

FAIR Software tools

Carlos Martinez-Ortiz

"How FAIR are you?" webinar, February 24, 2021



Outline

- FAIR principles (in general)
 - Why is this important for researchers?
- FAIR principles (for software)
 - Why is it different from data?
- Guidance
- Questions?











2016: "The FAIR guiding principles for scientific data management and stewardship" Wilkinson et al., doi:10.1038/sdata.2016.18

2018: "Central to the realization of FAIR are FAIR Digital Objects, which may represent data, software or other research resources." (European Commission)

2019: "Towards FAIR Principles for Research Software" Lamprecht et al., doi: 10.3233/DS-190026





My personal view on FAIR











Why is reuse important for science?

Antonie van Leeuwenhoek

- 70x to over 250x magnification
- Crafted his own lenses
- Pioneered the study of microscopy



Image credit: Rijksmuseum Boerhaave





Why is reuse important for science?

- Lab equipment is readily available
- Microscope specifications allow us to find an equivalent device

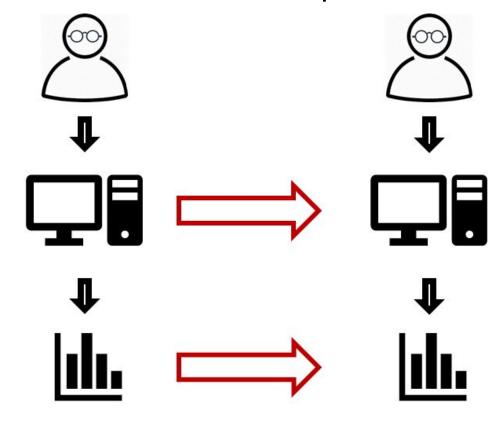
The devices we use for science are important!



Image credit: Wikimedia commons

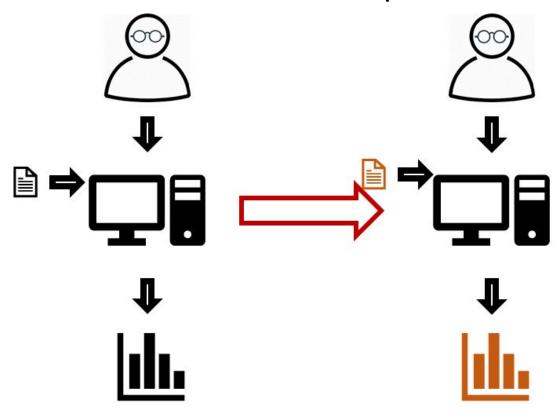


Software is our modern "microscope"



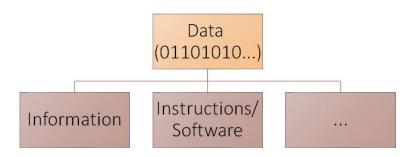


Software is our modern "microscope"



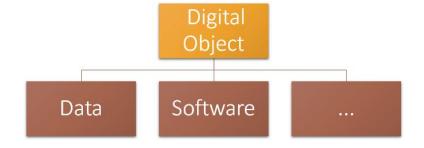


Can we use the FAIR principles for software?



Software is not data!

VS.







Some previous FAIR software initiatives/outcomes

"Applying FAIR Principles to Software" at the 2017 Workshop on Sustainable Software Sustainability (WOSSS17)

"Sharing Your Software – What is FAIR?" at the 2018 American Geophysical Union (AGU) Fall Meeting

"FAIR Software" Birds of a Feather meeting at deRSE 2019

Towards FAIR principles for research software 2019 DOI: 10.3233/DS-190026

From FAIR research data toward FAIR and open research software

"Making Software FAIR" at the DTL Communities@Work 2018 Conference

Top 10 FAIR Data & Software Global Sprint, including "10 easy things to make your software FAIR" 2019

"Five recommendations for FAIR software" at NL-RSE 2019

FAIR Computational Workflows 2020 DOI: 10.1162/dint_a_00033

Lorentz Workshop 9-13 March 2020 (<u>Automated Workflow Composition in</u> <u>the Life Sciences</u>) "FAIRness assessment for software" at the 2018 DBCLS/NBDC BioHackathon

"FAIR principles for Software" at 2019 Workshop on Sustainable Software Sustainability (WOSSS19)

TIB Training workshops on FAIR Data and Software 2018 - 2019

FAIRsFAIR T2.4: FAIR assessment for research software

BRDI NAS Washington 16-17 March 2020





Ongoing effort on defining FAIR principles for software

FAIR4RS WG

- https://www.rd-alliance.org/fair-principles-research-software
- https://www.rd-alliance.org/group/software-source-code-ig/wiki/fair4software-reading-materials
- https://github.com/force11/FAIR4RS/blob/master/CommunityEngagementChannels.md







Let's talk about you...

Which countries do you or your organisation represent?

Cyprus	France	UK
Spain	Finland	South Africa
UK	Ireland	Ghana



Which countries do you or your organisation represent?

brexited UK	Germany	South Africa
Cyprus		



Mentimeter

What is your role? (PhD, researcher, ...)

researcher

PhD student

Researcher

Researcher

Postdoc

Application Scientist

PhD Student

staff bioinformatician



What is your role? (PhD, researcher, ...)

Mentimete

Community Manager	Researcher	coder
Researcher	coordinator	Bioinformaticist



Bioinformatics & Genomics

What is your field of research?

I'm a PhD student in bioinformatics

Bioinformatics

Genomics & bioinformatics

SE

biocuration

The research software community

bioinformatics / comp sci

Clinical Research Informatics



What is your field of research?

Data Management Software

Bioinformatics, Applied ML

bioinformatics

PopGen/Reference

Graphs/Genomics

genomics

genomics



How do you use research software?















What type of research software do you use?

Excel

depends on which task and which project

R

Depends on the project

R, docker, Make, standalone tools

Packages in R for analysing sequencing data, Galaxy Workflows

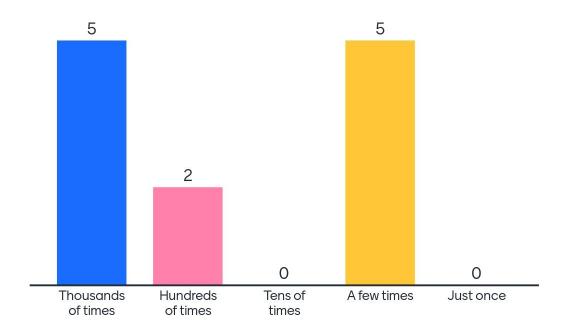


What type of research software do you use?

Command-line, Python, R, Nextflow



How many times does research software gets reused





How do I get started?



ELIXIR

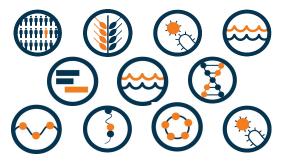


Summary of ELIXIR's key opportunities for participation



Platforms

(cross-domain, focusing on one aspect of service delivery)



Communities

(domain or technology specific)



Focus groups

(Topic specific, most open to all)



The Tools Platform: Developing a tools ecosystem





To support efforts around software packaging & containers, e.g. BioConda/BioContainer and support sustainable integration into bio.tools and OpenEBench



Bio.tools, a
discovery portal
for
bioinformatics
software
information,
providing
curated
description of
tools and data
services



OpenEBench, an infrastructure providing services for hosting scientific benchmark activities and technical monitoring of bioinformatics tools and service

= Galaxy

To drive the development of execution platforms (eg Galaxy) and ensure integration with bio.tools, OpenEBench and workflows using CWL



Raise software quality and sustainability, by producing and promoting software best practices and developing training activities

Tools Interoperability, guidelines and resources for guaranteeing platforms integration at the ELIXIR Tools platform ecosystem, with other platforms at ELIXIR and beyond.

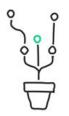




Visit fair-software.eu



#1
USE A PUBLICLY
ACCESSIBLE
REPOSITORY
WITH VERSION
CONTROL



#2 ADD A LICENSE



#3
REGISTER YOUR
CODE IN A
COMMUNITY
REGISTRY

#4
ENABLE CITATION
OF THE
SOFTWARE



#5 USE SOFTWARE QUALITY CHECKLIST





Where can I learn more?

https://quide.esciencecenter.nl/

https://the-turing-way.netlify.app/

https://softdev4research.github.io/4OSS-lesson/

https://elixir-europe.org/platforms/tools/software-best-practices











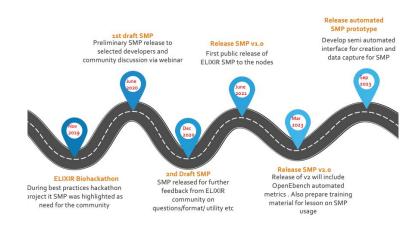
What else can I do to make my software more FAIR?

- Software management plans
 - O What are they?
 - How would that help me?
 - Where do I start?

Why use Software Management Plans (SMPs) in ELIXIR?

Towards unified approach to software development and management in ELIXIR.

- Similar to Data Management Plans, SMP is an awareness tool.
- Helps group think in advance about the software that will be developed
- SMP questions help you think about most important parts of project
- Help assign roles and responsibilities in software project
- Can be used as a guide for everyone involved in the project
- Community driven development planned within ELIXIR







More software management plans

- SSI software management plans:
 - https://www.software.ac.uk/software-management-plans
- eScience Center Software Sustainability Protocol
 - http://doi.org/10.5281/zenodo.1451751

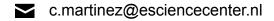




Tools for FAIR sharing of data

- https://github.com/FAIRDataTeam/FAIRifier
- https://github.com/fair-data/fairdatapoint
- https://github.com/fair-workflows/nanopub





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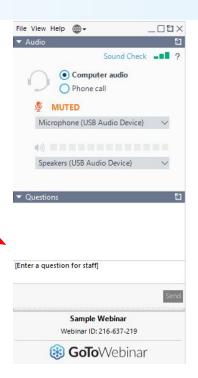




Questions?

Please write your questions in the questions window of the GoToWebinar application

Title: FAIR software tools
Presenter: Carlos Martinez Ortiz







Next CINECA webinars

Title: Practically Fair

Presenter: Andrew Stubbs

Date: Thurs 4th March 2021

Time: 3:00 PM GMT / 4:00 PM CET

Registration and details:
https://www.cineca-project.eu/news
-events-all/practically-fair

Title: How to make training FAIR

Presenter: Anna Swan & Sarah Morgan

Date: Thurs 18th March 2021

Time: 3:00 PM GMT / 4:00 PM CET

Registration and details:

https://www.cineca-project.eu/news-eve nts-all/how-to-make-training-fair