

**RESEARCH ARTICLE** 



# Removing Barriers to Reproducible Research in Archaeology

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Cite as

Karoune, E., and Plomp, E. (2022) Removing Barriers to Reproducible Research in Archaeology. Zenodo, ver. 5 peer-reviewed and recommended by Peer Community in Archaeology. https://doi.org/10.5281/zenod o.7320029.

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Recommender Ben Marwick

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Cheng Liu, Lisa Lodwick, Sam Leggett, Cyler Conrad. This version of the article has been peer-reviewed and recommended by

\*Peer Community in Archaeology\*\*

\*\*This version of the article has been peer-reviewed and recommended by Peer Community in Archaeology\*\*

https://doi.org/10.24072/pci.archaeo.100022

#### ABSTRAC<sup>®</sup>

Reproducible research is being implemented at different speeds in different disciplines, and Archaeology is at the start of this journey. Reproducibility is the practice of reanalysing data by taking the same steps and producing the same or similar results. Enabling reproducibility is an important step to ensure research quality and validate interpretations. There are currently many barriers to moving towards reproducible research such as the skill level of researchers in the practices, software and infrastructure needed to do reproducible research and concerns relating to opening up research such as how to share sensitive data.

In this article, we seek to introduce reproducible research in an understandable manner so that archaeologists can learn where and how to start improving the reproducibility of their research. We describe what reproducible archaeological research can look like and propose three different computational skill levels of reproducible workflows with examples. Finally, in an extensive appendix, we address common questions about reproducible research to remove the stigma about these issues and suggest ways to overcome them.

Keywords: Reproducibility, Archaeology, Open Research, Computational Skills

#### Introduction

The move towards reproducible research has been accelerating in recent years in all research disciplines, especially in terms of increased data and code sharing (Peng & Hicks 2021). Developments such as the UNESCO recommendation on open science are driving forward open science practices including reproducibility (UNESCO 2021). The adoption of open science practices has been happening even faster since the COVID-19 pandemic with major publishers making COVID-19 articles freely open access, a surge in preprints (Besançon et al. 2021, Kadakia et al. 2021) and initiatives to make data openly

available (OpenSAFELY 2022). The pandemic forced researchers to work out how to conduct research in distributed teams and move research activities online such as the surge on virtual conferences (Viglione 2020) and increase in global collaborations (Maher & Van Noorden 2021). These online research activities have adopted the collaborative and computational methods common in open science communities, pushing this approach further into the mainstream of research.

Reproducible research practices benefit us as archaeologists as they increase the sustainability and reuse potential of archaeological data for future research. Increased access to research outputs can make participation in archaeological research more equitable for researchers and the wider public. Furthermore, archaeology is not just one discipline; it includes many sub-disciplines with different methodologies, practices and data standards. This inherent interdisciplinarity can make it difficult to understand each other's research, let alone build on research by using someone else's data, especially if the research is not conducted in a transparent and reproducible manner.

Nevertheless, there is still a long way to go for all archaeological research to be reproducible and there are many barriers that archaeological researchers face when trying to implement reproducible research (Carney & Davies 2020, Marwick 2017a, Marwick *et al.* 2017, Strupler 2021, Strupler & Wilkinson 2017). Barriers include limited time due to pressure to publish, lack of resources and integrated infrastructure, increasingly complex datasets and analysis scripts, and not knowing how to conduct reproducible resource (Alston & Rick 2020; Barker 2016; Peng 2011; Peng & Hicks 2021). In this article, we are seeking to address the barrier of not knowing how to conduct reproducible research, by explaining what is typically meant by *reproducible*, describing why reproducible research is important for archaeological research, giving some examples of what *reproducible workflows* look like. We define three levels of creating reproducible workflows to provide entry points for any archaeologist to improve their work, even those with only basic computer skills. We also include an appendix that answers common concerns and questions about reproducible research (Appendix A). A second appendix includes a glossary that defines words appearing in the main text in bold and italic typeface (Appendix B).

We propose that researchers take small-steps to implementing a reproducible workflow. We recommend starting by applying open science practices to one aspect of your research and then keep adding another skill or practice. Conducting reproducible research involves learning knowledge and skills about many different open science practices and this can take time. By taking small steps in learning new skills, reproducible research and open science practices seem less daunting and archaeologists can gradually move towards fully reproducible workflows (Figure 1).



**Figure 1:** Taking incremental steps to improve your reproducible workflow will help you to increase your skills in transparently sharing your research. The Turing Way project illustration by Scriberia. Used under a CC-BY 4.0 licence. DOI: <a href="https://doi.org/10.5281/zenodo.3332807">10.5281/zenodo.3332807</a>.

#### What is reproducible research?

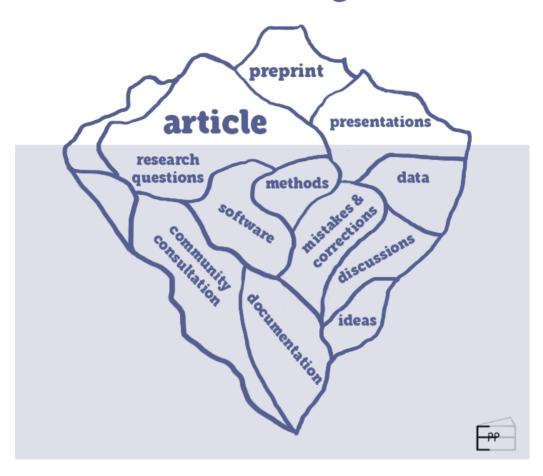
Reproducible research is when data from the original study can be reanalysed taking the same steps and producing the same or similar results (**Figure 2**). This can only be achieved with a transparent record of the research, also known as a reproducible workflow. Therefore, the data, methods, and analysis should be made available to allow other researchers to review and reproduce the study. This means the research results can be validated and reused more easily. Research that is not reproducible, or not shared in a transparent manner, is more difficult to assess and is not representative of the full extent of the research that has been conducted. It is much like **Figure 3**: when we just look at publications and presentations, we are only able to see the tip of the 'Research Iceberg', and we may not understand the entire nature of the research as it was originally conducted. Closely associated with reproducible research is the term 'computational reproducibility', which is used more specifically for obtaining the same results despite different computing hardware or software compiler configurations (see for example Marwick *et al.* 2018 and Strupler & Wilkinson 2017).

		Data		
		Same	Different	
Analysis	Same	Reproducible	Replicable	
	Different	Robust	Generalisable	

**Figure 2:** Reproducibility and replicability terminology explained. Image by The Turing Way Community (2021) under CC-BY 4.0. Same and different refer to the data and analysis being used - in all cases you are looking for same or similar results.

Reproducibility is distinct from *replication* (Figure 2). Replication is where a study is conducted independently using the same analysis as the original study to produce different data and resulting in the same or similar answers (Graham & Huffer 2020). In archaeology, direct replication of results (from the same samples) is relatively uncommon due to the limited availability of remains to investigate and status traditionally attached to collection of new data through fieldwork. As Strupler (2021) suggests, replication of archaeological investigations does occasionally take place by returning to earlier excavated sites and carrying out further investigations, re-analysing museum collections, or revisiting earlier publications. Replication may also occur as part of teaching archaeology, for example when students are assigned to reanalyse published research (Marwick *et al.* 2020b). A result is *robust* when a dataset is analysed using different analysis approaches that provide similar answers. *Replicable* and robust findings then allow us to establish *generalisable* results, where the result is not dependent on a particular dataset or specific workflow (The Turing Way Community 2021).

# Research Iceberg



**Figure 3:** The Research Iceberg, where only the article, *preprint*, and presentations on research are visible. The components of the research on which these visible outputs are based remain invisible (research questions, methods, data, mistakes and corrections, discussions, community consultation, documentation and ideas). Image by Esther Plomp.

#### Why is reproducible archaeology important?

Archaeology is the study of the materials of past human life and activities (Daniel 2021). Archaeological research is extremely varied involving many different sub-disciplines and crossing the humanities and the sciences. It produces many types of data, both quantitative and qualitative, and as a discipline we are just becoming aware of what this means in terms of open data sharing and other transparent practices that enable reproducible research (Kansa *et al.* 2013; Marwick *et al.* 2017).

There are three key reasons for moving towards reproducible research in archaeology. First, the limited remains available for study (limited by the destructive nature of archaeological research, financial, location and ownership limitations). Second, equal access to knowledge generated by these remains; and third, the sensitive remains that we study.

Most archaeological research involves the destruction of materials (Harris 2006) - whether this is during excavations or scientific investigations. The data and *metadata* (see also *paradata*) collected during excavations is often all that is left of the *in-situ* archaeological remains. We use the stratigraphic method to record information about archaeological sites. The artefacts and ecofacts removed from archaeological sites are changed during the process of our studies through sampling, cleaning, conservation and analysis. Hence, we need to implement ways of working to preserve the data and metadata of these processes in the most sustainable manner possible to allow future generations to reuse this information for reinterpretation of

archaeological remains. Kansa & Kansa (2021) suggest that broadening data literacy skills in archaeology will result in realising the full potential of archaeological data such as data reuse across projects and large-scale data integrations. We must therefore concentrate on facilitating reuse of physical and digital artefacts, data and metadata, with as much care as we do with recording sites stratigraphically to preserve the archaeological record.

Compounding this destructive methodology is the finite remains that we study. Archaeological excavations are limited by the amount of funding for archaeological research and the finite number of locations that can be excavated. All excavations are limited in the time allowed for excavations and often the areas on archaeological sites that can be excavated. Therefore, we typically do not excavate the whole surface of archaeological sites. Nevertheless, the process of excavation requires destruction of the specific locations that we do excavate.

The artefacts and ecofacts that we sample are altered or destroyed through analysis and often only studied in a limited way - limited by restraints on money for scientific analysis in terms of specialists' time, paying for time using scientific equipment, technician time, consumable costs, as well as limited by restricting the number of people who can study the material. Often only one or very few specialists examine each type of material from one site. Consequently, it is of paramount importance that our research is reproducible to enable (re)assessments of archaeological data by other specialists who may not be able to directly physically access the original finds.

Reproducible research practices can help to ensure equitable access to archaeological research. *Transparent recording* of the whole research lifecycle makes research more accessible to anyone, allowing them to participate in the research process. Transparent recording also allows credit to be given fairly for the work that is done in the whole research project. To move to a more sustainable and inclusive future for archaeological research, we need to move away from the idea of sole ownership of research kept on our local computers that only benefits ourselves or few researchers. We must move to a more altruistic way of working for collective benefit by opening up our data (when possible) and processes for increased validation and reuse.

A third reason to move to reproducible ways of working is that some types of archaeological data and research focuses are sensitive. For example, studies involving human skeletal remains and excavations conducted on sites belonging to Indigenous groups. We need to consider carefully who owns the remains and the data we produce from these studies (Carroll *et al.* 2020). We need to consider questions such as who should have access to these resources for research, and how they are best preserved in the long term. The CARE principles (Carroll *et al.* 2020) are a framework to help researchers address data stewardship and governance around Indigenous data (see **Appendix A**). It is also imperative to work out how the physical artefacts and the digital outputs can be stored to make them accessible to the appropriate audiences and for sustainable future use. Sensitive data does not preclude reproducibility. In fact, it is more important to establish validation processes as there may be limitations with sharing data.

#### What does reproducible research look like?

Most published research articles are currently not reproducible - they are not transparent records of research. This is demonstrated by a survey conducted by Nature of 1,576 researchers from multiple disciplines, which found 70% of the respondents had tried and failed to reproduce another scientist's experiments (Baker 2016).

It is difficult or impossible to reproduce research from research articles as most are stand-alone papers that contain brief methods and minimal data. The potential for robust validation and reuse of these pieces of research is therefore rather limited.

The differences between stand-alone articles and articles that contain the details for full reproducibility can be found in Peng's (2011: Figure 1) reproducibility spectrum (Figure 4). This diagram shows the addition of data and code files as well as details of the *computational environment* linked to the paper to move towards full reproducibility of the research. In fact, more detail is needed than stated in Peng's spectrum because the full methodological details (*protocols*) used for data collection would be required for replication of any experimental work included in the article (Figure 4). These methodological details could also be called *metadata* or *paradata*. Large meta-analysis studies need computational reproducibility to enable merging and reuse of datasets and pose unique challenges (Ellis *et al.* 2021; Morrison *et al.* 2021). Computational

reproducibility is vital for studies that want to reuse the same methods for additional analysis of samples from the same or similar archaeological sites.

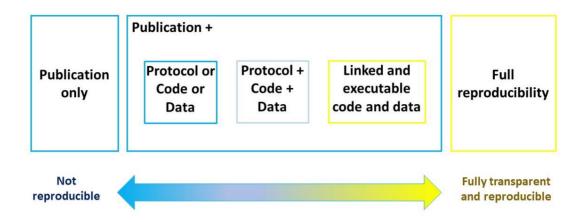


Figure 4: An adapted reproducible spectrum (Peng 2011) with the addition of protocols.

In archaeology, authors often only state they will share their data upon request. However, research in other fields has found that authors rarely respond to these requests (Tedersoo *et al.* 2021, Vines *et al.* 2014). Consequently it is now recommended practice, for greatest sustainability and findability, that researchers deposit research files (data, code and methods) in a *data repository*. Institutions might provide access to a data repository, there may be a discipline specific data repository (such as IsoArcH for isotopic data - <a href="https://isoarch.eu">https://isoarch.eu</a>), or researchers can make use of general data repositories, such as <a href="Zenodo-https://zenodo.org">Zenodo - https://zenodo.org</a> (European Organization For Nuclear Research and OpenAIRE, 2013), <a href="Open Acience Framework">Open Acience Framework - https://osf.io</a>, or <a href="Figshare - https://figshare.com">Figshare - https://figshare.com</a> (see <a href="Appendix A">Appendix A</a> for more information on <a href="how to make your data accessible">how to make your data accessible</a>). Using data repositories will provide a <a href="Digital Object Identifier">Digital Object Identifier</a> (DOI) for all files as a whole, or for each research output, depending on where and how it is archived. It is then important to use the DOI(s) to write a <a href="data">data</a> and code <a href="availability statement">availability statement</a> (The Turing Way Community 2022a) at the end of a article - this links the article with the rest of the research outputs. Thanks to the DOI assigned to the research outputs, the transparency of the research record is improved and benefits such as increased visibility and citations are obtained (Piwowar <a href="eac. 2007">eac. 2007</a>; Piwowar & Vision 2013; Christensen <a href="eac. 2019">eac. 2019</a>; Colavizza <a href="eac. 2019">eac. 2019</a>; Colav

Computational tools such as *version control* and open-source software for data analysis can facilitate transparency of research. Version control is a systematic approach to record changes made in a file, or set of files, over time. It creates a history of the changes made to the file(s) that can be transparently reported. Version control can be achieved simply by using naming conventions, such as file-v0.1 and file-v0.2, to name your files. Software such as Google Drive can also be used to automatically track the history of files. There are more advanced version control systems such as <u>GitHub</u> (<a href="https://github.com">https://github.com</a>) or <u>GitLab</u> (<a href="https://github.com">https://github.com</a>) or <u>GitLab</u> (<a href="https://github.com">https://github.com</a>), which use the **Git** software. **GitHub** accounts can be linked to a data repository for archiving, such as <u>Zenodo</u> and the <u>Open Science Framework</u>. These computational tools create a richly detailed history of any research files that can even be used to assign credit for each individual researcher's work during the project. Please see <u>The Turing Way for an example of how contributors can be recognised</u> (The Turing Way Community 2022b).

Open-source software is software that is released under a license that gives users the rights to freely use, study, change, and distribute the software and its source code to anyone and for any purpose. Commonly used open source software languages are R and Python. When analysis scripts are written in R and Python, all of the steps taken in the data analysis are transparent and traceable and can easily be shared with others. Other researchers can reuse the code for their own needs and it could potentially enable others to reproduce your analysis, if accompanied with other research outputs (for example, method, data and computational environment). Marwick (2022a) provides a list of archaeological articles that both use R for data analysis and

visualisation, and also have R code openly available to facilitate reproducibility. *Proprietary software* (software that requires a paid license to be able to use it, for example SPSS or Excel) may in some cases be more user friendly, but using these tools typically prohibits the examination and reproduction of methods. This is because unless the publications is accompanied with detailed written documentation of analysis steps, the reader is unable to examine the analysis code. Furthermore, others may not have access to the paid software that you have used (Nust & Pebesma 2020).

Although advanced version control systems and open-source software can help to create a transparent reproducible workflow, they often have a steep learning curve creating a barrier to some researchers. However, a transparent reproducible workflow doesn't have to include theuse of advanced computational skills to achieve reproducibility and there are many levels of reproducible workflows. We describe three different levels of creating reproducible workflows here, listed in order of computational skill required (**Table 1**). Following one of the skill levels of reproducible workflows proposed here will help to produce a more transparent record of research that can be linked to in the text of research articles.

Table 1: Three levels of reproducible workflows based on computational skills (least skilled to most skilled).

Level	Needed	Computational skill required	Examples	Tools
1. Transparent recording	- Documentation of data collection and analysis steps - Raw data - Analysis output file	Yes, basic (non- coding)	Karoune (2021a, 2022); Strupler & Wilkinson (2017)	Excel, Google docs and sheets, SPSS, Data Repository
2. Research Compendium	- Documentation (README) - Data - Code	Yes, intermediate	Plomp (2021), Leggett (2022a), Crema et al. (2022), DiNapoli et al. (2019), Florin et al. (2021), Huffer & Graham (2017), Kim et al. (2021), Lancelotti (2018), Lewis (2021), McLaughlin et al. (2021), Reidsma et al. (2021), Reynolds et al. (2019), Selden et al. (2021), Spake & Cardoso (2021), Timbrell et al. (2022), Timpson et al. (2021), Utting (2022)	GitHub, GitLab, R, Data Repository
3. Executable Article	- Documentation (README) - Data - Code - Computational environment	Yes, advanced	Wang & Marwick (2020a), Bartholdy & Henry (2022), Paixão <i>et al.</i> (2021), Gantley <i>et al.</i> (2018)	GitHub, GitLab, R, Binder, Data Repository

## Level 1: Transparent recording of all sampling, laboratory methods, data and analysis through documentation

Transparent recording requires the least computational skills but produces a full transparent record of what has been done. It does not include any computational code, as the analysis steps can all be written down in a simple document and linked to an open dataset. This means any type of analysis software such as Excel, Google sheets, SPSS, etc, can be used. To achieve reproducibility, it is important to write down all the analysis steps taken in a document in a way that another person could understand and reproduce what was done.

Files to include with an article can be deposited in a data repository and referred to in the text of the article by their DOIs, the *data availability statement* and the references.

These files might include a:

- Document file clearly written data collection methods (sampling and laboratory methods) and analysis steps;
- Raw data files csv format is the best for reuse for tabular data. For image files JPEG, TIFF, PNG, SVG
  are generally recommended. The DANS file format recommendations contain guidance on other
  recommended formats (DANS 2022).
- Analysis output file SPSS output file or analysis version of Excel file. This is a cleaned version of the
  data files with analysis performed on it. It should show outputs such as figures produced for an article
  and include calculations.

You can version control all your work using a file naming system or choose a software that contains a simple history tracking system such as using Google Docs and sheets. This will help you to document your data collection and analysis steps fully.

#### Level 2: Research compendium linked to your article

A research compendium contains extensive documentation about the methods used, code files, details of the computational environment and raw data files (Marwick *et al.* 2018). A set of folders can be set up from the beginning of your research project and continually added to throughout the project. There is no single correct folder structure, but there are several widely used templates for research compendium folder structures, for example, <u>Project Tier</u> (2022) and <u>The Turing Way Community</u> (2022c).

Files and folders to include in a research compendium, deposited in a data repository and linked to using a DOI in an article (in the text and in the data/code availability statement):

- README file contains clearly written data collection methods (sampling and laboratory methods), information about the computational environment
- Data raw data, cleaned data, analysis data,
- Code scripts used to analyse data.
- An output folder could also be included for the final article tables and figures.

**Table 1** lists several examples of research compendia, here we describe two compendia in detail to illustrate their typical contents: Plomp (2021a) and Leggett (2022a).

Plomp (2021a) provided a detailed description of the dataset in the article (Plomp 2021b), with links to the dataset on a disciplinary specific data repository, IsoArcH (Salesse *et al.* 2018), and R scripts used in data analysis are publicly available on GitHub/Zenodo. The dataset on IsoArcH is available in .xlsx format and includes more detailed geographical information of the samples (latitude, longitude, altitude and distance from sea) as well as a .ris file containing the relevant research articles (Plomp 2021b). The figures in the data article were produced using R, and the scripts (with documentation and installation instructions) are shared on GitHub and archived on Zenodo (Plomp & Peterson 2021).

Leggett (2022a) provided a detailed description of the dataset context in the article, with links to the dataset and the R scripts used to analyse the data made available on OSF (Leggett 2022b) and GitHub (Leggett 2022c). Leggett provides examples of the outputs as figures, which are clearly and consistently numbered in

the same way as the publication. These figures are provided in a higher quality .TIFF format, and in a compressed format if files are too large (.JPG). The R script is available and contains comments to facilitate reuse. Leggett indicated the license clearly in the README file of the GitHub repository and in the metadata information at OSF.

#### Level 3: Executable research compendium

In an executable research compendium, the author provides a computational environment so that readers can easily reproduce the figures and other computational results presented in the article. This makes it easy for others, such as peer reviewers and readers, to reproduce the results even if their computers lack the exact packages that was used.

The files and folders to include with an article are the same as with a level 2 research compendium, but they are packaged up to run the code. This will typically include a:

- README file contains data collection methods (sampling and laboratory methods), information about the computational environment
- Data raw data, cleaned data, analysis data
- Code scripts used to analyse the data
- Container using a tool such as Binder

Binder is a web application that allows users to create sharable, interactive, browser-based computational environments from code repositories (Gibson 2021, Graham 2018). For example, if you have your research compendium on GitHub, Binder can allow a reader to run the R code in their web browser. No downloading or installation is required by the reader. The examples listed in **Table 1** have a Binder button on their GitHub repository. A reader can click this button to activate RStudio in their web browser, and run code from the author's repository.

As for the level 2 research compendium, files are deposited in a data repository. Then the DOI link is added within the article's data and code availability statement and in the text of the methods section to help the reader find the additional materials.

#### **Conclusions**

Archaeology is a complex science with many sub-disciplines encompassing many different methods and producing diverse datasets. This multi-disciplinarity coupled with the limitations imposed on archaeological research by its destructive practices, the need for equal access to knowledge generated by these remains and the sensitive nature of archaeological remains makes it incredibly important for our research to be comprehensible. This can only be achieved through improving the reproducibility of our research.

We recognise that for many archaeologists reproducibility is an unfamiliar concept. Our article addresses this by defining core terms relating to reproducible research (**Appendix B**) and describing a variety of options for increasing the reproducibility of our research. **Appendix A** contains many additional definitions and answers to frequently asked questions about making research reproducible. We have attempted to motivate you to start your reproducibility journey and remove some of the barriers that may have previously prevented you from starting.

In our opinion, our three levels of creating reproducible workflows provide entry points for any archaeologist, even those with basic computer skills. Remember that you do not have to start with a fully computational reproducible workflow as done in the executable research compendium. The most important thing is to start making your materials available in a transparent manner, which can be achieved by transparent recording and documentation, our Level 1 workflow. Each time you complete a project and obtain more experience with making your research available in a more transparent way, then for your next project you can take a further step to improve the computational reproducibility of your work.

#### **Acknowledgements**

Thanks to *The Turing Way* community. Many thanks to Yo Yehudi for her work on indexing references on data science and to Ben Marwick for his list of reproducible archaeological articles. Thank you to Bjorn Bartholdy and Celine Kerfant for comments on the drafts of this article. Many thanks to Ben Marwick, our recommender, and our reviewers Lisa Lodwick, Sam Leggett, Cyler Conrad, Cheng Liu, for their kind and constructive reviews which improved the clarity and quality of this work.

Preprint version five (v) of this article has been peer-reviewed and recommended by Peer Community In Archaeology of the PCI (https://doi.org/10.24072/pci.archaeo.100022).

#### Data, scripts, code, and supplementary information availability

There is no new data or code used for this article.

#### Conflict of interest disclosure

The authors declare that they comply with the PCI rule of having no financial conflicts of interest in relation to the content of the article. Emma Karoune is on the managing board of Peer Community in Archaeology. Esther Plomp is the Secretary General and Open Research Ambassador of IsoArcH.

#### **Funding**

Emma Karoune's FAIR Phytoliths Project is funded through EOSC-Life from the European Union's Horizon 2020 research and innovation programme under grant agreement No 824087 and hosted by Historic England and Universitat Pompeu Fabra.

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#### **Appendix**

#### Appendix A: Frequently asked questions about reproducible research in archaeology

You probably have many questions about different aspects of reproducible research. Therefore, we want to discuss the most frequently asked questions that we hear from archaeologists about reproducibility to try to remove barriers and help you make progress along your reproducible research journey. The easiest way to use this Appendix is to go to the question that is currently on your mind:

- How do you decide if you should publish data and/or code openly?
- Where do you start training skills in open science and reproducibility?
- What should you consider when you publish an article with underlying data/code?
- How do you share data to make it more accessible to others?
- How do you clean up the data and code before sharing this publicly?
- How do you share research methods more openly?
- What is metadata?
- What about licenses/copyright?
- Isn't reproducible archaeology more expensive?
- What if people misinterpret data or find a mistake?
- Is archaeology suitable for preregistration?
- A supervisor doesn't want to work reproducibility, how do you convince them?
- Will reproducible research be taken into account when looking for a next job?
- Do platforms like SciHub, ResearchGate, Academia.edu count as Open Access?

#### How do you decide if you should publish data and/or code openly?

There may be several reasons that you cannot share your data or code publicly. The data you work with may belong to a community you are collaborating with, you may be dealing with personal data, sharing the data may have consequences on biodiversity, you might not be sure if you have any data to begin with, you may not have the rights to share the data or software, or you may be concerned about people 'scooping' your results.

#### You collaborate with a community

To ensure that you do not harm the community that the data belongs to, it is important to follow the CARE principles. The CARE principles facilitate Indigenous control in data governance and reuse, promoting equitable participation (Carroll *et al.* 2020). They address historical inequities and ensure that value from Indigenous data is created in a way that is grounded in Indigenous worldviews and by creating opportunities for Indigenous Peoples.

- 'Collective benefit' for Indigenous Peoples must be facilitated when Indigenous data is used, to
  achieve inclusive and equitable innovation, as well as to improve governance and citizen
  engagement.
- 'Authority to control' and govern data is the right of Indigenous People.
- 'Responsibility' is achieved through nurturing respectful relationships with Indigenous peoples when working with their data.
- 'Ethics' in data practices is representation and participation of Indigenous Peoples, who must be
  the ones to assess benefits, harms, and potential future uses based on community values and
  ethics.

The CARE principles require engagement with people and purpose to address the cultural, ethical, legal, and social dimensions associated with the intended uses of the dataset (Carroll *et al.* 2020; 2021, see also Marwick *et al.* 2020a). The CARE principles address issues of relevance for many populations (such as privacy, future use, reuse, stewardship) and can be used as a standard in crafting policies on data acquired

about communities or populations (Carroll *et al.* 2020). When collaborating with communities it is also important to use infrastructure that centers their needs (for example, <u>Mukurtu</u> 2022).

#### You work with personal data

The CARE principles are also aligned with privacy laws, which can place additional requirements on the public sharing of personal data. This may be less relevant for archaeological remains, but can play a role in more recent cases or when your research is based on interviews such as ethnographic studies. These privacy laws differ per country and it is important to check which laws apply. If you are based at a larger institution there are generally experts available that can provide advice.

When following the CARE principles, or privacy laws, it may not always be possible to make the data publicly available, which could hamper reproducibility. The CARE principles and privacy laws should be prioritised in these cases but this does not mean you should not try to work reproducibly. There are alternative methods to fully open data that you could take: restricting data access by providing private repository links, providing access to synthetic data (synthetic data is a fake dataset produced to have the same qualities as your real dataset and therefore would produce similar results using your analysis - see Jordan *et al.* 2022, Noble 2022 or Allen 2021; for introductory articles and a video on synthetic data, and also Shannon & Walker 2018; for a case study in geographic research), or anonymising/generalising datasets by erasing personal/location data. Sharing part of your data or a dataset that is very similar to the original allows others to understand, evaluate and verify the used methods.

#### You work with sensitive location data

It might be harmful to share certain types of locational data and you should weigh the risks versus the benefits of sharing these types of data. Freely releasing GIS coordinates online as part of your dataset could potentially help looters and illegal excavators find sites (Strupler & Wilkinson 2017). This could lead to destruction of archaeological sites. Location data can always be omitted from a dataset if you think this is a potential problem. The Portable Antiquities Scheme in the UK provides a guide (Publishing find spots as a third party - Portable Antiquities Scheme 2012) to publishing find spots to prevent issues with find spot security. Guidance includes not publishing location to a greater resolution than 1km square or 4 figure grid reference and if security is imperative then a pseudonym is given to the location so that the actual location will only be published with the Finds Liaison Officers advice.

The biodiversity community has similar potential problems with sharing the location data of endangered plant and animal species. However, the majority of this community feel there is more benefit using open data as its future reuse could lead to greater conservation opportunities, promote community engagement and reduce duplication of survey efforts (Tulloch *et al.* 2018).

#### You work with qualitative or theoretical data

If your research is more theoretically focused or based on other resources you may not have a dataset to share. Reproducibility may not directly translate to qualitative data given the unique importance of interpretation and subjective nature of qualitative data collection (Tsai *et al.* 2016). Instead, you can focus on providing information about the context of these resources and make your publications and/or books openly available.

#### You do not have the rights to share the data/code

When reusing the materials that others have created, or when you are using proprietary software or hardware, it is important to check if you have the right to share the resulting data and code. Another example is when you are using analysis code, such as R scripts, that has been developed by team members and passed on for use in your lab group. It also may not always be possible to share your results if license restrictions are in place (see **Appendix A**: 'What about licenses/copyright?').

In these cases you should be as transparent as possible about the procedures or processes followed and about the limitations of making your outputs available. In the long term you can consider moving away from proprietary software, if possible, towards open source software such as R or Python so that you can make your code publicly available. You can also encourage others to archive their code and make it citable so that you can properly credit them in your own outputs.

#### What if people will 'scoop' you?

You may wonder what will happen to the data once it is openly available and fear that someone will use the data for their next publication. This is something which has not yet been reported and there are several reasons for this. Generally, when you share your data through a repository, there is a timestamp associated with the work (similar to *preprints* or published articles). With *version control* on platforms such as GitHub it is even clearer who contributed what to the work as there are timestamps and records of all contributions. As you are the expert of the data and/or code, it will also be easier for others to collaborate with you instead of trying to reinvent the wheel themselves, so it's a good idea to make your contact details available to enable collaborators to contact you. Making your data available sets you up for these collaborations because your work is easier to find and having access to the data/code facilitates collaboration.

#### Where do you start training skills in open science and reproducibility?

Upskilling yourself can be time consuming so take it a step at a time and remember it does not have to be costly. There are lots of free and open educational resources for you to use.

Start by looking in these places:

- **1. Your own institution**: Investigate what courses your own institution offers. This could be through your own department, student services, a research software engineering group or library services.
- **2.** Open science online courses to work through at your own speed: Most online courses are not specific to archaeology, but focus on general skills or knowledge for open science that can then be applied to your own research. There are dedicated open science online training platforms that have courses you can work through at your own speed such as <a href="FOSTER">FOSTER</a> (<a href="https://www.fosteropenscience.eu">https://www.fosteropenscience.eu</a>), <a href="Open Science">Open Scholarship Knowledge Base</a> (<a href="https://www.oercommons.org/hubs/OSKB">https://www.oercommons.org/hubs/OSKB</a>).
- **3.** Attend an online course or workshop: This has the benefit of providing you with training materials but also an instructor you can ask for help. For computational skills courses, The Carpentries (https://carpentries.org) runs lots of different courses on data, library and software skills. There are also courses on platforms such as Coursera and EdX, for example, 'Reproducible Templates for Analysis and Dissemination' (https://www.coursera.org/learn/reproducible-templates-analysis). There are also many open science focused workshops some of which are archaeology focused such as recent efforts by the Association for Environmental Archaeology that ran an open science focused conference and a workshop (Karoune *et al.* 2021) and a workshop on Reproducible Archaeology held at Durham University (Clarke *et al.* 2021).
- **4. Apply to join a training programme:** For a more in-depth training experience, you could join an open science training and mentoring programme such as <a href="Open Life Science">Open Life Science</a> (<a href="https://openlifesci.org">https://openlifesci.org</a>) or <a href="Open Life Science">Open Life Science</a> (<a href="https://openlardware.space">https://openlardware.space</a>). These programmes are a mix of seminars, hands on training and mentorship to allow you to gain the skills and support to start or complete an open science focused project related to your own research. There have been a number of archaeological projects within the Open Life Science programme such as "<a href="FAIR Phytoliths">FAIR Phytoliths</a>" (Karoune *et al.* 2022) and "Intellectual Property, Indigenous Knowledges, and the Rise of Open Data in Australian Environmental Archaeology".
- **5. Join a community or association:** There are a number of archaeological associations focused on this way of working such as Computer Applications and Quantitative Methods in Archaeology (https://caa-international.org). There are also online communities around the R programming language, such as Rchaeology (https://rchaeology.github.io) and R-Ladies (https://rladies.org). The Software Sustainability Institute (SSI) is a large community of Research Software Engineers and researchers that use software. They run a fellowship programme for those using computational methods in their research and also offer lots of great resources for those wanting to learn computational skills. Examples of SSI blogs for beginner codes are: Resources for using spreadsheets in research and moving to other tools (Laird 2021) and Training resources for researchers who want to code (Hulme et al. 2021). Your local institution might also have a network that can support you, such as the UK Reproducibility Network (UKRN, https://www.ukrn.org) in the UK and the Open Science Communities in the Netherlands, Sweden, Arabia, Portugal, Serbia and Ireland (https://www.osc-international.com).

#### More free educational resources:

- <u>Teaching Reproducible Collaborative Data Analysis to Undergraduates Using Compendia</u> (Wang 2021)
- Introduction to R Programming for Historical Archaeologists (Bollwerk et al. 2021)
- <u>Tidyverse for Archaeologists A Guide for Beginners</u> (Marwick 2018)
- How To Do Archaeological Science Using R (Marwick 2017b)
- CRAN Task View: Archaeological Science (Marwick 2022a)
- The Open Digital Archaeology Textbook (Graham et al. 2019)
- There are many e-books on R such as <u>Big Book of R</u> (Baruffa et al. 2022) and <u>R for Data Science</u> (Wickham & Grolemund 2017)
- A Beginner's Guide to Conducting Reproducible Research (Alston & Rick 2020)
- Geocomputation with R (Lovelace et al. 2019)
- R style guides like Google's R Style Guide (Google 2022) and ISAAK's R Style Guide (ISAAK 2018)

### You can make your workflow more open by ...

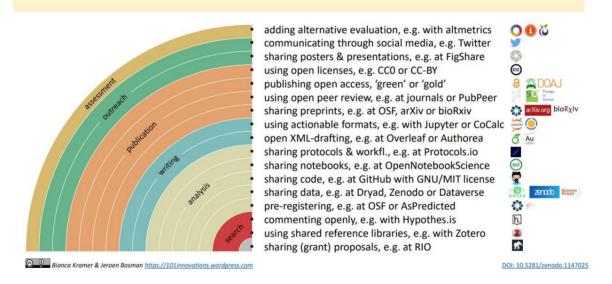


Figure S1: The rainbow of Open Science practices by Kramer & Bosman 2018.

You could take a look at the rainbow of open science practices to get some ideas (Kramer & Bosman 2018, **Figure S1**).

#### What should you consider when you publish an article with underlying data/code?

To help you with publishing your research in a more transparent way, you can follow this checklist:

- 1. Have you set up a *Data Management Plan* (The Turing Way Community 2022d)? In this <u>Data Management Plan</u> you can think about how you will store, document and share your research.
- 2. Ask feedback on how you documented and structured your data and code from someone that is not a co-author: are they able to trace your steps?
- 3. Decide in what order you want to publish your research objects: do you want to publicly share the data and code before the article is published or would you rather wait? see The Turing Way (The Turing Way Community 2022e) for more information on how to <a href="link your research objects">link your research objects</a> during the sharing process.

- 4. Make sure a README file is included with your data/code: even if there is an associated article it is important to provide all relevant information to interpret the data or use the code in the same place as the data/code. You can also let others know how to cite you in this README file (Chue Hong *et al.* 2019a).
- 5. Ensure that you selected a license for your data and code: without it your research objects cannot be reused by others (see 'What about licenses and copyright?').
- 6. Provide the DOI of your research objects where it is relevant (for example, table/figure descriptions, materials and methods section, data/code availability statement).
- 7. Check whether you cited your data and code in the references: this is where citations are counted from not the acknowledgements or data availability statements. If you reused any data/software you should also cite these (Chue Hong *et al.* 2019b).

#### How do you share data to make it more accessible to others?

To make your data accessible and reusable you should share your data according to the FAIR principles. The FAIR principles (Wilkinson *et al.* 2016; Lamprecht *et al.* 2020, **Figure S2**) facilitate the reproducibility of the research undertaken. The principles recommend that scientific data and software are:

- 'Findable' thanks to their *persistent identifier* that is assigned to the dataset via a data repository or through a data article.
- 'Accessible' so that the data and metadata can be examined. Note that for data to be Accessible it
  does not necessarily need to be open: if only the metadata about the dataset is available, the data
  is still considered to follow the FAIR principles.
- 'Interoperable' so that data can be analysed and integrated with other data using common vocabulary and formats.
- 'Reusable' data is appropriately documented and licensed. A license defines what others may or
  may not do with your data. Open licenses, such as those of the <u>Creative Commons</u> (Creative
  Commons 2022) or the <u>Open Data Commons</u> (Open Data Commons 2022), allow others to reuse
  the data without limiting restrictions (see for more detail: '<u>What about licenses/copyright?</u>'
  below).

When choosing where to disseminate your data or code you can choose between two routes: 1) choose one platform or 2) use multiple platforms based on their different functionalities and link the persistent identifiers in the documentation. For example, you could share your data and code on Zenodo (https://zenodo.org, European Organization For Nuclear Research and OpenAIRE, 2013) and your research protocol on protocols.io (https://www.protocols.io). Both Zenodo and protocols.io allow you to add the persistent identifiers to other research outputs in the metadata, making it easy for others to find the related outputs. Note that it is not recommended to share the same outputs multiple times on different platforms, as it will be difficult for reusers to interpret which version they should use and cite. To find the appropriate data repository for your data, you can use FAIRsharing (https://fairsharing.org) or re3data (https://www.re3data.org).



**Figure S2**: The FAIR principles. The Turing Way project illustration by Scriberia. Used under a CC-BY 4.0 licence. DOI: <u>10.5281/zenodo.3332807</u>.

#### How do you clean up the data and code before sharing this publicly?

Before you share the data or code you want to make sure that the dataset is complete and that variables are explained (**Figure S3**). Similarly, for code it will be needed to remove unnecessary parts and make sure functions and variables are adequately documented.

There is often a psychological barrier to sharing data and code due to our own perception that it is too messy or could be criticised by others. See <a href="the-blog by Mombrea">the-blog by Mombrea</a> (2013) about reasons developers are scared of making their code public for more details. However, even researchers with many years of programming experience will constantly seek help to structure data and refine code including using online platforms such as Stack Overflow. Seeking out this help and sharing solutions is more efficient than constantly reinventing the wheel.

For both data and code, it can help to have a colleague or collaborator review your work (see Reimer *et al.* 2019 for an example on how to set this up). They can provide you with feedback on the readability and completeness, and reproduce your results. Any feedback on where your collaborators get stuck or struggle will benefit the outputs that you will eventually share with a wider public.

There are several resources that delve deeper in how you can structure and document your data (Borer et al. 2009; Briney et al. 2020; Hart et al. 2016; Fuchs & Kuusniemi 2018) or code (Sandve et al. 2013; Ram 2019). Some of them go deeper into the specifics of a programming language, such as R (Wickham 2014; Krystalli 2021; Navarro 2021).

It can be helpful to have a folder structure set up and explained in a README file if your dataset/code is very complex. For folder structure examples, see templates set up by Nikola Vukovic (Vukovic 2017), Chelsea Beck (Beck 2016) and Barbara Vreede (Vreede 2020). You can structure folders based on the person that has generated the data/folder, chronologically (month, year, sessions), per project, or based on analysis method/equipment/type of data.

In data management it is important to stay consistent, avoid leaving empty values (use NA instead) as it is not always clear what an empty cell actually means (no value, a value of zero, not measured?). If you use consistent file naming it is easier for you to find your files - see <a href="Jenny Bryan's work">Jenny Bryan's work</a> (Bryan 2015) and <a href="Caltech's guide">Caltech's guide</a> (Briney 2020). For example, you can include the date in the format YYYMMDD in your file name so that your files order chronologically. This also makes it easier to see if you have any duplicate files.



**Figure S3**: Cleaning up your data and code using research Data Management practices is recommended before sharing your data and code. The Turing Way project illustration by Scriberia. Used under a CC-BY 4.0 licence.

DOI: 10.5281/zenodo.3332807.

In spreadsheets, put as little information as possible in a single cell and only one observation per row (Broman & Woo 2018). You can share additional information in a README file or in a data dictionary (Buchanan et al. 2021) or code book (Kent State University 2022) that describes the spreadsheet and any cleaning steps you took. In your data, avoid formatting to describe the data (colours, font, bolding). Instead, add additional cells for the information that this formatting should be conveying. You can also use data validation to avoid errors. Excel and OpenRefine (https://openrefine.org) have several options that you can use. For more spreadsheet tips see the Carpentries curriculum on spreadsheets for ecologists (Hoyt et al. 2019) and social scientists (Data Carpentry 2021), Hao Ye's work (Ye 2020) and information on The Turing Way (The Turing Way Community 2022g).

To manage your code it can be helpful to use Git/GitHub to keep better track of any modifications made (and by whom) (Perez-Riverol et al. 2016). If you share your research software from the start you will also structure it differently and more readable to others than you would if you would if you kept it closed. For software it should be clear what language and environment you are using, and if there are any dependencies and/or packages needed to process the data in a similar fashion as the analysis conducted for the study. See 'Make sure that your code is in a sharable state' (Schlauch 2021) and Krafczyk et al. (2021, p5-11) for more details about how to ensure your code is ready to be shared. Also see the article by Perkel (2020) on the irreproducibility of code written years ago. Once you're more experienced with using GitHub you can use GitHub actions to prevent errors in data collection (Kim et al. 2022)

Add a README file to your dataset or your software repository. README text files should describe the methods used for data collection and analysis and include data/software-specific information (parameters, variables, column headings, symbols used, etc.). See <a href="Make a README">Make a README</a> (Guo 2021) for more information on why README files are important and how you can set up your own. You can use README files from existing projects and datasets as examples or inspiration (for <a href="example for data">example for data</a> (Cornell University 2022) and a <a href="mailto:general">general</a> (Drew 2022) and <a href="mailto:archaeological example">archaeological example</a> (Marwick 2022b) for code).

#### How do you share research methods more openly?

Research methods are the processes that generate research data. Using different methods, or adapting certain steps of a procedure, can affect the resulting research data. To increase the reproducibility of your work it is therefore crucial to make methods more openly available. Methods can include wet lab protocols, software analyses, strategies for surveys (see Strupler & Wilkinson 2017) and may involve various types of equipment. Methods shared on platforms such as <a href="mailto:protocols.io">protocols.io</a> (<a href="https://protocols.io">https://protocols.io</a>) can facilitate reuse of the data or the method you used, as these platforms allow anyone to set up a copy of the method (forking).

#### Examples are

- Article by Cerasoni (2021a) and accompanying protocol (Cerasoni 2021b) on stone tool illustrations and Matzig (2021) on an R-package for artefact processing.
- Protocol by Thaler & Gneisinger (2021) on use-wear experiments.
- Protocols by Brown et al. (2021) on **ZooMS Spectra**.
- Protocols by Plomp et al. (2019 and 2020) on isotope analysis (neodymium and strontium respectively).
- Protocols on dental calculus sampling by Warinner et al. (2020), Sabin & Fellows Yates (2020),
   Wilkin et al. (2021).
- Protocols on **3D models** by Tang et al. (2022), Falcucci (2022) and Göldner et al. (2022).
- Protocol on **DNA** sampling by Orfanou et al. (2020).

#### What is metadata?

Metadata is information about the data. These could range from your notes about data collection and processing to the information that you are required to fill in when you deposit data in a data repository. The last type of metadata is machine readable and will facilitate data discovery (see <u>FAIR</u>). Most data repositories, such as Zenodo and Figshare, will use standardised schemes of these information fields - such as <u>Dublin Core</u> (Wikipedia 2022). Standardised metadata, or a metadata standard, will enhance the interoperability of information as similar descriptions are used which should make it easier to integrate data. The integration of studies would allow archaeologists to address research questions on a larger scale. You can start small by searching for metadata standards using <u>FAIRsharing</u> (<a href="https://fairsharing.org">https://fairsharing.org</a>) or start discussions in your subfield about how to standardise data documentation.

To our knowledge, archaeology has these specific metadata standards:

- CIDOC CRM for **field surveys** (de Haas & van Leusen 2020).
- Monument Inventory DAta Standard Heritage (MIDAS Heritage), for recording **heritage** information on buildings, archaeological sites, shipwrecks, parks and gardens, battlefields, areas of interest and artefacts (FAIRsharing Team 2018c).
- Art and Archaeology Vocabulary employed for indexing bibliographical records for the "Art and Archaeology" FRANCIS database (FAIRsharing Team 2018a).
- FISH Archaeological Sciences Thesaurus (FISH-AST) for recording **techniques**, **recovery methods** and materials (FAIRsharing Team 2018b).
- CARARE Metadata Schema for an **organisation's online collections**, heritage assets and their digital resources (FAIRsharing Team 2022).
- Thesaurus of Geographic Names (TGN) terminology that focuses on recording names, relationships, place types, dates, notes, and coordinates for current and historical cities, nations, empires, archaeological sites, lost settlements, and physical features (FAIRsharing Team 1987).
- MetaShARK for **ecological data** (Earnaud *et al.* 2021).
- For Ecology metadata standards, also see Michener et al. (1997).

Other metadata standards that could be useful are:

- The <a href="RFC-3339">RFC-3339</a> (Kleyne et al. 2002) or ISO 8601 standards, which specify the order in which dates are written: YYYY-MM-DD.
- ISO 19115 (ISO 2014) for **geographic** information.

To learn more about metadata, visit the Archaeology Data Services website (Niven 2011).

#### What about licenses and copyright?

Licenses govern what someone else can do with data and software that you share. The various licenses have different criteria about what is allowed when the data/software is reused, and there are different types of licenses available for data and software.

- For data the <u>Creative Commons Licenses</u> (Creative Commons 2022), or <u>Open Data Licenses</u> (Open Data Commons 2022) are most often used. For example, the CC-BY license for data requires that the reuser provides attribution for data re-use through, for example, citation.
- For **software** the <u>Choose a License</u> website (Choose a License 2022) provides an overview of the available licenses. An often used license for software is the MIT license, that similarly to the CC-BY licence, requires attribution for reuse.

For both data and software it is important to follow the license requirements. Sometimes these requirements are in conflict, or incompatible. Incompatible licenses can get especially complex when you want to reuse software created by others. This makes combining datasets or software difficult, which is something to keep in mind when you choose a more restrictive license for your outputs. The fewer restrictions a license has, the easier it is for others to reuse your work (for data CCO or CC-BY, for software MIT). If you are unsure whether you are complying with license requirements, check if your institution provides any advice on this. Generally this type of support is available from the Library or a copyright support desk.

#### Isn't reproducible archaeology more expensive?

It is a misconception that working with an open science approach is more expensive. This idea of higher cost stems from the well-known high costs of **gold open access** journal articles and also dedicated archaeological data repositories being commercial businesses that charge for data deposition. See 'Getting started with open repositories - part 1 - what you might think' for more information about this misconception - (Karoune 2021b).

In fact, everything that you would want to do openly with your research can be done for free using free open-source software, free tools and apps such as GitHub and Google Drive, and free data repositories such as Zenodo, Open Science Framework, Figshare and Dataverse.

- Depositing data and other research outputs: There is a wide choice of free and open repositories
  for depositing data and other research outputs. This might be through your own institution or one
  of the large public infrastructure repositories such as Zenodo, Open Science Framework, Dataverse
  or Figshare.
- **Software for open analysis:** To use the R coding language for analysis, you can use RStudio. It is free to download and there are many packages that allow you to do the types of statistical analysis, which you would have done in expensive proprietary software such as SPSS.
- Publishing open access: You can make your articles open access for free using the green or diamond open access routes:
  - Green open access is where you deposit a version of your article (not the final formatted version that will be in the journal but a preprint or postprint version) on an open repository such as a preprint server (some examples are arXiv, bioRxiv, or EarthArXiv) or one of the data repositories mentioned above. This can be done at no cost to you or the reader. The majority of journals allow you to do this, but carefully check the journal's guidelines. See <a href="Sherpa Romeo">Sherpa Romeo</a> for details of journal policies (Jisc 2022).
  - O You can also use diamond open access, which is free for authors to publish and free for readers type of open access that some journals offer such as those paid for by societies, associations or communities such as Peer Community in Archaeology and PaleoAnthropology.
- Version control for open reproducible workflows: For simple version control, you can use Google
  Drive. There are free advanced version control tools that you can use based on Git GitHub or
  GitLab. An alternative to Git is <u>Subversion</u> (<a href="https://subversion.apache.org">https://subversion.apache.org</a>) also a free and opensource software.

#### What if people misinterpret data or find a mistake?

To avoid misinterpretation of the data you should provide sufficient information about your dataset and all the data required for appropriate reuse (**Figure S4**). You can also list your contact details in the documentation or readme file so that reusers can contact you with questions or concerns. You could, for example, set up an <a href="ORCID">ORCID</a> (<a href="https://orcid.org">https://orcid.org</a>), a persistent identifier for researchers that you own and control, with your contact details to ensure that reusers are able to find you. ORCIDs are particularly beneficial if you have a common name or if you expect to switch between institutions in the future.

# DOCUMENTATION



**Figure S4**: Clearly written and available documentation will allow others to follow the steps you took in the research process, preventing misinterpretation. The Turing Way project illustration by Scriberia. Used under a CC-BY 4.0 licence. DOI: 10.5281/zenodo.3332807.

As you are only human it is entirely possible that there is a mistake in your data or script. Keep in mind that if anyone would find an error in your data that this means that your dataset is engaging and relevant (Strupler 2021). To prevent errors you can use the guidance from <a href="Error Tight">Error Tight</a> to set up a workflow in the lab that makes it more likely that mistakes made in the lab are caught early (Strand 2021). You can also minimise mistakes in your own research outputs by asking someone from your lab to check your data or code before making it more widely available, for example, by trying to reproduce your work (a co-pilot, see Reimer et al. 2019).

Even after close scrutiny by a colleague it could be that someone discovers a mistake after you shared the data or code publicly. Most data repositories allow you to upload a new version of the data/code where you can explain in the documentation what has changed in this new version and why. Correcting this mistake may save the re-users of your data and code, and yourself, a lot of time and may increase the trustworthiness of your data and code as you facilitate the self-correcting nature of the scientific process. Research shows that improving the original work can have a beneficial effect on your reputation (Ebersole *et al.* 2016).

#### Is archaeology suitable for preregistration?

A *preregistration* is a document in which the research design, and sometimes hypotheses, is specified before research is carried out. This could also be done through a *registered report*. Preregistering your research may structure your data collection, management and analysis which can result in more robust research, reusable datasets and reduce the time spent managing problems and data cleaning on a more ad hoc manner (Ross & Ballsun-Stanton 2021). Ross & Ballsun-Stanton (2021) argue that preregistration is beneficial for archaeologists. Preregistration encourages a more thoughtful approach to research design, better management of biases through making approaches and assumptions more explicit, and it encourages good practices in research transparency (Ross & Ballsun-Stanton 2021). Good practice around archaeological preregistration is still emerging, but Ross & Ballsun-Stanton (2021) offer some helpful pointers.

#### A supervisor doesn't want to work reproducibility, how do you convince them?

There are several strong arguments to make for moving to a reproducible research workflow (**Figure S5**). Many funders are now requiring more open practices. The <u>UK Research and Innovation</u> (2021) and the <u>European Research Council</u> (2022) both have policies requiring immediate open access publishing through Gold or Green Open access for all grant holders. These publications must be linked to all research outputs to validate research. This means that your supervisor will have to start opening up their work to some extent and it would be good to learn how to do this well now.

Additional links to funding policies:

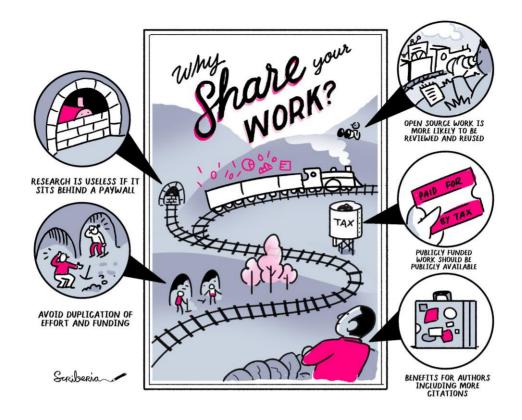
- <u>Wellcome Trust funding guidance</u> (Wellcome 2022) including open access policy, responsible conduct of research and data, software and materials management and sharing.
- Australian open access policy (ARC 2021).
- National Science Foundation from the USA (NSF 2015).
- Draft Open Access Policy from India (Open Access India 2017).

Similarly to changing funding requirements, the importance of the published research articles is likely to change in the upcoming years. Several individuals have already called the stand-alone scientific paper outdated (Marwick *et al.* 2017; Krumholz 2015), obsolete (Somers 2018), or dead (Robert Terry during the second UNESCO Conference on Open Science - Link to video (Terry 2021)). While the scientific paper has not yet died, the journals have requirements that your work should fulfil before it will be published. Increasingly, this includes making the underlying data and code available (Hrynaszkiewicz 2019), see for example the American Journal of Physical Anthropology requirements (Turner & Mulligan 2019, Wiley 2022). Even if journals do not have these requirements, it may be that your reviewers ask to see the underlying code and data (Stark 2018). Sharing the data/code during the peer review process may thus result in improvements of your work or faster acceptance as the reviewer does not have to wait for access (Markowetz 2015).

Having a reproducible workflow, which is transparent and open, has greater research impact. This has now been proven in a number of ways. Open access publications are known to have a citation advantage over publications behind paywalls (Langham-Putrow *et al.* 2021). It has also been found that linking open data to your article increases citations significantly (Piwowar *et al.* 2007; Piwowar & Vision 2013; Christensen *et al.* 2019; Colavizza *et al.* 2020).

You could also consider publishing more articles by writing a data or software paper for your project. There are specific journals for data and software papers (Journal of Open Research Software and Journal of Open Source Software). Data journals are subject specific such as the Journal of Open Archaeology Data or more general such as Scientific Data. Publishing these additional types of articles would give you credit for the extra work that you are doing to produce a reproducible workflow and also increase the overall outputs of the project, increasing the impact.

Moving to reproducible workflows is going to take time and it will help to talk about the benefits within your research group to encourage others to follow your example. Find allies within your department or other people within your subfield that do work reproducible to convince the supervisor that this is a good thing.



**Figure S5**: Benefits of sharing your work openly. Research is useless if it is not accessible and sitting behind a paywall. Through sharing your work you can avoid duplication of effort and waste of funding. Publicly funded work should be publicly available as it is paid for by taxpayers. Open source work is more likely to be reviewed and reused and can generate more citations. The Turing Way project illustration by Scriberia. Used under a CC-BY 4.0 licence. DOI: 10.5281/zenodo.3332807.

#### Will reproducible research be taken into account when looking for a next job?

Academic institutes are changing the focus of research evaluations, moving away from the impact factor of articles to a more broader evaluation that also takes into account education, open science practices and leadership. Examples are the <u>TRIPLE model at Utrecht University</u> in the Netherlands (Utrecht University 2021).

Making your work openly available will help build your reputation for being an honest and careful researcher (Markowetz 2015). Experience with Open Science practises is also increasingly asked for in vacancies (Schönbrodt *et al.* 2021).

Funding bodies are increasingly asking about data and software management and the sharing of these research outputs. Moving towards sharing these outputs will therefore outweigh the costs in the long term by increasing your chances for funding and by improving your sharing workflows earlier rather than later.

Next to improving your chances on the academic job market, open data and code can also be useful in positions elsewhere, such as in industry where the demand for computational skills is high (<u>Anaconda 2020</u>; Kim *et al.* 2018).

#### Do platforms like SciHub, ResearchGate, Academia.edu count as Open Access?

Platforms such as SciHub, ResearchGate, and Academia.edu do not count as sustainable Open Access. SciHub, while providing access to research more widely, is not a legal platform and is hosted by a single individual. This makes long term sustainability questionable, and the founder, Alexandra Elbakyan, is dealing with multiple lawsuits.

Academia.edu is not an educationally-affiliated organisation and instead monetising scholarly outputs. By agreeing to their privacy policy Academia.edu is furthermore able to sell your information to other companies (<u>Tóth Czifra 2020</u>). ResearchGate has been subjected to lawsuits that determined that the

platform is responsible for copyright infringement, which can result in the removal of the papers that they made openly available (Kwon 2022). ResearchGate and Academia.edu are also not open about their business and sustainability models, or interoperable with other services (Fitzpatrick 2020).

While Academia.edu and ResearchGate are good for advertising your research and networking like other social media platforms, you might be illegally sharing copyrighted work through these platforms. If your article has a CC-BY-NC-ND license, you are not allowed to share it on Academia.edu and ResearchGate as these are commercial platforms which are excluded by the NC part of the license (Non-Commercial). This can be circumvented by choosing a CC-BY license so that you are allowed to share it on these platforms, as you retain the rights to your work and there are no commercial reuse restrictions.

You can also share your work via a preprint or **postprint** version under an open license through more sustainable solutions such as data repositories and preprint servers. Institutions can also play a role here by retaining control of the infrastructures that provide access to research outputs.

An example of scholarly communities retaining control of all the infrastructure involved in making research available is the <a href="Peer Community">Peer Community in Archaeology</a> (<a href="https://archaeo.peercommunityin.org">https://archaeo.peercommunityin.org</a>) platform and <a href="IsoArcH database">IsoArcH database</a> (<a href="https://isoarch.eu">https://isoarch.eu</a>, Salesse et al. 2018). The Peer Community in Archaeology are openly reviewing and recommending preprints therefore increasing the transparency of quality control processes. Disciplinary specific repositories such as IsoArcH (for bioarchaeological isotope data) increase the impact of datasets, as they are curated by specialists and accompanied by the relevant metadata, which makes the data more reusable.

If you would like to learn more about Open Access in archaeology, see Kansa et al. (2013).

#### Appendix B: Glossary of terms used in this paper

Some definitions are adapted from The Turing Way Glossary (The Turing Way Community 2022f).

**Binder** - The Binder Project is a software project to package and share interactive, reproducible environments. It creates an interactive notebook or workspace for running code in the computational environment that the author of the analysis used. It should produce the ability to reproduce analysis and/or figures using one press of a button. A Binder or 'Binder-ready repository' is a code repository in GitHub that contains both code and content to run, and configuration files for the environment needed to run it.

**Computational environment** - Features of a computer which can impact the behaviour of work done on it, such as its operating system, what software it has installed, and what versions of software packages are installed.

**Container** - A container is a standard unit of software that packages up code and all its dependencies so the application runs quickly and reliably from one computing environment to another.

Data availability statement - A data availability statement (also sometimes called a 'data access statement') tells the reader where the research data associated with a paper is available, and under what conditions the data can be accessed. They also include links using a DOI to the data set, code and other documentation.

**Data Management Plan** - A Data Management Plan (DMP), or Output Management Plan, is a document that describes how your research outputs will be generated, stored, used and shared within your project.

**Data Repository** - A long-lived place on the internet where resources (be they data, software, publications or anything else) can be stored and accessed via a persistent identifier (such as a DOI). This keyword is often shortened to 'repo'.

**Diamond Open Access** - refers to academic texts (such as monographs, edited collections, and journal articles) published/distributed/preserved with no fees to neither reader nor author. Most diamond open access journals are managed by academic institutions, communities, government agencies, learned societies or sometimes individuals.

**Digital Object Identifier** - A Digital Object Identifier (DOI) is a persistent identifier or handle used to identify objects uniquely, standardized by the International Organization for Standardization (ISO). An implementation of the Handle System, DOIs are in wide use mainly to identify academic, professional, and government information, such as journal articles, research reports, data sets, and official publications. However, they also have been used to identify other types of information resources, such as commercial videos.

**Generalisable** - Combining replicable and robust findings allow us to form generalisable results. Note that running an analysis on a different software implementation and with a different dataset does not provide generalised results. There will be many more steps to know how well the work applies to all the different aspects of the research question. Generalisation is an important step towards understanding that the result is not dependent on a particular dataset nor a particular version of the analysis pipeline.

**Git** - is a free and open source distributed version control system. Git is the version control system (software) behind advanced version control tools such as GitHub and GitLab.

**GitHub** - is a code hosting platform for version control and collaboration. It is used for storing, tracking and collaborating on software and research projects. GitHub enables researchers and developers to upload their own data, code and documentation and to collaborate with others.

**Gold open access** - the publisher makes all articles and related content available for free immediately on the journal's website. In such publications, articles are licensed for sharing and reuse via creative commons licenses or similar. An article processing charge (APC) is paid by the authors.

**Green open access** - Independently from publication by a publisher, the author posts the work to a website controlled by the author, the research institution that funded or hosted the work, or to an independent central open repository, where people can download the work without paying. This can be a preprint (version of article prior to peer preview) or postprint (version that has been peer reviewed). This is free for the author.

**Metadata** - the data/information about the data. This can include information about who collected the data and when, and also the methods used for data collection.

**Paradata** - Paradata of a data set or survey are data about the process by which the data were collected. **Persistent Identifier** - A long-lived method for identifying a resource that is unique, and widely understandable by a community. This includes ORCIDs as an identifier of researchers and digital object identifiers (DOI) as identifiers of research objects.

**Postprint** - is the version of an article that incorporated changes from the peer review process, but does not yet have publication formatting or layout applied. It is usually uploaded by the authors to a public or institutional server where it is available openly.

**Preprint** - is a version of an article that precedes formal peer review and publication in a peer-reviewed journal. Like postprints, authors generally upload this version of the article themselves using a public/institutional server where it is available openly.

**Preregistration** - is the practice of registering the research design of the research project before it is conducted. This includes details of hypotheses, methods and proposed analysis steps.

**Proprietary software** - is software that requires a paid license to be able to use it and it is closed-source (the code behind the software and the code that you produce in your analysis is not available to see).

**Protocol** - is another term for method. Protocols are often step-by-step instructions that could be used in wet-lab or other practical research. In principle it can be applied to any method that contains several steps.

**Python** - is a high-level, interpreted, general-purpose programming language. Its design philosophy emphasises code readability with the use of significant indentation.

**R** - is a programming language for statistical computing and graphics supported by the R Core Team and the R Foundation for Statistical Computing.

**Registered report** - is a type of publication that is written before the research is conducted and includes the research question/s, methodology and proposed analysis steps. It is then peer reviewed prior to data collection.

**Replicable/Replication** - A result is replicable when the same analysis performed on different datasets produces qualitatively similar answers.

**Reproducible** - A result is reproducible when the same analysis steps performed on the same dataset consistently produces the same answer.

**Reproducible workflow** - a transparent record of the research that includes data, methods, and analysis to allow other researchers to review, reproduce and replicate the study.

**Robust** - A result is robust when the same dataset is subjected to different analysis workflows to answer the same research question and a qualitatively similar or identical answer is produced. Robust results show that the work is not dependent on the specificities of the programming language chosen to perform the analysis.

**Transparent recording** - the complete recording of every step of the research lifecycle. This can be done through thorough documentation or by using advanced version control systems such as Git and GitHub.

**Version Control** - is a systematic approach to record changes made in a file, or set of files, over time.