



# FAIR Data and You: Are you ready for 2023?

Learn how to succeed with your next NIH grant:  
New Data Mandates and Opportunities

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# Topics for Today's Webinar

1. Open science and data sharing
2. Data as a research output
3. What are the FAIR principles?
4. Why were the FAIR data principles introduced?
5. A deeper dive into some of the FAIR principles
  - a. Use of persistent identifiers
  - b. Rich metadata
  - c. Data licenses
  - d. FAIR vocabularies
  - e. Community standards
6. Introduction to data repositories
7. FAIR data and you: Importance of Data Management



# About dkNET

- Research resource information portal for biomedical researchers
- Information network to connect DK researchers and NIH-funded resources and centers
- Funded by National Institute of Health (NIH) - National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK)
- Developed and maintained by the FAIR Data Informatics Laboratory (fdilab.org) at UCSD (Supports major informatics projects in biomedicine)





# Get involved in the dkNET Community

## dkNET Homepage: dkNET.org

Blog, Calendar

The screenshot shows the dkNET homepage with the following elements circled:

- Navigation Bar:** A red circle highlights the top navigation menu with links: ABOUT, RESOURCE REPORTS, DISCOVERY PORTAL, AUTHENTICATION REPORT, and HYPOTHESIS CENTER.
- Resource Reports:** An orange circle highlights the 'Resource Reports' section, which includes a question about antibody specificity and links to tools like Cell lines, Antibodies, Organisms, Plasmids, Biosamples, and Protocols.
- Authentication Reports & FAIR Data:** An orange circle highlights the 'Authentication Reports & FAIR Data' section, which discusses NIH's new policies and provides links to reports, research management, and data repositories.
- Discovery Portal:** An orange circle highlights the 'Discovery Portal' section, which offers a search across 100s of biomedical databases for funding, images, phenotypes, literature, and more.
- Hypothesis Center:** An orange circle highlights the 'Hypothesis Center' section, which allows users to analyze diverse 'omics data to generate or test research hypotheses.
- Webinar Series:** An orange circle highlights a 'dkNET Webinar Series' card for 'Appyters: Turning Jupyter Notebooks into Data-Driven Web Apps' on Friday, May 28, 2021.
- Newsletter:** An orange circle highlights a 'Newsletter' card announcing that this month's newsletter is now available, offering funding opportunities, events, and community news.
- Virtual Booth:** An orange circle highlights a 'Visit dkNET Virtual Booth Anytime, Anywhere' card with the 'Ask dknet.org' button.
- Footer:** A green circle highlights the footer navigation: Community News, Blog, News, Live Webinars, On-Demand Webinars, and Subscribe to dkNET mailing list. Another green circle highlights a tweet from @dkNET\_info.

Join Webinar

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# Open science

...the movement to make scientific research (including **publications, data, physical samples, and software**) and its dissemination **accessible** to all levels of an inquiring society, amateur or professional. Open science is transparent and accessible knowledge that is shared and developed through collaborative networks. It encompasses practices such as publishing **open research**, campaigning for **open access**, encouraging scientists to practice **open notebook science**, and generally making it easier to publish and communicate scientific knowledge. -adapted from [Wikipedia](#)







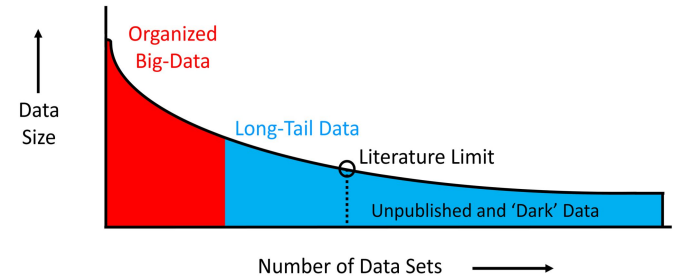
# Motivations for open science

- Consequence of digital revolution: access and computation
- Two main arguments in biomedicine:
  - Research is largely funded by the public and the public should have access to the results
  - Open science = better science
    - Transparency
    - More eyes
    - Increasing computational capacity that can mine biomedical literature and data for new insights



# Trends in Open Data

- “Long tail data”
  - Make data underlying claims in a paper available
  - Publish data as a primary product of research, like you would a paper
    - Transparency
    - Reuse
    - Aggregation to increase analytic power
- “Big science” : generate large pools of freely available and more usable data “AI/ML Ready”
  - Spur data science in biomedical science
- Funders and journals are increasingly requiring data sharing







# January 25, 2023

- US National Institutes of Health new data sharing policy goes into effect
- All data must be managed; most data should be shared
- “As open as possible; as closed as necessary”
- Mandates the ***inclusion, approval and execution*** of a Data Management and Sharing Plan
  - (DMP + S = DMS)

## Final NIH Policy for Data Management and Sharing

### Notice Number:

NOT-OD-21-013

### Key Dates

Release Date: October 29, 2020  
Effective Date: January 25, 2023

### Related Announcements

[NOT-HG-21-023](#) - Notice Announcing NHGRI Guidance for Third-Party Involvement in Extramural Research

[NOT-HG-21-022](#) - Notice Announcing the National Human Genome Research Institute's Expectation for Sharing Quality Metadata and Phenotypic Data

[NOT-OD-21-014](#) – Supplemental Information to the NIH Policy for Data Management and Sharing: Elements of an NIH Data Management and Sharing Plan

[NOT-OD-21-015](#) – Supplemental Information to the NIH Policy for Data Management and Sharing: Allowable Costs for Data Management and Sharing

[NOT-OD-21-016](#) – Supplemental Information to the NIH Policy for Data Management and Sharing: Selecting a Repository for Data Resulting from NIH-Supported Research

[NOT-OD-20-013](#) - Request for Public Comments on a DRAFT NIH Policy for Data Management and Sharing and Supplemental DRAFT Guidance

### Issued by

Office of The Director, National Institutes of Health (OD)

### Purpose

#### Summary

The National Institutes of Health (NIH) is issuing this final NIH Policy for Data Management and Sharing (DMS Policy) to promote the management and sharing of scientific data generated from NIH-funded or conducted research. This Policy establishes the requirements of submission of Data Management and Sharing Plans (hereinafter Plans) and compliance with NIH Institute, Center, or Office (ICO)-approved Plans. It also emphasizes the importance of good data management practices and establishes the expectation for maximizing the appropriate sharing of scientific data generated from NIH-funded or conducted research, with justified limitations or exceptions. This Policy applies to research funded or conducted by NIH that

<https://grants.nih.gov/grants/guide/notice-files/NOT-OD-21-013.html>



# The Details...

The effective date of the DMS Policy is January 25, 2023, including for:

- Competing grant applications that are submitted to NIH for the January 25, 2023 and subsequent receipt dates;
- Proposals for contracts that are submitted to NIH on or after January 25, 2023;
- NIH Intramural Research Projects conducted on or after January 25, 2023; and
- Other funding agreements (e.g., Other Transactions) that are executed on or after January 25, 2023, unless otherwise stipulated by NIH.



# More Details...

## What

- Defines Scientific Data as: “The recorded factual material commonly accepted in the scientific community as of sufficient quality to validate and replicate research findings, regardless of whether the data are used to support scholarly publications. Scientific data **do not include** laboratory notebooks, preliminary analyses, completed case report forms, drafts of scientific papers, plans for future research, peer reviews, communications with colleagues, or physical objects, such as laboratory specimens.”
- Even those scientific data not used to support a publication are considered scientific data and within the final DMS Policy’s scope

## When

- “[s]hared scientific data should be made accessible as soon as possible, and no later than the time of an associated publication, or the end of the award/support period, whichever comes first.”
- Researchers may share data underlying publication during the period of award but may share other data that have not yet led to a publication by the end of the award period.

## Where

- Encourages the use of established repositories to the extent possible.



# More Details...

## How

- NIH encourages data management and data sharing practices consistent with the FAIR data principles

## Funding

- Fees for long-term data preservation and sharing are allowable, but funds for these activities must be spent during the performance period, even for scientific data and metadata preserved and shared beyond the award period.

## Repercussions

- After the end of the funding period, non-compliance with the NIH ICO-approved Plan may be taken into account by NIH for future funding decisions for the recipient institution

The DMS Policy applies to all research, funded or conducted in whole or in part by NIH, that results in the generation of scientific data. This includes research funded or conducted by extramural grants, contracts, Intramural Research Projects, or other funding agreements regardless of NIH funding level or funding mechanism.



# Good data management is the gateway to data sharing

	<b>Ad Hoc</b>	<b>One-Time</b>	<b>Active and Informative</b>	<b>Optimized for Re-Use</b>
<b>Planning your project</b>	When it comes to my data, I have a "way of doing things" but no standard or documented plans.	I create some formal plans about how I will manage my data at the start of a project, but I generally don't refer back to them.	I develop detailed plans about how I will manage my data that I actively revisit and revise over the course of a project.	I have created plans for managing my data that are designed to streamline its future use by myself or others.
<b>Organizing your data</b>	I don't follow a consistent approach for keeping my data organized, so it often takes time to find things.	I have an approach for organizing my data, but I only put it into action after my project is complete.	I have an approach for organizing my data that I implement prospectively, but it not necessarily standardized.	I organize my data so that others can navigate, understand, and use it without me being present.
<b>Saving and backing up your data</b>	I decide what data is important while I am working on it and typically save it in a single location.	I know what data needs to be saved and I back it up after I'm done working on it to reduce the risk of loss.	I have a system for regularly saving important data while I am working on it. I have multiple backups.	I save my data in a manner and location designed maximize opportunities for re-use by myself and others.

Borghi J, Abrams S, Lowenberg D, Simms S, Chodacki J (2018) Support Your Data: A Research Data Management Guide for Researchers. Research Ideas and Outcomes 4: e26439.

<https://doi.org/10.3897/rio.4.e26439>



# Changing the culture around data management and sharing

- **Science and Society**

- Transparency
- Reproducibility
- Reduced waste
- *Driving discovery*

- **Me**

- Answer to the underpowered study
- Data sharing and good data management are closely aligned
- Compliance with mandates
- Credit for the totality of my work

- **Future me**

- One most likely to benefit from good data management and sharing through stable archives
- No one ever regretted annotating too much

- **My colleagues (and PI)**

- Easy to engage with colleagues over well annotated data and associated code
- What happens when the post doc leaves?



## But how do I do that?

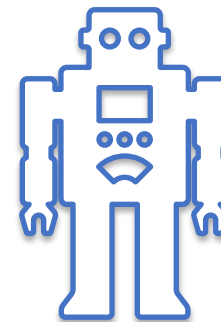
- “NIH encourages data management and data sharing practices consistent with the FAIR data principles.”
- “NIH strongly encourages the use of established repositories to the extent possible for preserving and sharing scientific data”



# The FAIR Guiding Principles for scientific data management and stewardship

High level principles to make data:

- Findable
- Accessible
- Interoperable
- Re-usable



...for humans *and* machines

Mark D. Wilkinson et al. The FAIR Guiding Principles for scientific data management and stewardship, *Scientific Data* (2016). [DOI: 10.1038/sdata.2016.18](https://doi.org/10.1038/sdata.2016.18)





# Definition: Machine readable

- "(meta)data in a format that can be easily processed by a computer without human intervention while ensuring no semantic meaning is lost." ...

There are two types of machine-readable data: human-readable data that is **marked up** so that it can also be read by machines (e.g. **microformats**, **RDFa**, **HTML**) and data file formats intended principally for processing by machines (**CSV**, **RDF**, **XML**, **JSON**).

Again, these formats are only machine readable if the data contained within them is formally structured; exporting a CSV file from a badly structured spreadsheet does not make the data machine-readable. -adapted from [Wikipedia](#)



# CSV file...or is it?

SubjectID	Subject Group	Species	Age	Unit	Sex	Strain
F001	1	Mouse	25 days	F	C57Bl/6	
F002	1	Mouse	23 days	M	C57Bl/6	
F003	1	Mouse	26 days	M	C57Bl/6	
F004	1	Mouse	25 days	M	C57Bl/6	
F005	1	Mouse	24 days	F	C57Bl/6	

SubjectID	Subject Group	Species	Age	Unit	Sex	Strain
F001	1	Mouse	25 days	F	C57Bl/6	
F002	1	Mouse	23 days	M	C57Bl/6	
F003	1	Mouse	26 days	M	C57Bl/6	
F004	1	Mouse	25 days	M	C57Bl/6	
F005	1	Mouse	24 days	F	C57Bl/6	

Good resource: <https://schoolofdata.org/courses/>



# Metadata

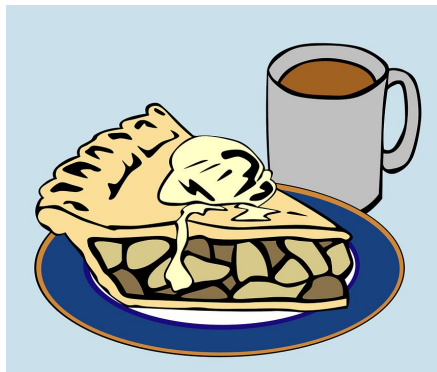
- “Data about about data” e.g., file format, file size
- ***Descriptive metadata*** describes a resource for purposes such as discovery and identification. It can include elements such as title, abstract, author, keywords, study organism, condition studied etc

<https://en.wikipedia.org/wiki/Metadata>



MOM

A closer look at FAIR





## Findable

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

## Interoperable

- I1. (meta)data use a formal, accessible, shared, and broadly applicable language for knowledge representation.
- I2. (meta)data use vocabularies that follow FAIR principles
- I3. (meta)data include qualified references to other (meta)data

## Accessible

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

## Re-usable

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



# Findable



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



# Huh?

- “Principle F1 is arguably the most important because it will be hard to achieve other aspects of FAIR without globally unique and persistent identifiers. Hence, compliance with F1 will already take you a long way towards publishing FAIR data”-[GoFAIR](#)
  - Unique in the world + Stable (persistent)
    - Identify only a single object for all time (never reused)
    - Only persistent and unique because organizations stand behind them\*\*\*
  - Can be resolvable, i.e., you can plug it into a web browser and be taken to the object ***independent of its location***
  - Allows a digital object to be reliably tied to its metadata

F1. (meta)data are assigned a ***globally unique and persistent identifier***



# Some examples of PIDs

Front Neuroinform. 2016 Apr 19;10:11. doi: 10.3389/fninf.2016.00011. eCollection 2016. [Paperpile](#)

## Brain-Wide Mapping of Axonal Connections: Workflow for Automated Detection and Spatial Analysis of Labeling in Microscopic Sections.

Papp EA<sup>1</sup>, Leergaard TB<sup>1</sup>, Csucs G<sup>1</sup>, Bjaalie JG<sup>1</sup>

### Author information

#### Abstract

Axonal tracing techniques are powerful tools for exploring the structural organization of neuronal connections. Tracers such as biotinylated dextran amine (BDA) and Phaseolus vulgaris leucoagglutinin (Pha-L) allow brain-wide mapping of connections through analysis of large series of histological section images. We present a workflow for automated detection of neuronal labeling in large image series, aligned to a Waxholm Space (WHS) atlas, which different parts of the rat primary somatosensory cortex were used to automate detection of labeling in individual images. For high to medium labeling densities, automatic detection of neuronal labeling in large image series, aligned to a Waxholm Space (WHS) atlas, match individual sections. Based on the alignment, WHS coordinates. The new workflow modules increase the efficiency of labeling in individual sections, and enable anchoring to anatomical atlases for individual sections.

**KEYWORDS:** automated image processing; axonal tract tracing;

PMID: 27148038 PMCID: [PMC4835481](#) DOI: [10.3389/fninf.2016.00011](#)

PMID: 27148038 PMCID: [PMC4835481](#) DOI: [10.3389/fninf.2016.00011](#)

The screenshot shows the ORCID iD profile for Maryann Elizabeth Martone. The profile includes a biography, a list of keywords, and a list of works. The biography states that she received her BA from Wellesley College in biological psychology and her Ph.D. in neuroscience from the University of California, San Diego. She is currently the principal investigator of the Neuroinformatics Framework project. The keywords listed are Neuroinformatics, neuroscience, FORCE11, Neuroscience Information Framework, and ontologies. The works section shows 50 works, with a list of categories: Education (2), Employment (3), Funding (10), and Works (50 of 83).

- DOI: Digital object identifier
- ORCID: Researcher identifier
- RRID: Resource Identifier
- **Globally unique: identifies one thing only**
- Unlike URL's or catalog numbers, may **NOT** be re-used
- **Issued by registries who track and identify unique entities**





# PIDS in action: DOI

- Identifies an object regardless of its location
- Issued by authorities: DataCite and CrossRef who work with data repositories to make sure that the links don't break
- Issued by data repositories who agree to keep the links up to date
- A PID is a social contract!



**HIRN RESOURCE BROWSER**

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[Home](#) / [Dataset](#) / [Epigenomics](#) / Raw and processed imaging mass cytometry (IMC) data from an assay using a panel of 37 antibodies, of pancreas sections from 12 human donors at different stages of type 1 diabetes

**Resource Name**  
Raw and processed imaging mass cytometry (IMC) data from an assay using a panel of 37 antibodies, of pancreas sections from 12 human donors at different stages of type 1 diabetes

**Export**  
**Canonical ID / Source** [doi:10.17632/cydmwfsztj.1](https://doi.org/10.17632/cydmwfsztj.1)

Name	Organization	Consortium	Contact
Clive Wasserfall	University of Florida		<a href="#">Contact</a>
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DOI: 10.17632/cydmwfsztj.1

<https://doi.org/10.17632/cydmwfsztj.1>



# Lesson: You cannot assign PIDs to your data yourself

You obtain a PID from a trusted repository that provides PIDs

<https://dknet.org/about/Suggested-data-repositories-niddk>



## Suggested data repositories

Home / Suggested data repositories

### Where can I deposit my data?

We have organized a list of data repositories that are recommended by the following sources: NIDDK domain experts, Nature Scientific Data, PLOS One, NLM NIH Science. It is generally best practice to deposit data into a discipline-specific and community recognized repository if one is available, or into an institutional or generalist repository if available.

- NIDDK-specific repositories
- NIH-supported repositories
- Institutional repository
- Other NIDDK Project-specific or consortium-specific data or sample repositories

### By scientific disciplines

- All research data types
- Chemistry and chemical biology and biochemistry
- Cytometry and Immunology
- Functional Genomics
- Imaging
- Metabolomics
- Molecular and supramolecular structure
- Neuroscience
- Nucleic acid sequence
- Other domain-specific repositories
- Protein sequence
- Proteomics

### De-identified human clinical research data

Clinical trial data is encouraged to be submitted to the [ClinicalTrials.gov](https://clinicaltrials.gov) even if it is not required. For studies include human genomic and associate phenotype database of Genotypes and Phenotypes (dbGaP). Another repository that you can consider is ICPSR, which hosts a variety of human data, including many of science studies. Information on uploading data to ICPSR can be found [here](#). Before uploading data, please note that the data should be de-identified, and you institutional IRB's requirements and receive approvals. For completed phase I-IV interventional studies, you can also share anonymized data at [Vivli](https://vivli.org).

### NIH-supported repositories (for complete and current list of NIH repositories click here)

Repository Name	RRID	Description	Type of Data	Recommended By
Accelerating Medicines Partnership Type 2 Diabetes Knowledge Portal (AMP-T2D)	RRID:SCR_003743	Portal and database of DNA sequence, functional and epigenomic information, and clinical data from studies on type 2 diabetes and analytic tools to analyze these data. <a href="#">Provides data and tool...[more]</a>	Array, exome sequencing, whole genome sequencing data	NLM, NIDDK
Analysis, Visualization, and Informatics Lab-space (AnVIL)	RRID:SCR_017469	Portal to facilitate integration and computing on and across large datasets generated by NHGRI programs, as well as initiatives funded by National Institutes of Health or by other agencies th... <a href="#">[more]</a>	Genomic data	NLM, NIDDK
Biological General Repository for Interaction Datasets (BioGRID)	RRID:SCR_007393	Curated protein-protein and genetic interaction repository of raw protein and genetic interactions from major model organism species, with data compiled through comprehensive curation efforts.	Molecular interaction data	NLM, NIDDK, PLoS ONE, Sci Data
Cancer Imaging Archive (TCIA)	RRID:SCR_008927	Archive of medical images of cancer accessible for public download. All images are stored in DICOM file format and organized as Collections, typically patients related by common disease (e.g. ... <a href="#">[more]</a>	Primary DICOM image datasets from cancer patients and analysis datasets	NLM, NIDDK, PLoS ONE, Sci Data
Cancer Nanotechnology Laboratory (caNanoLab)	RRID:SCR_013717	Data sharing portal designed to facilitate information sharing across international biomedical nanotechnology research community to expedite and validate use of nanotechnology in biomedicine.	Physico-chemical, in vitro and in vivo assays data that characterize nanomaterials. *This is a curated resource which may not accept direct submission of data. Contact the database directly f... <a href="#">[more]</a>	NLM, PLoS ONE, Sci Data
Cell Image Library (CIL)	RRID:SCR_003510	Freely accessible, public repository of vetted and annotated microscopic images, videos, and animations of cells from a variety of organisms, showcasing cell architecture, intracellular funct... <a href="#">[more]</a>	Microscopic imaging data	NLM, NIDDK
ClinicalTrials.gov	RRID:SCR_002309	Registry and results database of federally and privately supported clinical trials conducted in United States and around world. Provides information about purpose of trial, who may participat... <a href="#">[more]</a>	Clinical trial	NLM, NIDDK, Sci Data
DNA DataBank of Japan (DDBJ)	RRID:SCR_002359	Maintains and provides archival, retrieval and analytical resources for biological information. Central DDBJ resource consists of public, open-access nucleotide sequence databases including r... <a href="#">[more]</a>	Gene sequence	NLM, NIDDK, Science, PLoS ONE, Sci Data
Database of Interacting Proteins (DIP)	RRID:SCR_003167	Database to catalog experimentally determined interactions between proteins combining information from a variety of sources to create a single, consistent	Protein interaction data	NLM, NIDDK, PLoS ONE, Sci Data



## F2: Data are described with rich metadata

- What exactly are rich metadata?
- “I shall not today attempt further to define the kinds of material I understand to be embraced within that shorthand description, and perhaps I could never succeed in intelligibly doing so. But *I know it when I see it...*”- Justice Stewart Potter





# We know how to publish papers so they can be found by search engines, understood and reused by people we don't know

**concise and brief title**

**Safety of Propofol Sedation for Pediatric Outpatient Procedures**

Reagan Larsen, MD, David Galloway, MD, Sheetal Wadera, MD, Dean Kjar, MS, David Hardy, MD, Curtis Mirkes, DO, Lori Wick, MD, and John F. Pohl, MD

**abstract**

Propofol sedation is used more frequently in pediatric procedures because of its ability to provide varying sedation levels. The authors evaluated all outpatient pediatric procedures using propofol sedation over a 6-year period. All sedation was provided by pediatric intensivists at a single institution. In all, 4716 procedures were recorded during the study period; 15% of procedures were associated with minor complications,

whereas only 0.1% of procedures were associated with major complications. Significantly more major complications associated with propofol occurred during bronchoscopy ( $P = .001$ ). Propofol administered by a pediatric intensivist is a safe sedation technique in the pediatric outpatient setting.

**Keywords:** propofol; sedation; intensivist; pediatric

**I. Introduction:**

Propofol (2,6-diisopropylphenol) is a global central nervous system depressant that has a sedative effect by activating the GABA<sub>A</sub> receptor, inhibiting the NMDA receptor, and modulating calcium influx through slow calcium ion channels. It is an effective medication as it can provide varying levels of sedation using the physician's ability to titrate by intravenous infusion its amnesia effect while providing a minimum level of analgesia.<sup>1</sup> Propofol has a rapid onset of anesthetic action and rapid recovery time, which makes this medicine ideal for use in the outpatient setting. Also, propofol can be useful for pediatric sedation in which procedural goals should include patient safety, maximum physical and emotional comfort, improved patient cooperation, and successful completion of the procedure.<sup>2</sup> Increasing numbers of pediatric patients are undergoing complex diagnostic and therapeutic procedures, which makes sedation necessary. Studies have shown that children who undergo such procedures can find the experience to be worse than the actual disease process, suggesting that propofol use may be beneficial in the pediatric population.<sup>3,4</sup>

There are few studies that have evaluated the efficacy and safety of propofol given by nonanesthesiology staff. Wheeler et al<sup>5</sup> evaluated 91 children who underwent propofol sedation for outpatient procedures administered by a pediatric intensivist. Only a small number of these patients had minor adverse events, suggesting that intensivists can safely administer propofol in an outpatient setting. Guenther et al<sup>6</sup> demonstrated that emergency room physicians can also administer propofol safely for short pediatric procedures.

The increased demand for pediatric sedation may require nonanesthesiology physicians to use propofol sedation. At our institution, a pediatric sedation team has been established, which consists of a board-certified pediatric intensivist who administers propofol as well as a sedation nurse who monitors the patient during and after each procedure. We conducted a retrospective study evaluating outpatient propofol sedation for pediatric procedures to determine this medication's safety when administered by pediatric intensivists.

→ **method of investigation/experiment**

**nature of problem**

**purpose**

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http://jcp.sagepub.com

From the Departments of Pediatrics (RL, DG, SW, DH, LW, JP), Biostatistics (DK), and Internal Medicine (CM), Scott and White Memorial Hospital, Texas A&M University Health Science Center, Temple, Texas.

Author disclosure: None of the authors of this manuscript has financial relationships relevant to this case.

Address correspondence to: John F. Pohl, MD, Section, Pediatric Gastroenterology, Department of Pediatrics, Scott and White Hospital, Texas A&M Health Science Center, 2401 S. 31st, Temple, TX 76708; e-mail: jpohl@swmail.sw.org.

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- Standard structure
  - Abstract
  - Introduction
  - Methods
  - Results
  - Discussion
  - Figures/tables
  - References
  - Acknowledgements

# PIDs + Rich Meta = FAIR<sup>1</sup>

**Meaningful title**

**Full author list**

**How to cite data**

**DOI**

SPARC Dataset Structure

APA Chicago IEEE Bibtex More on Crosscite.org

Larauche, M., Wang, Y., Wang, P.-M., Dubrovsky, G., Lo, Y.-K., Hsiang, I., Dunn, J., Liu, W., Tache, Y., & Mulgeta, M. (2021). Influence of direct colon tissue electrical stimulation on colonic motility in anesthetized male Yucatan minipig (Version 3) [Data set]. Pennisive Discover: <https://doi.org/10.26275/HOWG-TBH>

Copy Citation

**Data description**

**Link to experimental protocol**

**Usage notes**

**Subjects and samples**

**Link to code**

**Description** About Files Gallery

**Study purpose:** Neuromodulation is emerging as an alternative therapy to bowel dysfunctions. However, knowledge of optimal stimulation modalities is lacking. This study aims to map the colonic wall impedance and luminal pressure changes to electrical stimulation of the colon and autonomic nerves in a porcine model.

**Data collection:** Direct electrical stimulation of proximal (pC), transverse (tC) and distal (dC) colon was done using planar flexible multi-electrode array panels. In all instances, pC, tC, and dC wall bioimpedance, and luminal pressure (*manometry*) changes were monitored before, during and after stimulation. The regional colonic motility index was quantified by measuring the area under the curve of the motility index. A regional colonic motility index pressure map was generated.

**Primary conclusion:** Direct colon wall electrical stimulation caused changes in the motility of the proximal, transverse and distal regions motility.

**Protocol Links:**  
<https://www.protocols.io/api/v3/protocols/21857>

**Curator's Notes**

**Experimental Design:** Anesthetized male Yucatan minipigs (n=28) were used and assigned to 2 experimental groups by stimulation organ: (1) direct stimulation - transverse colon, (2) direct stimulation-proximal colon, and (3) direct stimulation-distal colonic regions. The probes were used to measure the effect of electrical stimulation on the serosal side of the colonic tissue at multiple sites, in an acutely anesthetized preparation. The effect of stimulation was used to create a functional map of colonic motor response to local electrical stimulation before, during, and after stimulation, and the

**Completeness:** Complete

**Subjects & Samples:** Young adult male Yucatan pigs (n=28), 27 - 32 kilograms, were used for this study.

**Primary vs derivative:** The primary folder is organized by subject identification and with results in xlsx, pl2, smrx, and s2rx formatted files. A summary of all experiments as an xlsx formatted document

**Important notes:** (1) There is a Docs folder attached to this study. This folder contains metadata for the data set, a list of files, and a summary of this data (pg).

**Code Availability:** The compressed Matlab code (rar) used to extract data from .pl2 format data folder.

Rich metadata aids in **search**, provides **provenance**, **credit** for those who produced it and valuable **context** for understanding the data

**Grant**

**Key words**

**Originating publication**

Description About Files Gallery

Last Updated: June 1, 2021

Corresponding Author: Yvette Tache (ytache@mednet.ucla.edu)

Dataset DOI: <https://doi.org/10.26275/howg-tbh>

NIH Award: OT2D0024899

Tags: autonomic, pelvic, gastrointestinal, electrocortical-therapy, pre-clinical

References: Larauche, M., Wang, Y., Wang, P.-M., Dubrovsky, G., Lo, Y.-K., Hsiang, E., ... Millon, M. (2020). The effect of colonic tissue electrical stimulation and celiac branch of the abdominal vagus nerve neuromodulation on colonic motility in anesthetized pigs. *Neurogastroenterology & Motility*, 32(11). doi:10.1111/nmo.13925



# Lesson: More metadata is better than less metadata

Think about publishing data like you would publish a paper; what information do people need to understand it and use it?



# Accessible



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

“As open as possible, as closed as necessary”



# Structured vs unstructured (meta)data

- Structured (meta)data = Data organized according to a data model—an abstract model that organizes elements of **data** and standardizes how they relate to one another and to properties of the real world entities.
- Unstructured (meta)data = information that either does not have a pre-defined **data model** or is not organized in a pre-defined manner. Unstructured information is typically **text**-heavy, but may contain data such as dates, numbers, and facts as well. This results in irregularities and **ambiguities** that make it difficult to understand using traditional programs as compared to data stored in fielded form in databases or **annotated (semantically tagged)** in documents. ([Wikipedia](#))





## Example

- Unstructured: “subjects comprised male (N=6) and female (N =6) C57BL/6 mice, aged 25 days”
- Structured:
  - Subject Group: 1
  - Organism: mouse
  - Age: 25
  - Age unit: days
  - Strain: C57BL/6
  - Sex: Male
  - Number: 6
  - Subject Group: 2
  - Organism: mouse
  - Age: 25
  - Age unit: days
  - Strain: C57BL/6
  - Sex: Female
  - Number: 6



# Interoperable



- I1. (meta)data use a formal, accessible, shared, and broadly applicable language for knowledge representation.
- I2. (meta)data use vocabularies that follow FAIR principles
- I3. (meta)data include qualified references to other (meta)data



# Typical Research Scenario

Three different laboratories study a rat model of Type 1 diabetes

- Data set 1: Physiological measurements correlated with gene expression in kidney
- Data set 2: Gene expression in peripheral organs
- Data set 3: Gene expression in major organs



# One of these things is not like the other. Maybe.

	Data set 1		Data set 2		Data set 3
Age	15-30 days	Age range	10-20 days	Age	adult
Organism	WF.BBDR-(D4Arb 29-D4Rat44)/Wor	Subject	N8 WF.iddm4	Animal	6
Gene name	Calretinin	Gene	Calb2	Exp	CR
Location	Ct Kidney	Region	Renal cortex	Sample	Cortex

- What are the problems here if you wanted to combine these data?
- What would a computer not “know”
- What could have been done differently?



# FAIR vocabularies

- Have the characteristics of FAIR data:
  - PIDs
  - Metadata
  - Relationships
- Cortex of kidney = [UBERON:0001225](https://www.ebi.ac.uk/ontology/uberon/0001225)
- Can be read by both humans and machines



# PIDs are for more than dataset IDs an articles

- Best practice: Use common identifiers to unambiguously identify the same entity across multiple data sets
  - Reagents and tools: RRIDs
  - Genes: Gene IDs
  - People: ORCIDs
  - Articles: DOIs, PMIDs
  - Concepts: Ontologies/controlled vocabularies



# PID Power!

Attribute	Data set 1	Data set 2	Data set 3
<b>Age</b>	15-30 days	10-20 days	adult
<b>Organism</b> <a href="#">OBI_0100026</a>	WF.BBDR-(D4Arb29-D4Ra t44)/Wor <a href="#">RRID:RGD_1357172</a>	N8 WF.iddm4 <a href="#">RRID:RGD_1357172</a>	6 <a href="#">RRID:RGD_1357172</a>
<b>Gene</b> <a href="#">SO_0000704</a>	Calretinin <a href="#">Gene ID: 117059</a>	Calb2 <a href="#">Gene ID: 117059</a>	CR <a href="#">Gene ID: 117059</a>
<b>Anatomical structure</b> <a href="#">UBERON:0000061</a>	Ctx kidney <a href="#">UBERON:0001225</a>	Renal cortex <a href="#">UBERON:0001225</a>	Cortex <a href="#">UBERON:0000956</a>



# What are RRIDs?

- A persistent identifier for research resources: antibodies, digital tools, cell lines, organisms, plasmids, biosamples
- Required by many journals: Supplied by authors to identify resources in the materials and methods section
- Designed to answer two simple questions:
  - What resources were used in a study
  - Who else has published with this resource?

[RRID:AB\\_1855196](#)



Anti-PYY (ab1)  
antibody produced in chicken,  
affinity isolated antibody  
Catalog Number **GW22771**  
Formerly listed as GenWay Catalog Number 15-288-22771,  
Peptide YY Antibody.

Secondary antibody	Rabbit IgG	No. 21200, Alexa Fluor 488	Molecular Probes
	Sheep IgG	No. A21448, Alexa Fluor 647	Molecular Probes
	Chicken IgG	No. 703-605-155, Alexa Fluor 647	Jackson ImmunoResearch Laboratories (West Grove, PA)
	Guinea pig IgG	No. 11073, Alexa Fluor 488	Molecular probes
	Rabbit IgG	No. 31573, Alexa Fluor 647	Molecular probes

*RRID* Research Resource Identifier (<https://dknet.org/>)

[Fothergill LJ et al. Cell and Tissue Research. 375 \(2\) 359-69, 2019.](#)





# Where can I find RRIDs? Ask dkNET!

ABOUT ▾ RESOURCE REPORTS DISCOVERY PORTAL AUTHENTICATION REPORT HYPOTHESIS CENTER



## dkNET: Connecting Researchers to Resources



### Resource Reports

Is my antibody specific? Who else is using my software tools? Answer these questions and more using Research Resource Identifiers (RRIDs) and Digital Object Identifier (DOIs).

[Tools](#) | [Cell lines](#) | [Antibodies](#) | [Organisms](#) | [Plasmids](#) | [Biosamples](#) | [Protocols](#)



### Authentication Reports & FAIR Data

View resources on how to comply with NIH's new policies on authentication of key biological resources, using our authentication reports, and making data FAIR.

[Authentication reports](#) | [Research data management](#) | [Suggested data repositories](#)

Appyters: Turning Jupyter Notebooks into Data-Driven Web Apps  
Friday, May 28, 2021, 11 am - 12 pm (PDT)  
dkNET Webinar Series

Newsletter  
Find out full events, news, and community



Discovery

Search

Funding



Hypothesis

Analysis

hypoth



## Resource Summary Report

New Search

Previous Search Results

Home / Resource Reports / Organisms / Resource Summary Report



Organism Name

**WFBBDR-(D4Arb29-D4Rat44)/Wor**

RRID:RGD\_1357172



Organism Information

URL: [http://rgd.mcw.edu/tools/strains/strains\\_view.cgi?id=1357172](http://rgd.mcw.edu/tools/strains/strains_view.cgi?id=1357172)

Proper Citation: (RGD Cat# 1357172;RRID:RGD\_1357172)

Description: Rattus norvegicus with name WFBBDR-(D4Arb29-D4Rat44)/Wor from RGD.

Database: Rat Genome Database Strain List RGD

Notes: This congenic was generated by the marker-assisted protocol where a segment of BBDR/Wor is transferred to WF background and the animals were screened using microsatellite markers. U Medical School, Worcester, MA

References: PMID:12401717

Organism Name: WFBBDR-(D4Arb29-D4Rat44)/Wor

Database Abbreviation: RGD

Species: Rat

Phenotype: type 1 diabetes mellitus, increased susceptibility to autoimmune diabetes

Availability: live

Catalog Number: 1357172

Background: congenic strain

Collapse



Usage and Citation Metrics

We have not found any literature mentions for this resource.



Collaborator Network

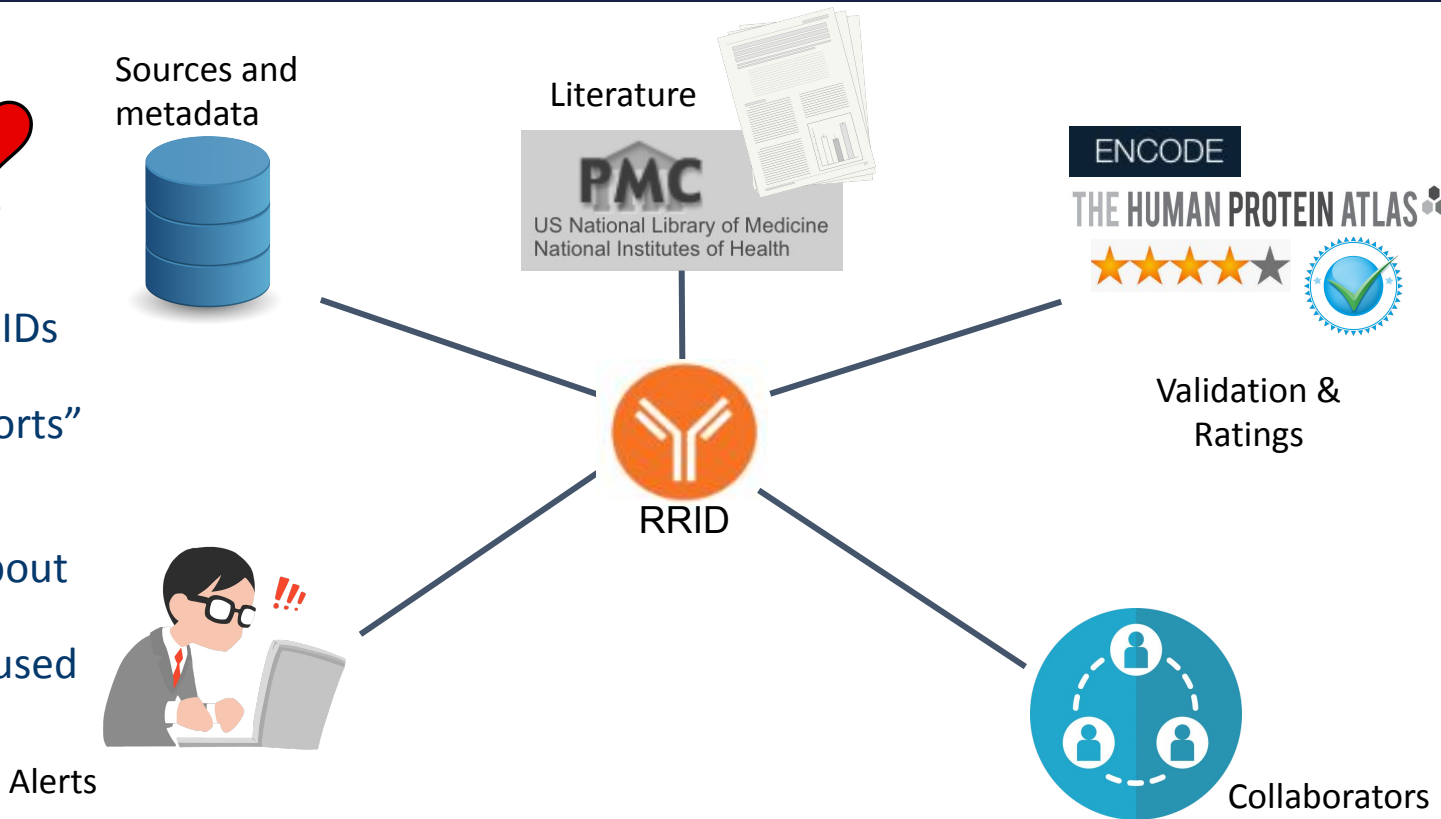
A list of researchers who have used the resource and an author search tool that have literature mentions

<https://dknet.org/rin/rrids>



# PIDs make it easier to combine data across sources

- Computers ❤️ PIDs like RRIDs
- dkNET uses RRIDs to create “Resource reports” which provide additional information about how these resources are used





# Lesson: Use RRIDs and other IDs in your lab notebooks, spreadsheets, data dictionaries and papers!

Helps you and others (including computers) identify the tools that you use



# Re-usable



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



# Lesson: Think about where your data will end up in the beginning

Best practice: Submit your data to repository specialized for your type of data or your domain

..if there isn't one, then there are also general purpose repositories available



# Where Can I Deposit My Data?

- List of DK relevant repositories, recommended by NLM and various journals

- Created in conjunction with NIDDK

- Coming soon: FAIR data wizard

- FAIR Standards
- Clinical Repositories Information
- Data maintenance
- Data size limit and cost
- Dynamic database

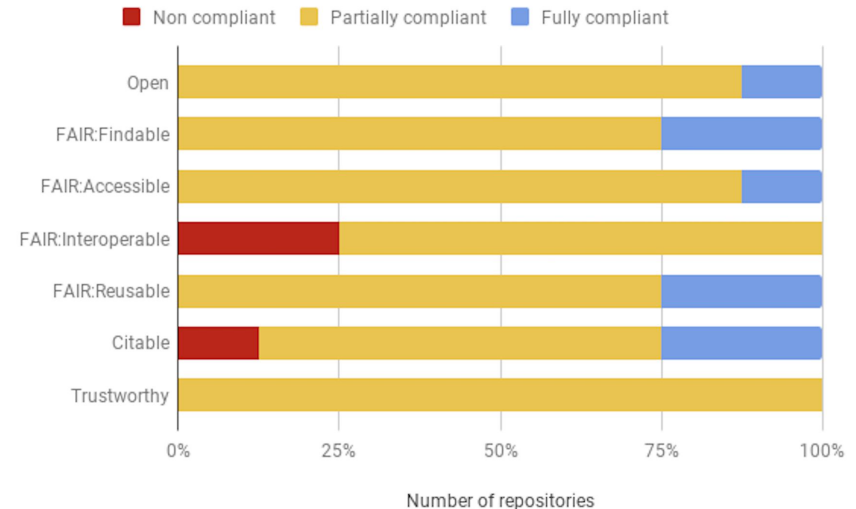
Repository Name	RRID	Description	Type of Data	Recommended By
ArrayExpress	RRID:SCR_002964	International functional genomics data collection generated from microarray or next-generation sequencing (NGS) platforms. Repository of functional genomics data supporting publications. Prov ...[more]	Microarray; next-generation sequencing (NGS)	NIDDK
Biological General Repository for Interaction Datasets (BioGRID)	RRID:SCR_007393	Curated protein-protein and genetic interaction repository of raw protein and genetic interactions from major model organism species, with data compiled through comprehensive curation efforts.	Molecular interaction data	NLM, NIDDK
Database of Interacting Proteins (DIP)	RRID:SCR_003167	Database to catalog experimentally determined interactions between proteins combining information from a variety of sources to create a single, consistent set of protein-protein interactions ...[more]	Protein interaction data	NLM, NIDDK
Gene Expression Omnibus (GEO)	RRID:SCR_005012	Functional genomics data repository supporting MIAME-compliant data submissions. Includes microarray-based	Microarray; next-generation sequencing	NLM, NIDDK

Repository Name	RRID	Description	Type of Data	Recommended By
Dataverse Network Project	RRID:SCR_001997	Project portal for publishing, citing, sharing and discovering research data. Software, protocols, and community connections for creating research data professional ...[more]	All research data	NIDDK



# Data repositories

- “Publishing platform” for data
- Run by governments, researchers, commercial entities, non-profit entities, universities
- Hundreds of them across biomedicine
- Four major types:
  - Data type specific
  - Domain specific
  - Generalist
  - Institutional
- A stated goal of NIH’s strategic data plan is to modernize biomedical repositories
- Ensure persistence, searchability, FAIRness etc



Murphy et al., PLoS One, in press. Evaluation of biomedical data repositories against Open, FAIR, Citable and Trustworthy criteria

Preprint:

<https://www.biorxiv.org/content/10.1101/2021.01.19.427362v2>




# What standards should I use?

- Repositories often enforce specific standards for metadata and data
- Thinking about where your data will end up before you start your experiments will help you determine how to collect, annotate and organize your (meta)data
- [Fairsharing.org](http://Fairsharing.org) maintains a database of standards and policies across biomedicine

Repository Name	RRID	Description	Type of Data	Submitting Data	Accessing Data	Guidelines/Standards	Recommended by
<a href="#">ArrayExpress</a>	<a href="#">RRID:SCR_002964</a>	Archive of Functional Genomics Data stores data from high-throughput functional genomics experiments, and provides these data for reuse to the research community.	<a href="#">Microarray; next-generation sequencing (NGS)</a>	<a href="#">How to submit to ArrayExpress</a>	<a href="#">How to access ArrayExpress Data</a>	<a href="#">MIAME standard for microarray data.</a>	NIDDK; Nature Scientific Data; PLOS; Science
<a href="#">Biological General Repository for Interaction Datasets (BioGRID)</a>	<a href="#">RRID:SCR_007393</a>	BioGRID is a curated biological database of protein-protein interactions, genetic interactions, chemical interactions, and post-translational modifications from major model organism species. All interaction data are freely provided through our search index and available via download in a wide variety of standardized formats.	<a href="#">Molecular interaction data</a>	<a href="#">How to submit data to BioGRID</a>	<a href="#">How to access BioGRID data</a>	Molecular interaction data should be deposited with a member of the International Molecular Exchange Consortium (IMEX), following the <a href="#">MIMIX recommendations</a> .	NIDDK; Nature Scientific Data; PLOS
<a href="#">database of Genotypes and Phenotypes (dbGaP)</a>	<a href="#">RRID:SCR_002709</a>	The database of Genotypes and Phenotypes (dbGaP) was developed to archive and distribute the data and results from studies that have investigated the interaction of genotype and phenotype in Humans.	<a href="#">Genotyping and phenotyping information in human subject</a>				
<a href="#">Database of Interacting Proteins (DIP)</a>	<a href="#">RRID:SCR_003167</a>	DIP catalogs experimentally determined interactions between proteins. It combines information from a variety of sources to create a single, consistent set of protein-protein interactions.	<a href="#">Protein interaction data</a>				
<a href="#">Gene Expression Omnibus (GEO)</a>	<a href="#">RRID:SCR_007303</a>	Gene Expression Omnibus is a public functional genomics data repository supporting MIAME-compliant submissions of array- and sequence-based data. Tools are provided to help users query and download experiments and curated gene expression profiles.	<a href="#">Microarray; next-generation sequencing (N)</a>				
<a href="#">GenomeRNAi</a>	<a href="#">RRID:SCR_013088</a>	A database containing phenotypes from RNA	<a href="#">RNAi screening</a>				



Home About Us Mission Projects Conferences


### MIAME

#### Minimum Information About a Microarray Experiment

MIAME describes the **Minimum Information About a Microarray Experiment** that is needed to enable the interpretation of the results of the experiment unambiguously and potentially to reproduce the experiment. [Brazma et al., (2001), *Nature Genetics*]

The six most critical elements contributing towards MIAME are:

1. The raw data for each hybridisation (e.g., CEL or GPR files)
2. The final processed (normalised) data for the set of hybridisations in the experiment (study) (e.g., the gene expression data matrix used to draw the conclusions from the study)
3. The essential sample annotation including experimental factors and their values (e.g., compound and dose in a dose response experiment)
4. The experimental design including sample data relationships (e.g., which raw data file relates to which sample, which hybridisations are technical, which are biological replicates)
5. Sufficient annotation of the array (e.g. gene identifiers, genomic coordinates, probe oligonucleotide sequences or reference commercial array catalog number)
6. The essential laboratory and data processing protocols (e.g., what normalisation method has been used to obtain the final processed data)



standards, databases, policies

Standards Databases Policies Collections Add/Claim Content Stats Log in or Register

### Databases

Contribute by adding a database Any problems? Please tell us!

A catalogue of databases, described according to the [BioDBcore guidelines](#), along with the standards used within them; partly compiled with the support of Oxford University Press ([NAR Database Issue](#) and [DATABASE Journal](#)).

Search Databases  Search Reset Advanced





# Data Management and Sharing Plan

- Creating a good data management and sharing plan allows you to:
  - Comply with NIH mandates
  - Ensure that you allocate enough resources for preparing and sharing your data
  - Ensure that you collect your data in a FAIR manner
  - Easily share data with yourself, future you, your colleagues and the scientific community
- dkNET provides links to resources that can help

<https://dknet.org/rin/rigor-reproducibility-about>

The image shows two overlapping web pages. The top page is the DMPTool homepage, which includes the logo 'DMPTool' with the tagline 'Build your Data Management Plan'. It features a 'Welcome to the DMPTool' section and a grid of navigation buttons for various services like 'Research data in context', 'File formats & transformation', and 'Protecting sensitive data'. The bottom page is the MIT Libraries 'Data management' page, which has a navigation bar with 'Home', 'Services', 'Make a plan', 'Store your data', and 'Share your data'. The main content area on this page includes the heading 'Data management' and a list of links under the 'Make a plan' section: 'Why manage your data?', 'Write a data management plan', 'Other guides to data management', and 'Meet funder requirements'.



# R1.2: Provenance

- Metadata supplied with a dataset should include information about where and how the data were obtained
- Detailed experimental protocols
- Associated code
- Associated publications
- Standard authoring metadata
  - Also makes the data citable!

Effects of a fecal transplant from anxious donors on rehabilitative training, microbiota composition, systemic inflammation and behaviour following a unilateral cervical spinal contusion (C5, 125kdyn) in female Lewis rats

DOI:10.34945/F5XW2P

#### DATASET CITATION

Schmidt E. K. A., Raposo P. J. F., Madsen K. L., Fenrich K. K., Kabarchuk G., Fouad K. (2021) Effects of a fecal transplant from anxious donors on rehabilitative training, microbiota composition, systemic inflammation and behaviour following a unilateral cervical spinal contusion (C5, 125kdyn) in female Lewis rats. ODC-SCI:578 <http://doi.org/10.34945/F5XW2P>

#### ABSTRACT

**STUDY PURPOSE:** The purpose of this study was to determine whether optimal donor selection would influence the outcome of a fecal microbiota transplant (FMT) and the efficacy of rehabilitative training after a unilateral C5 spinal contusion (infinite horizons impactor, 125 kdyn) in female Lewis rats. Uninjured, genetically identical FMT donors (n=10) were selected as rats who displayed naturally reduced baseline activity levels and increased anxiety-like behaviour. Experimental groups consisted of a group that was gavaged with the FMT solution for 3 days following SCI (SCI+FMT, n=15) and a group that received a vehicle control solution (SCI+Vehicle, n=15).

**DATA COLLECTED:** Data collected for this study include:

16S rRNA sequencing of fecal matter collected pre-injury, on the day of injury, then 3, 7, 14 and 56 days post injury. This data includes the bacteria bacteroidetes, firmicutes, proteobacteria, cyanobacteria, lactobacillus, as well as the alpha diversity.

Rehabilitation training on a single pellet reaching task. This includes the number of attempts and success rate at baseline (pre-injury) and throughout 8 weeks of rehab training after SCI (10 min per day, 5 days per week, analyzed once per week).

Von Frey test at 1 and 9 weeks post SCI (normalized to baseline values).

The percentage of ipsilesional paw placements in the cylinder test (9 weeks post SCI).

The total distance travelled in the open field (9 weeks post SCI).

The total distance travelled and the percentage of time spent in the open arms of an elevated plus maze (9 weeks post SCI).

The time spent in the light component of the light-dark box test (9 weeks post SCI).

The percentage of sucrose solution consumed over 2 hours (9 weeks post SCI).

The time spent interacting in the social interaction test (9 weeks post SCI).

Lesion analysis.

IBA1 density immediately caudal to the lesion, at and immediately rostral to the lesion.

Plasma analytes were measured at baseline, 3, 21 and 77 days post SCI.

FITC dextran assay for intestinal permeability (7 days post SCI displayed as a fold change from baseline, this was a separate experiment which also includes lesion analysis and plasma analytes at 7 days post SCI).

**DATA USAGE NOTES:** The data presented in this dataset show that a fecal transplant from anxious donors (with decreased levels of Lactobacillus in their stool) does not prevent SCI-induced dysbiosis. This fecal transplant had some long-term side effects on systemic and local inflammation and also increased anxiety-like behaviour in the recipient rats. Overall, this study shows that optimal donor selection is critical for successful FMT treatment following SCI.

#### KEYWORDS

Spinal Cord Injury; Inflammation; Anxiety; Fecal transplant

#### PROVENANCE / ORIGINATING PUBLICATIONS

#### DATASET INFO

Contact: Fouad Karim (kfouad@ualberta.ca)

Lab: Karim Fouad

ODC-SCI Accession:578

Records in Dataset: 431

Fields per Record: 134

Files: 2

#### LICENSE

Creative Commons Attribution License (CC-BY 4.0)

#### FUNDING AND ACKNOWLEDGEMENTS

#### CONTRIBUTORS

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Kabarchuk, Gillian

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Fouad, Karim [ORCID:0000-0003-3654-7852]

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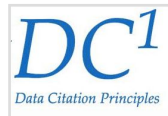
[DOI:10.34945/F5XW2P](https://doi.org/10.34945/F5XW2P)



# Data as a Research Product

*Sound, reproducible scholarship rests upon a foundation of robust, accessible data. For this to be so in practice as well as theory, data must be accorded due importance in the practice of scholarship and in the enduring scholarly record..."*

1. Data should be considered *legitimate, citable products of research*. Data citations should be accorded the same importance in the scholarly record as citations of other research objects, such as publications.
2. Data citations should facilitate giving scholarly credit and normative and legal attribution to all contributors to the data, recognizing that a single style or mechanism of attribution may not be applicable to all data.
3. In scholarly literature, whenever and wherever a claim relies upon data, the corresponding data should be cited.



# A data citation looks like a regular citation

DOI



[DOI:10.34945/F5XW2P](https://doi.org/10.34945/F5XW2P)

Full citation



## DATASET CITATION

Schmidt E. K. A., Raposo P. J. F., Madsen K. L., Fenrich K. K., Kabarchuk G., Fouad K. (2021) Effects of a fecal transplant from anxious donors on rehabilitative training, microbiota composition, systemic inflammation and behaviour following a unilateral cervical spinal contusion (C5, 125kdyn) in female Lewis rats. ODC-SCI:578 <http://doi.org/10.34945/F5XW2P>



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[DOI:10.34945/F5XW2P](https://doi.org/10.34945/F5XW2P)


# Proper data citation = data citation metrics

2 datasets found







Cervical (C5), unilateral spinal cord injury with diverse injury...  
scicrunch.org  
Updated Apr 25, 2019



Derivation of Multivariate Syndromic Outcome Metrics for...  
figshare.com  
avi, tiff  
Updated Jan 18, 2016



Not seeing a result you expected?  
[Learn](#) how you can add new datasets to our index.



Cervical (C5), unilateral spinal cord injury with diverse injury modalities, multiple behavioral outcomes, and histopathology  
Related Article  
[Explore at scicrunch.org](#)  
2 scholarly articles cite this dataset ([View in Google Scholar](#))  
**Unique Identifier**  
<https://doi.org/10.7295/W9T72FMZ>  
**Dataset updated** Apr 25, 2019  
**Dataset provided by**  
[University of California San Francisco Parnassus Campus](#)  
Burke Rehabilitation Institute, Weill Cornell Medical College  
**Authors**  
Adam R. Ferguson; Karen-Amanda Irvine; John C. Gensel; Amity Lin; Johnathan Ly; Mark R. Segal; Rajiv R. Ratan; Jacqueline C. Bresnahan; Michael S. Beattie

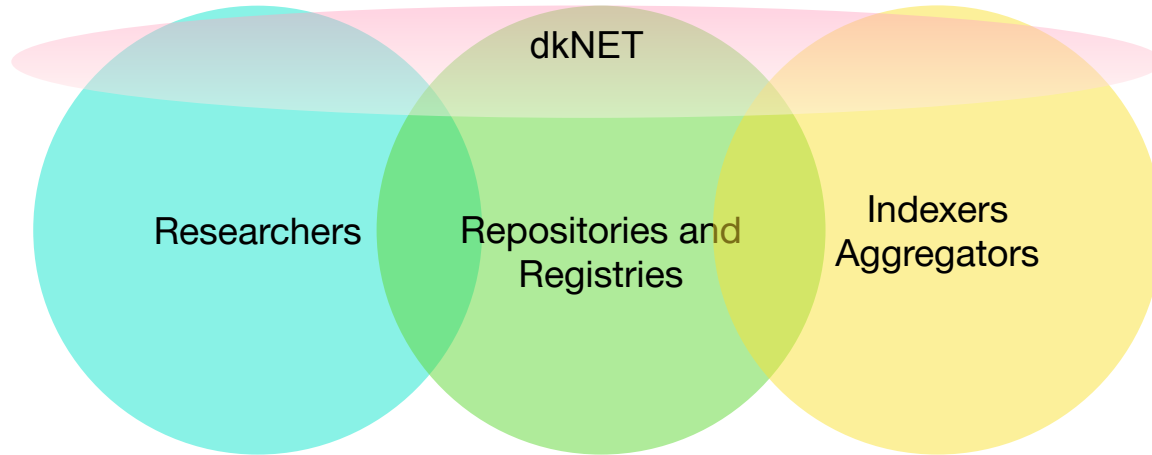


# OVERWHELMED?

When you're waist-deep in tribbles, it's a bit difficult to remember that your original objective was to guard the quadrotriticale.



# FAIR Partnership



- **Good data management**
- **Rich metadata**
- **Prepare to share**
- Open formats
- **Adopt/align to standards**
- **Submit to repository**

- Persistent identifier
- Machine based access
- Clear license
- Support for open, domain specific standards
- Machine readable metadata
- Future friendly formats
- Persistent metadata
- Bidirectional links
- Data citation

- Index
- Effective Search
- Persistent metadata



# Having trouble? Ask dkNET



## FAIR Data Resources

dkNET now offers information on how to manage data in compliance with the FAIR Data Principles.



- Find information for best practices in managing research data
- List of community approved repositories



- Request a data repository recommendation
- Tool for creating a FAIR Data Plan (Coming soon)

Coming soon: The FAIR data wizard!





# You are here



Changing the culture of biomedicine one step at a time...

