERS:



More visibility and citations.



Opportunities for collaboration.



Career recognition.



Decrease of non-compliance risks (legal, ethical, institutional and funders' policies).

#### FOR SCIENCE:



Facilitates data finding and re-use.



Enables new research and new insights on the data.



Valuable data is protected.



Supports research integrity and reproducibility.

#### FOR SOCIETY:



Efficient use of public resources.



Better quality research can lead to better decision-making.



Opportunities for citizen science.

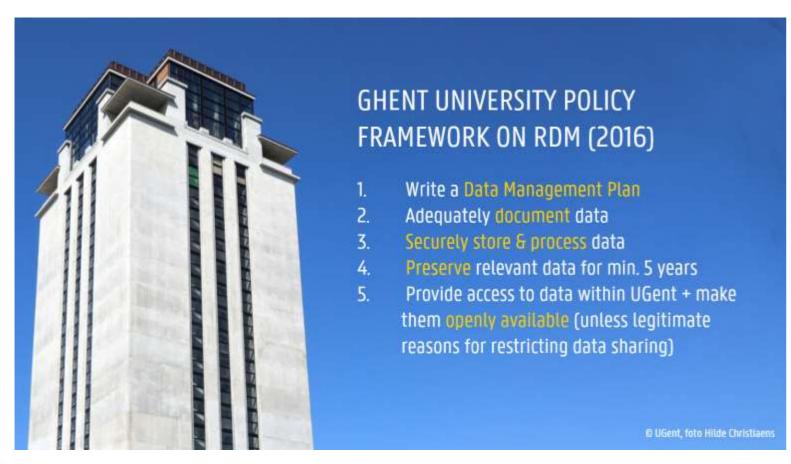


Increased transparency and trust in science.



## POLICIES AND REQUIREMENTS ON DATA SHARING

# Institutional policies





"PLOS journals <u>require authors to make all data necessary to replicate</u>

<u>their study's findings publicly available without restriction at the time of</u>

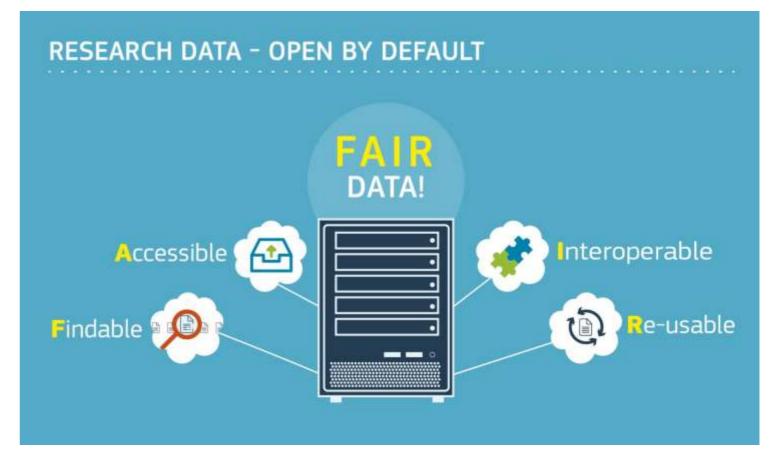
<u>publication</u>. When specific legal or ethical restrictions prohibit public

sharing of a data set, authors must indicate how others may obtain

access to the data." <u>PLOS ONE Data availability</u>



#### **HEU Open Science requirements**





# ALLEA European Code of Conduct for Research Integrity



- Access as open as possible, as closed as necessary
- Data as legitimate and citable products of research



- RDM = key part of FWO policy for all support channels
  - → "Sound data management is required for all research supported by the FWO."
- Specific expectations:
  - 1. preserve data for a minimum of 5 years after end of research
  - 2. submit a Data Management Plan
  - 3. encourages researchers to ensure that their data management complies as much as possible with the 'FAIR' principles
  - 4. advises researchers to apply the principle 'as open as possible, as closed as necessary'
- Reporting
  - → the FWO obliges to register all relevant output via the institutional 'CRIS' systems





# 3. Valuation of diverse research outputs

#### Research applications

- Adapted application form: Space for narrative, CV with a broad set of activities, 5 main publications and/or achievements, other scientific output and impact
- Adapted scoring grid for evaluation

#### Scientific reporting

- Specific question: 'Are one or more referable datasets linked to this publication or to the project?'
- Questions about metadata and accessibility



# WAYS OF SHARING DATA

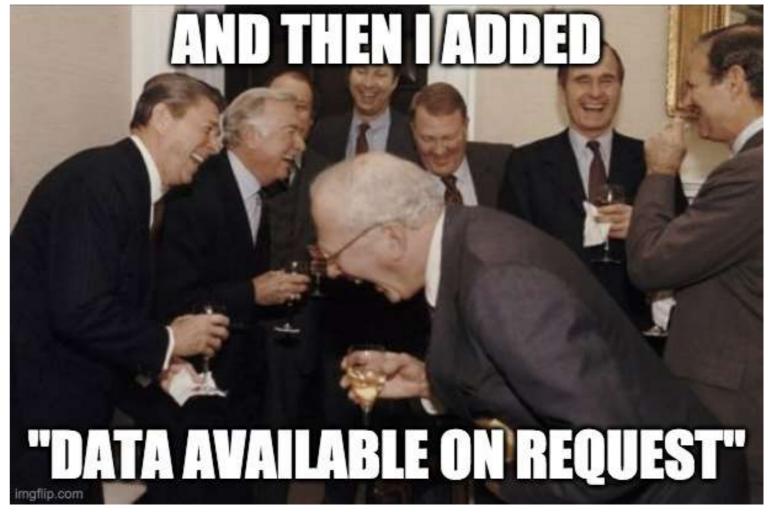


### DATA UPON REQUEST

Often unavailable

Recent research has found that as little as 6,8% of data was readily made available upon request

Only 14% responded



https://figshare.com/articles/presentation/Handson\_fMRI\_code\_and\_data\_sharing/12911165/1



## WAYS OF SHARING DATA







Article suppl. material



Data repository or archive





### DATA LINKED TO ARTICLE PUBLICATION

#### **Data availability**

Raw data of scRNA-seq (and output files from CellRanger) and spatial transcriptomics (Visium and stereo-seq) are deposited at the GEO under accession number <u>GSE207592</u>. Processed count matrices and cell annotations (scRNA-seq and Visium) are available online (<a href="https://marinelab.sites.vib.be/en">https://marinelab.sites.vib.be/en</a>). Molecular cartography data are available at Zenodo (<a href="https://zenodo.org/record/6856193#.Ytj-fnZBz-g">https://zenodo.org/record/6856193#.Ytj-fnZBz-g</a>). All other data supporting the findings of this study are available on reasonable request. <a href="mailto:Source data">Source data</a> are provided with this paper.



# FAIR DATA IN REPOSITORIES



## WHAT IS A DATA REPOSITORY?

A database infrastructure that collects, manages and stores data for preservation and sharing



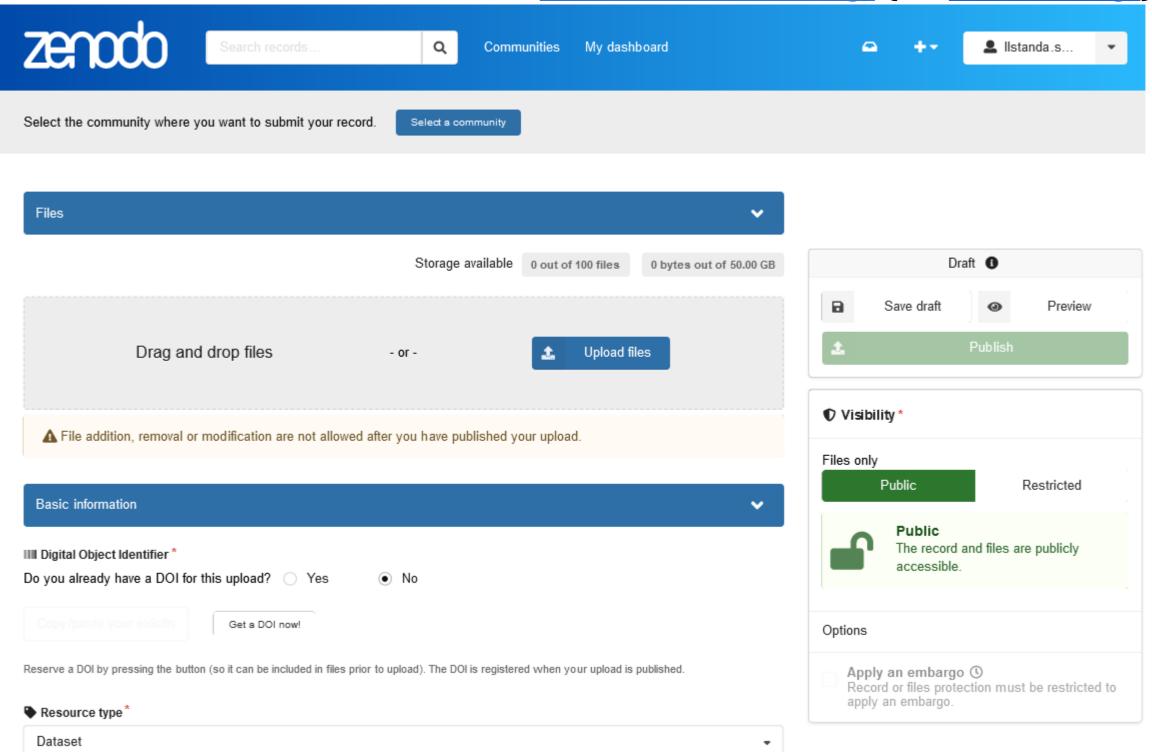






### **EXPLORING DATA REPOSITORIES**

Some repositories have a test environment: <u>sandbox.zenodo.org/</u> (vs. <u>zenodo.org/</u>)



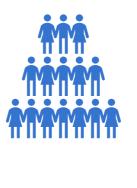


■ Title \*

### WHAT IS A DATA REPOSITORY?

A database infrastructure that collects, manages and stores data for preservation and sharing



















### DATA REPOSITORIES & FAIR DATA

- → Key infrastructures that provide functionalities to make your data FAIR.
- → Shared responsibilities between repositories and researchers in making data FAIR

FAIR principle	In practice	Repository	Researcher
FINDABLE	Assign a persistent identifier (PID)	•	
	Data are described with <u>rich</u> metadata	•	
	Metadata include the PID, are indexed & searchable online	•	
ACCESSIBLE	(Meta)data retrievable via PID using a standardised communications protocol (e.g. html)	•	
	Authentication and authorisation procedure	•	
INTEROPERABLE	Metadata standards	•	
	Controlled vocabularies	•	
	References to other (meta)data	•	
REUSABLE	(Machine readable) license / terms of use	•	•
	Detailed provenance documentation		
	Domain standards	•	



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	Controlled vocabularies			
	References to other (meta)data			
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	Detailed provenance documentation			
	Domain standards			



# PERSISTENT IDENTIFIERS

	Identifier	Globally unique identifier (guid, uuid)	Persistent identifier (PID)
What	a name that identifies a unique object gene identifier ENSG00000139618 book identifier ISBN 817525766-0	ensures that there are no two identical identifiers that point to different digital objects 123e4567-e89b-12d3-a456-426614174000	ensures that the digital object will remain findable over time and reduces the risk of broken links
How	can be (arbitrarily) assigned by anyone	by using a mechanism (e.g. using a registry service) for assigning identifiers	is determined by the commitment of the organisation that assigns and manages the identifier

#### Examples of globally unique, persistent identifiers

Identifier:



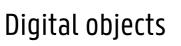




Organisation:







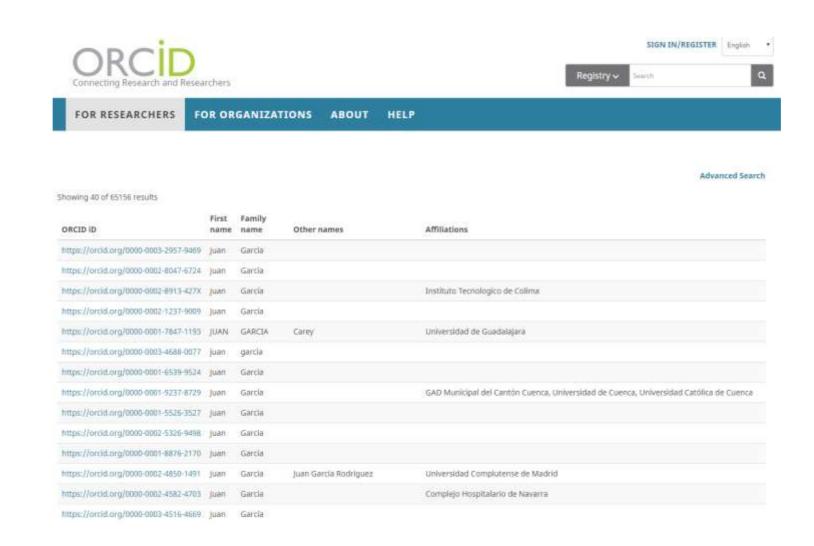
Research organisations

Researchers

# ORCID FOR IDENTIFICATION OF RESEARCHERS



Connection to contributions and affiliations



Unambiguous identity



Commonly used and required

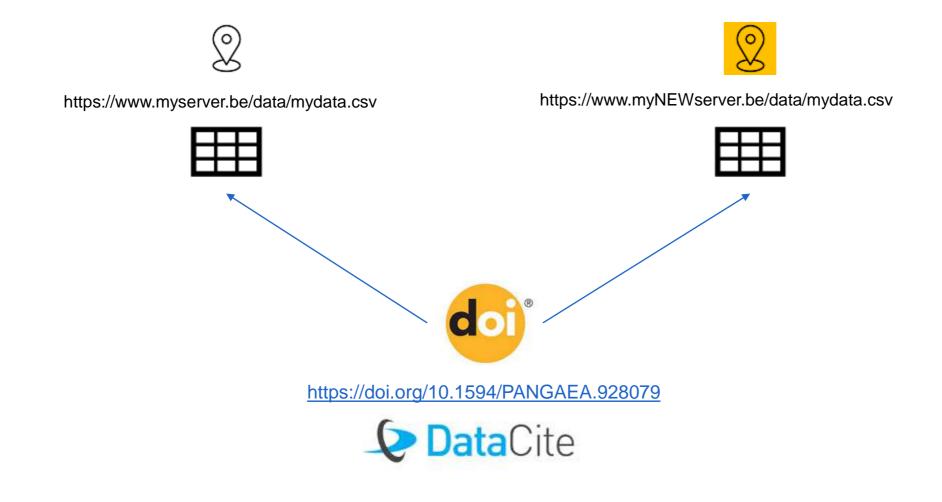


#### PID: PERSISTENT IDENTIFIER

**Globally unique:** an unambiguous reference for humans or machines, an identifier associated with only one resource at any time

**Persistent:** stable and resolvable identifier for the long term

e.g. DOI: Digital Object Identifier





### DATA REPOSITORIES & FAIR DATA

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	Authentication and authorisation procedure		•	
INTEROPERABLE	Metadata standards		•	
	Controlled vocabularies	Metadata	•	•
	References to other (meta)data			•
REUSABLE	(Machine readable) license / terms of use			•
	Detailed provenance documentation			•
	Domain standards			









WHERE





WH0



**DOCUMENTATION** 

Any contextual information that allows data to be understandable and reusable.

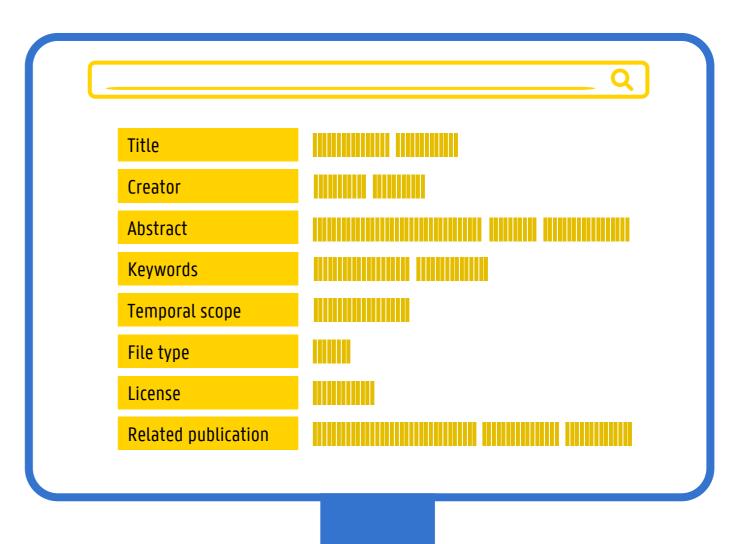


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**GENT** 

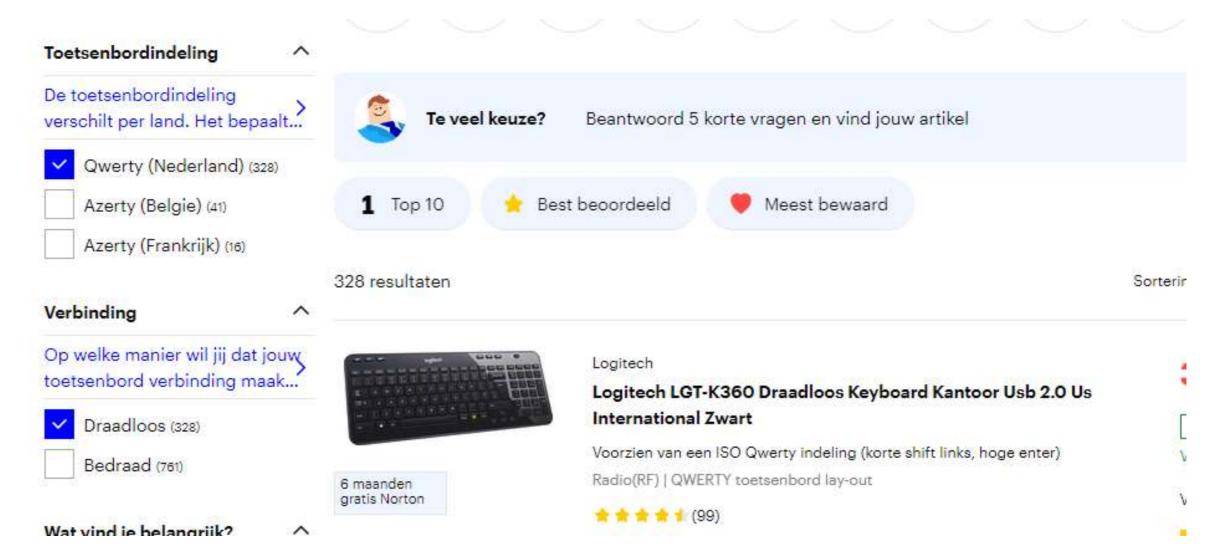
## **METADATA**

Information about data, recorded in a highly structured way, using a set of defined information fields or elements to be machine readable.



#### METADATA IS DATA...

- About data
- A (very) structured type of documentation: records information in sets of pre-defined fields or elements.
- Meant to be read & exchanged by machines
- To facilitate (online) data discovery
- To quickly judge usefulness of data

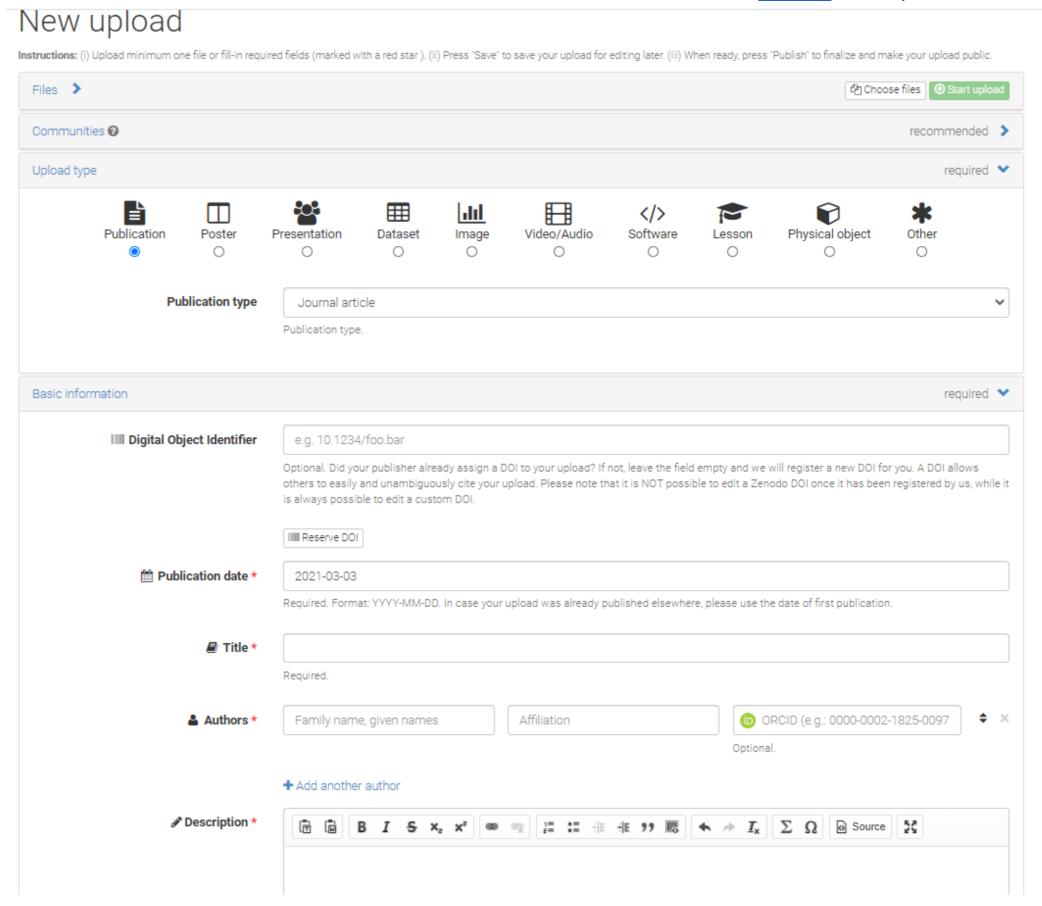




#### Zenodo data upload form

- Repositories: infrastructure to allow us to share & find data
- Provide services to create & manage metadata
- Typically, by filling a form with
  - Pre-defined fields
  - Lists of controlled terms to choose from (e.g. standard licenses)
- Metadata adheres to a specific metadata standard (interoperability)





### DATA REPOSITORIES & FAIR DATA

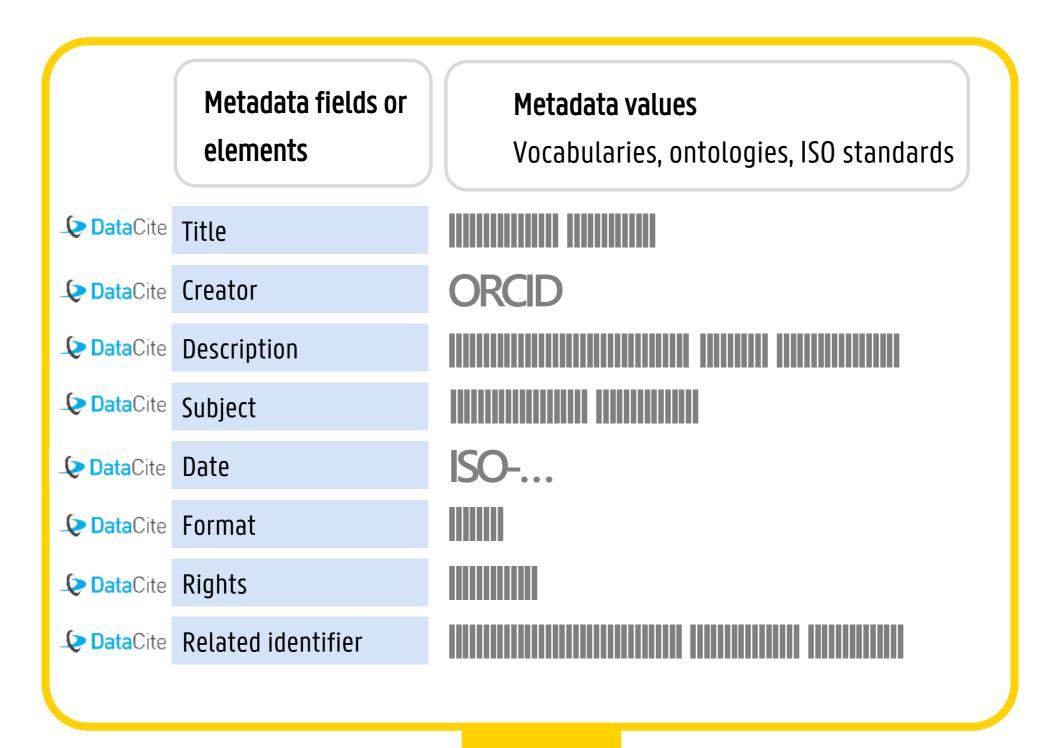
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	References to other (meta)data		•	•
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	Detailed provenance documentation			•
	Domain standards			



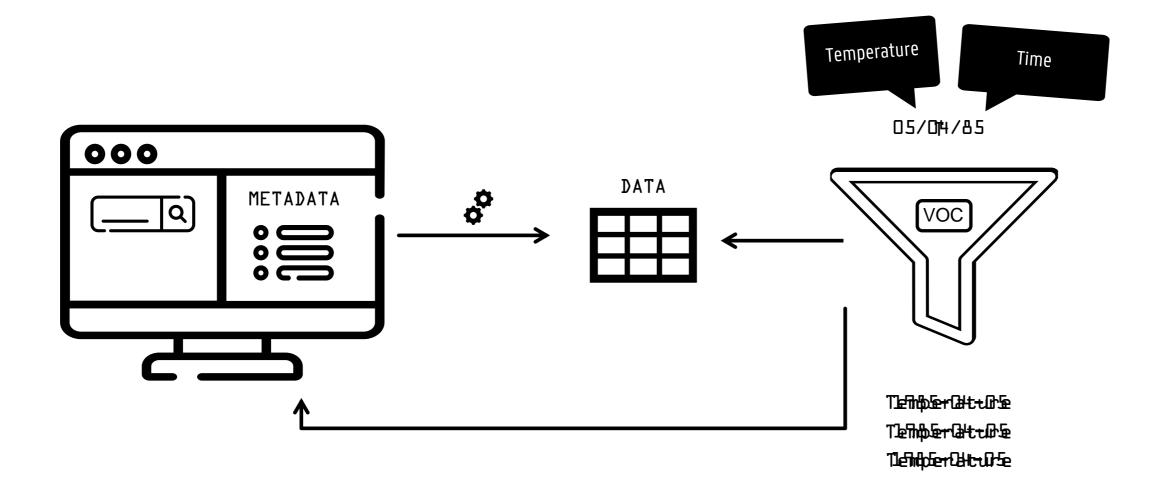
### **VOCABULARIES AND STANDARDS**

#### STANDARDIZED METADATA





## CONTROLLED VOCABULARIES TO STANDARDIZE TERMINOLOGY



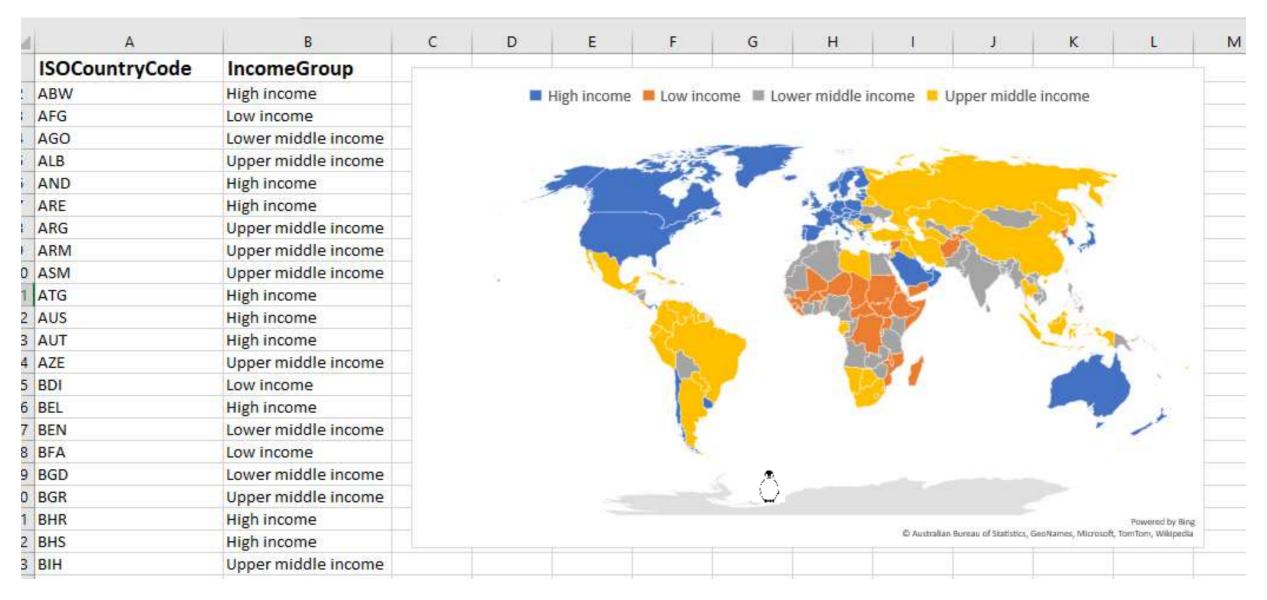
Temperature	temp	Т
Humidity	hum	Н

Controlled vocabulary



#### STANDARD VOCABULARIES & ONTOLOGIES

- Enable machine readability (use might be required in some domain repositories)
- Day-to-day example:





MS Excel recognizes <u>ISO Country codes</u> (or other place names in geonames.org) to automatically create maps

# FINDING REPOSITORIES



### WHICH TYPES OF REPOSITORIES EXIST?

#### Types of repositories

**GENERIC REPOSITORY** 













## **HOW TO SELECT A DATA REPOSITORY?**

#### Recommended repositories

RESEARCH DOMAIN'S CONSENSUS

RESEARCHERS'
BEST PRACTICES

JOURNAL RECOMMENDATIONS

DATA REPOSITORY REGISTRIES











e.g. <u>data repositories</u> recommended by Nature

re3data.org demo



### HOW TO SELECT A DATA REPOSITORY?

#### — Does your publisher or funder recommend a repository?

For

example:

#### **Data Repository Guidance**

Scientific Data mandates the release of datasets accompanying our Data Descriptors,

#### View data repositories

- Biological sciences: Nucleic acid sequence; Protein sequence; Molecular & supramolecular structure; Neuroscience; Omics; Taxonomy & species diversity; Mathematical & modelling resources; Cytometry and Immunology; Imaging; Organism-focused resources
- Health sciences
- Chemistry and Chemical biology
- Earth, Environmental and Space sciences: Broad scope Earth & environmental sciences; Astronomy & planetary sciences; Biogeochemistry and Geochemistry; Climate sciences; Ecology; Geomagnetism & Palaeomagnetism; Ocean sciences; Solid Earth sciences
- Physics
- Materials science
- Social sciences
- Generalist repositories



#### HOW TO SELECT A DATA REPOSITORY?

#### — Does your publisher or funder recommend a repository?





For

example:

**Deposit data in a trusted repository** and provide **open access** to it ('as open as possible, as closed as necessary')

#### **ELIXIR Deposition Database list**

Deposition Database	Data type	International collaboration framework <sup>1</sup>
ArrayExpress	Functional genomics data. Stores data from high-throughput functional genomics experiments.	
BioModels	Computational models of biological processes.	
BioSamples	BioSamples stores and supplies descriptions and metadata about biological samples used in research and development by academia and industry.	NCBI BioSamples database
BioStudies	Descriptions of biological studies, links to data from these studies in other databases, as well as data that do not fit in the structured	



#### GENERIC VS. DOMAIN-SPECIFIC REPOSITORIES

P All versions

File Type

□ Pdf (21859)

☐ Png (7726)

☐ Html (5956)

□ Jpg (3216)

□ Zip (1604)

☐ Hdf5 (624)



Partners Related resources Bulk downloads Submit data

Viral Sequences Host Sequences Expression Proteins Biochemistry **Imaging** Literature Viral sequences Q Raw and assembled sequence and analysis of SARS-CoV-2 and other coronaviruses Search Search Found 1318 results. 2 3 4 5 6 7 8 9 > Access Right Examples: ACE2, Severe acute respiratory syndrome 2... April 20, 2020 (Verson 1.0) Dataset Open Access ☑ Open (41069) COVID-19 CT Lung and Infection Segmentation Dataset □ Closed (8886) powered by ENA Showing 15 of 72,683 in Viral sequences > Sequences 🔞 Ma Jun; 👩 Ge Cheng; 🚳 Wang Yixin; 🚳 An Xingle; 🚳 Gao Jiantao; 🚳 Yu Ziqi; 🚳 Zhang Minqing; 🚳 Liu 🕽 ☐ Restricted (107) Shucheng; @ Wei Hao; @ Mei Sen; @ Yang Xiaoyu; @ Nie Ziwei; @ Li Chen; Tian Lu; @ Zhu Yuntao; Zhu Qi ☐ Embargoed (23) This dataset contains 20 labeled COVID-19 CT scans. Left lung, right lung, and infections are labeled by two Phylogeny Edit table view experienced radiologist. To promote the studies of annotation-efficient deep learning methods, we set up t Data types tasks based on All (737,152) Uploaded on April 20, 2020 Sequences (72,683) Accession Collection date Country Host Strain Isolate Location Reference sequences (2) April 24, 2020 (v2) Dataset Open Access Wuhan-Hu-1 MN908947 🚹 Dec, 2019 China Homo Raw reads (336,995) Linked COVID-19 Data: Ontology sapiens Sequenced samples Florian Thiery; (314,481)Linked COVID-19 Data Ontology maintained at https://github.com/Research-Squirrel-Engineers/COVID-19 L Studies (277) 10.5281/zenodo.3757283 10.5281/zenodo.3757279 Genes (22) Uploaded on April 24, 2020 Browser (1) LR991698 Sep 21, 2020 United Homo is record Variants (12,691) Kingdom sapiens B.1.1.7 COVID-19 datasets in Zenodo Organisms Severe acute respiratory MT612306 SARS-CoV-May 23, 2020 Australia Homo syndrome coronavirus 2 2/human/AUS/VIC1725/2020 sapiens (72,652)Canine respiratory coronavirus (23) Human coronavirus OC43 MW185086 COVID-19 data portal Aug 5, 2020 Australia Homo **GENT** sapiens

## EXAMPLE OF A DOMAIN-SPECIFIC REPOSITORY

### **COVID-19 DATA PORTAL**



About ▼ Partners Related resources Bulk downloads Submit data

Showing 15 of 4,050,433 in Viral sequences > Sequences



Data types All (8,859,658)	Down	load Phylog	eny					Edit table view
Sequences (4,050,433) Reference sequences (2)		Accession	Lineage	Cross-references (1)		Collection date	Country	Center name
Raw reads (3,840,601) Systematic Analyses		MN908947 🕦	В	Viral sequences > Genes (12)	See all 🗸	Dec, 2019	China	
(905,802) Studies (957)		LR991698 <b>(1)</b>	B.1.1.7 Alpha	BioSamples (2) ☑	See all 🗸	Sep 21, 2020	United Kingdom	COVID-19 Genomics UK Consortium
Genes (22)		MT539159	B.1.1.25	Proteins > Protein sequences (1)	See all 🗸	May 21, 2020	Bangladesh	
Browser (1) Variants (61,840)		MT558684	B.1	Proteins > Protein sequences (1)	See all 🗸	Apr, 2020	USA	
Release Date 🐧 🗸		LR898760	B.1.160.16	ENA Study (1) 년		2020	Switzerland	ETH ZURICH D-BSSE
		LR898761	B.1.177	ENA Study (1) 년		2020	Switzerland	ETH ZURICH D-BSSE
Collection date 🐧 🗸		LR898762	B.1.258	ENA Study (1) 년		2020	Switzerland	ETH ZURICH D-BSSE
Last modification		LR898763	B.1.177.23	ENA Study (1) 년		2020	Switzerland	ETH ZURICH D-BSSE
date 🐧		LR898764	B.1.177	ENA Study (1) 년		2020	Switzerland	ETH ZURICH D-BSSE
Organisms		LR898765		ENA Study (1) 년		2020	Switzerland	ETH ZURICH D-BSSE
Severe acute respiratory		LR898766	B.1.1.39	ENA Study (1) 🗗		2020	Switzerland	ETH ZURICH D-BSSE
syndrome coronavirus 2 (4,048,086)		MW331608	B.1	BioSamples (2) ☑	See all 🗸	May 5, 2020	USA	Texas Department of State Health Services-SARS-CoV-
Scotophilus bat coronavirus 512 (442)		KC117208		Literature > Coronaviruses (4)	See all 🗸	Jul, 2010	Mexico	PREDICT Consortium
☐ Murine coronavirus (407)		KC117209		Literature > Coronaviruses (4)	See all 🗸	Jul, 2010	Mexico	PREDICT Consortium
More >		KC117210		Literature > Coronaviruses (4)	See all 🗸	Sep, 2010	Mexico	PREDICT Consortium
Center name	<							>
COVID-19 Genomics UK	Showing	15 ▼ results						Previous Page 1 of 270,029 Next



Consortium (1,585,305)

CDC-OAMD (493,599)

Robert Koch Institute

## GENERIC VS. DOMAIN REPOSITORIES

- <u>Domain-specific repositories</u>
  - Visibility: where your peers will likely look for data
  - Use of discipline standards & vocabularies
    - Search filters, visualization tools tailored to domain
    - Interoperability: easier to integrate multiple datasets
  - Dataset curation by repository advisors often part of submission process
- Generic repositories:
  - Provide rich metadata to make it more findable & reusable







- Generic AND domain:
  - DOI, link to publication(s), dataset(s) or other outputs
  - Support minimum degree of FAIRness



## DOMAIN SPECIFIC: LIFE SCIENCES





Search...



Tips on how to search

ABOUT

SUBMISSION

**BROWSE** 

ACCESS

DOWNLOAD N

**METADATA** 

Helpdesk





## Study

#### WTCCC case-control study for Coronary Artery Disease

Study ID	Alternative Stable ID	Type
EGAS00000000003		GWAS

# PROTECTION DE DE LA COMPANSA DEL COMPANSA DE LA COMPANSA DEL COMPANSA DE LA COMPA

#### **Study Description**

WTCCC genome-wide case-control association study for Bipolar disorder (CAD) using the 1958 British Birth Cohort and the UK National Blood Service collections as controls.

## VIII) U.S. National Library of Medicine Clinical Trials.gov

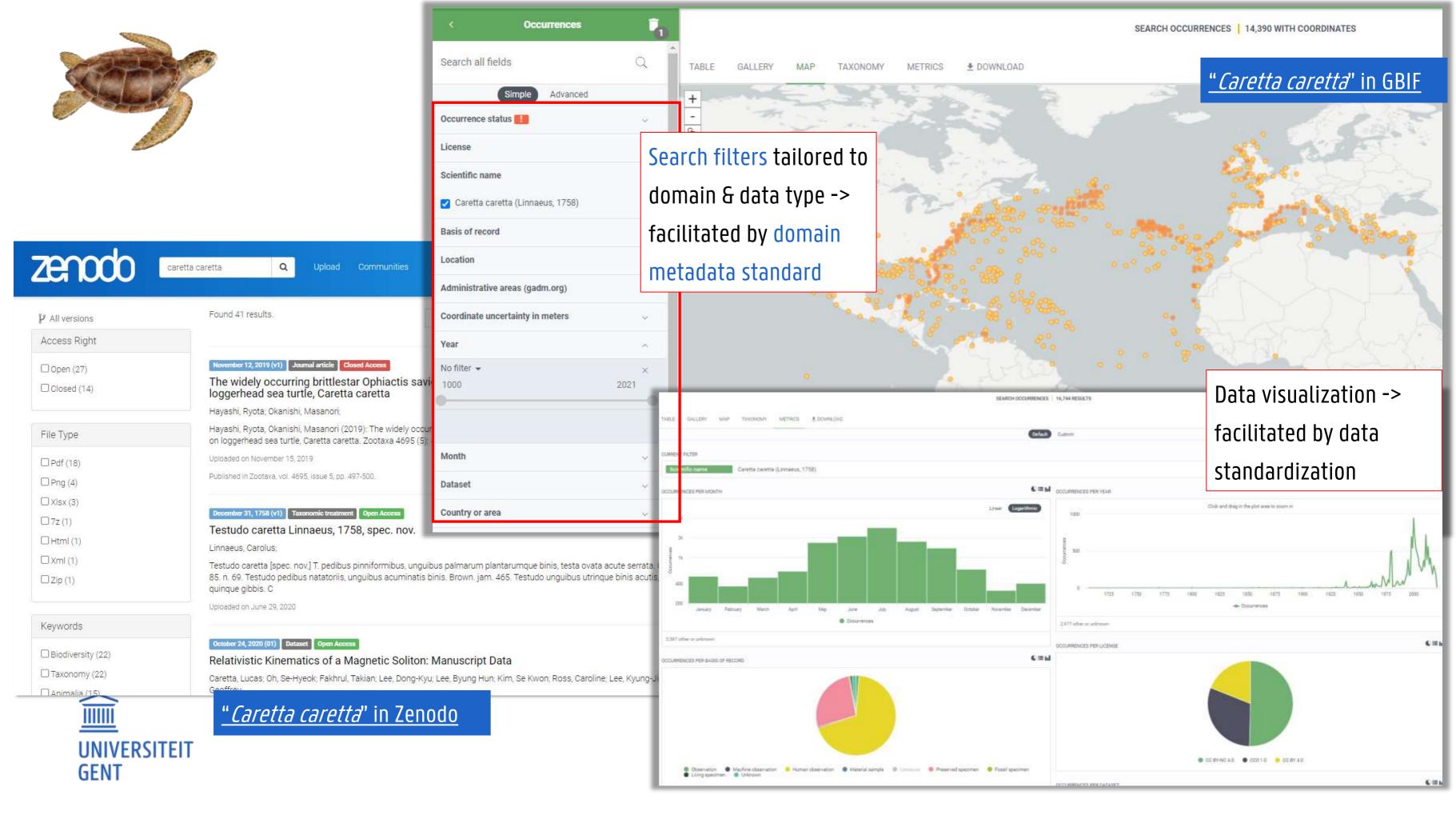
#### Study Datasets 3 datasets.

Click on a Dataset ID in the table below to learn more, and to find out who to contact about access to these data







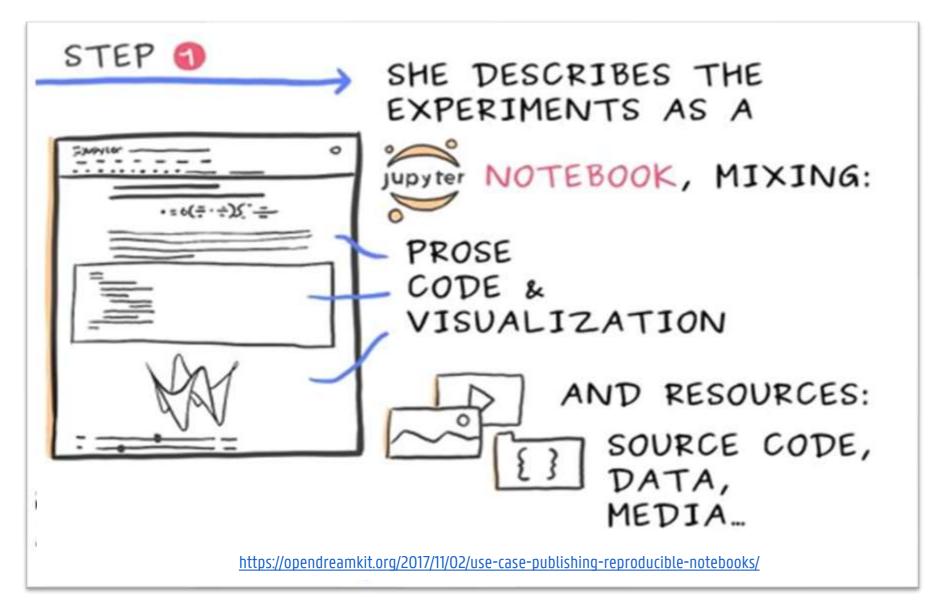


### REPOSITORIES FOR SOFTWARE AND DATA ANALYSIS

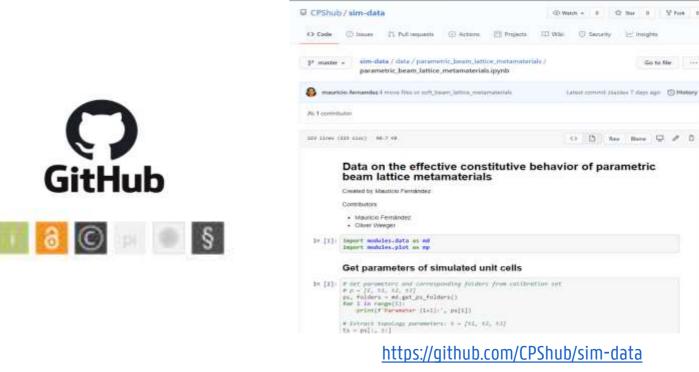
#### Run and publish notebooks

Search or jump to...

#### Jupyter notebooks



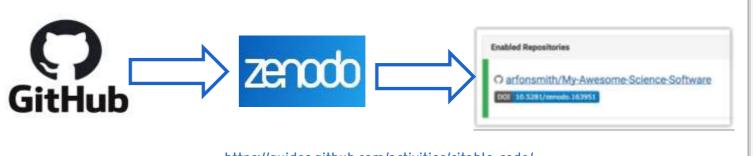








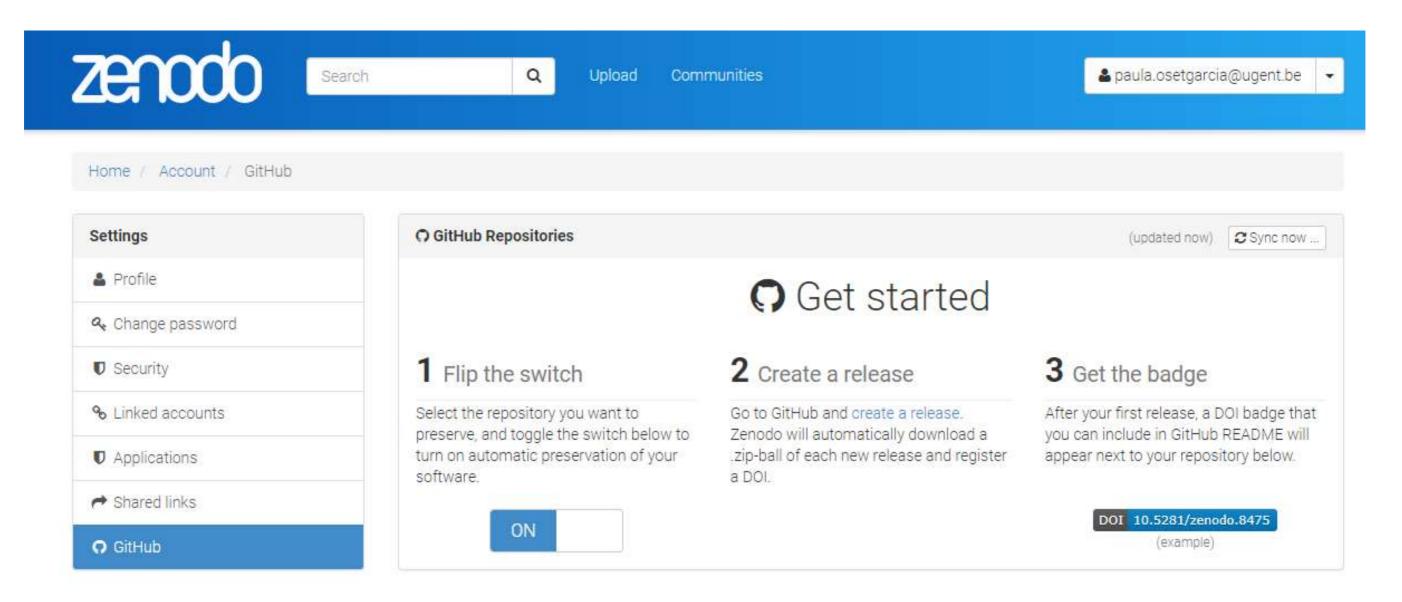
#### Publish git repositories



https://guides.github.com/activities/citable-code/

## SHARING CODE AND MAKING CODE CITABLE

— GitHub – Zenodo integration: <a href="https://guides.github.com/activities/citable-code/">https://guides.github.com/activities/citable-code/</a>





### THE JOURNEY TO FAIR DATA



Plan for data management, keep your DMP updated



Check regulations.

Explore potential data repositories and their requirements & guidelines.











#### Document as you go







- Protocols
- Readme files
- Data overview with metadata
- File organization and naming



Deposit data & documentation in repository; describe it with rich metadata, inc. access & reuse conditions





Adopt domain standards during data collection & processing Implement regulatory requirements



# PREPARING DATA DATA DEPOSITION CHECKLIST (GENERIC)

- Gather all relevant (meta)data needed for reproducibility:
  - experimental method details, raw data files, organized data tables, scripts
     and code, data visualizations, and statistical output
- Migrate data to recommended, reusable formats
- Organize files in a logical schema
- Provide a separate readme file



Jørgen Stamp -Digitalbevaring.dk



More info at: <a href="https://datadryad.org/stash/best\_practices">https://datadryad.org/stash/best\_practices</a>

## DATA LINKED TO ARTICLE PUBLICATION

#### https://doi.org/10.1098/rspb.2022.2456

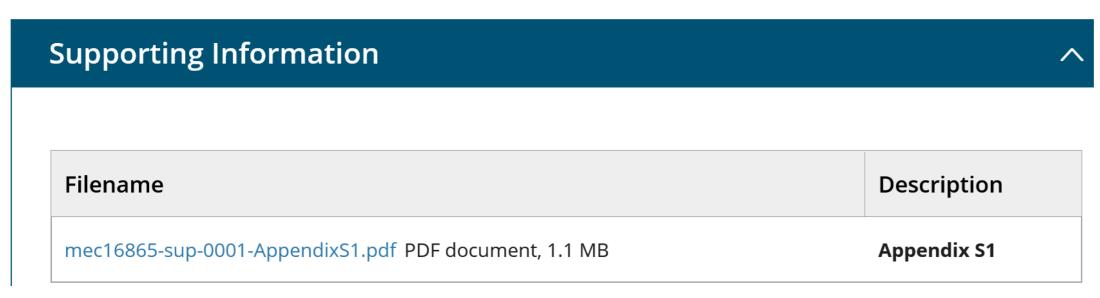
- Neutral processes underlying the macro eco-evolutionary dynamics of mixed-ploidy systems
- <u>Felipe Kauai, Frederik Mortier, Silvija Milosavljevic, Yves Van de Peer and Dries Bonte</u>
- All relevant data are within the manuscript and its electronic supplementary material. The code developed for running numerical simulations is available on a GitHub repository at <a href="https://github.com/KauaiFe/Polyploidy-V1.0">https://github.com/KauaiFe/Polyploidy-V1.0</a>. Readers will also find the code documentation within the electronic supplementary material
- Electronic supplementary material is available online at <a href="https://doi.org/10.6084/m9.figshare.c.6456230">https://doi.org/10.6084/m9.figshare.c.6456230</a>



## DATA LINKED TO ARTICLE PUBLICATION

#### https://doi.org/10.1111/mec.16865

- Runs of homozygosity reveal past bottlenecks and contemporary inbreeding across diverging populations of an islandcolonizing bird
- Claudia A. Martin, Eleanor C. Sheppard, Juan Carlos Illera, Alexander Suh, Krystyna Nadachowska-Brzyska, Lewis G.
   Spurgin, David S. Richardson
- The genomic data supporting this study and code used to perform the data analyses within this article are openly available in the Dryad Digital Repository: <a href="https://doi.org/10.5061/dryad.ksn02v75k">https://doi.org/10.5061/dryad.ksn02v75k</a> (Martin et al., 2022).





## DATA LINKED TO ARTICLE PUBLICATION

#### https://www.ncbi.nlm.nih.gov/pmc/articles/PMC9627755/

- At age 9, the methylome of assisted reproductive technology children that underwent embryo culture in different media is not significantly different on a genome-wide scale
- Rebekka M Koeck, Florence Busato, Jorg Tost, Heleen Zandstra, Sylvie Remy, Sabine Langie, Marij Gielen, Ron van Golde, John
   C M Dumoulin, Han Brunner, Masoud Zamani Esteki, and Aafke P A van Montfoort
- The raw and processed array data from this study have been uploaded to the Gene Expression Omnibus (GEO) database and are available under the accession number GSE196432 (<a href="https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE196432">https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE196432</a> ). The raw array data from the FLEHS study are available within the GEO database under the accession number GSE110128.

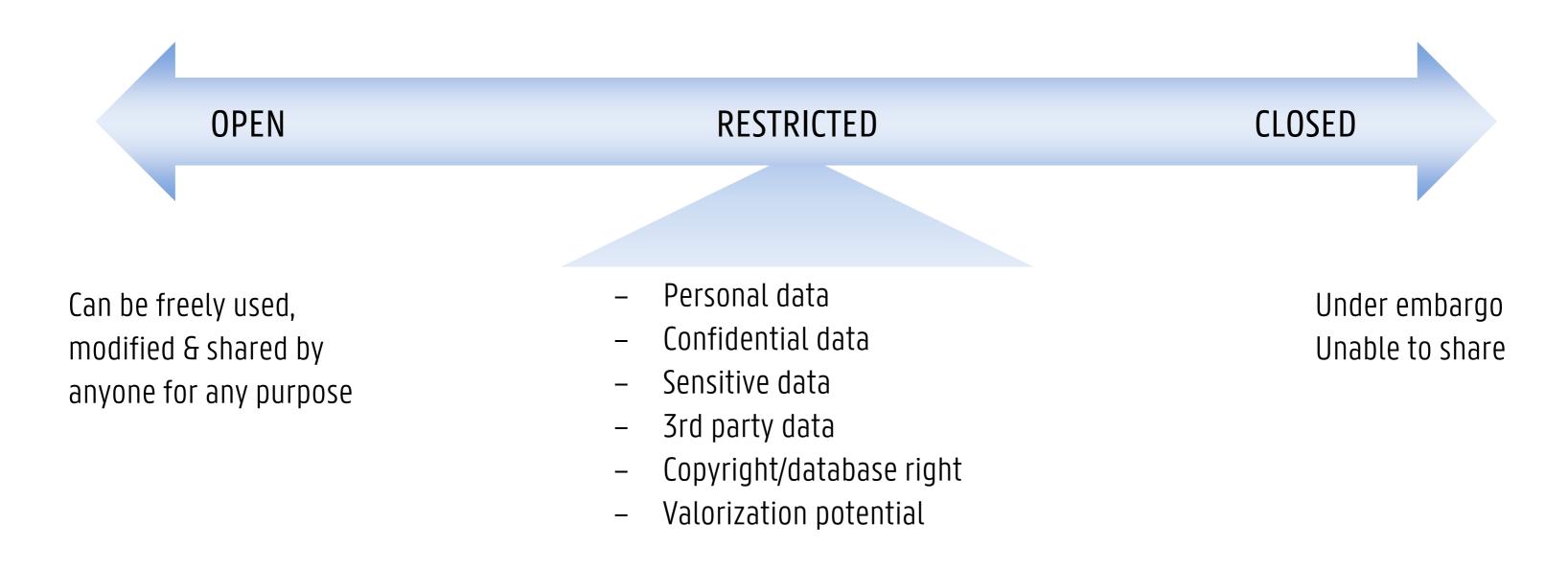


## ALWAYS OPEN SHARING?



## WHAT ABOUT SHARING OF SENSITIVE OR CONFIDENTIAL DATA?

"As open as possible, as closed as necessary"





## RESTRICTIONS ON DATA SHARING













Personal data

Confidential data

Sensitive data

Third-party data

Copyright

Valorization



## TAKE-HOME MESSAGES

- Publishing data can provide personal (career) advantages, and it benefits science and society at large
- Sharing data is a growing requirement from research institutes, funders and journals
- The recommended way to share data is using data repositories
  - → Data repositories provide essential functionalities to make your data FAIR
- Check repository requirements to prepare for deposit during research
- Putting your data in a repository  $\neq$  making data open



#### Ghent University Data Stewards

RESEARCH DEPARTMENT - UNIVERSITY LIBRARY

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