



GREI Generalist Repository
Ecosystem Initiative

Meet the GREI Repositories

October 12, 2022
GREI Collaborative Webinar Series





GREI

Generalist Repository
Ecosystem Initiative



OSF



figshare



DRYAD



MENDELEY DATA



Vivli

CENTER FOR GLOBAL CLINICAL RESEARCH DATA

zenodo



Meet our Speakers



*Ishwar
Chandramouliswaran
GREI Program
Director, NIH ODSS*



*Sonia Barbosa,
Manager of
Curation, Harvard
Dataverse*



*Jennifer Gibson,
Executive Director,
Dryad*



*Ana Van Gulick,
PhD, Government
and Funder Lead,
Head of Data
Review, Figshare*



Meet our Speakers



*Luca Belletti,
Product Manager,
Digital Commons
Data/Mendeley Data*



*Eric Olson,
Product Manager,
Center for Open
Science (Open
Science
Framework)*



*Ida Sim, MD, PhD,
Professor of
Medicine and
Computational
Precision Health,
UCSF, Vivli*



*Sara Gonzales,
Senior Data
Librarian,
Northwestern
University,
Zenodo*



Generalist Repository Ecosystem Initiative



GREI Program – Include GR’s in the NIH Data Ecosystem via the Concept of “Co-opetition”

work together on Common Capabilities & Best-practices



Expected Outcomes & Impact



Implement consistent capabilities (NOT-OD-21-016)



Create better access to & discovery of NIH funded data



Conduct outreach & train on FAIR data practices



Engage the research community



Make data sharing easier



Improve discoverability



Increase reproducibility of research



Encourage secondary use of data



MENDELEY DATA



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GREI Fact Sheet

<https://doi.org/10.6084/m9.figshare.21318270>

NIH Generalist Repository Ecosystem Initiative

The mission of GREI is to establish a common set of capabilities, services, metrics, and social infrastructure; raise general awareness and facilitate researchers to adopt FAIR principles to better share and reuse data.

This initiative will further enhance the biomedical data ecosystem and help researchers find and share data from NIH-funded studies in generalist repositories.

Goals of the Generalist Repository Ecosystem Initiative



Objectives



- Catalog Use Cases Supported
- Implement Open Metrics
- Prepare Training Materials
- Conduct Outreach
- Commit to 'Coopetition'
- Implement Best Practices for Data Repositories
- Support Discovery of NIH Funded Data
- Adopt Consistent Metadata Models
- Facilitate QA/QC
- Connect Digital Objects

Participating Repositories

The **Dataverse**[®] Project

 **DRYAD**

 **figshare**

 **MENDELEY DATA**

 **OSF**

 **Vivli**
CENTER FOR GENERAL SCIENTIFIC RESEARCH DATA

 **zenodo**

Contact

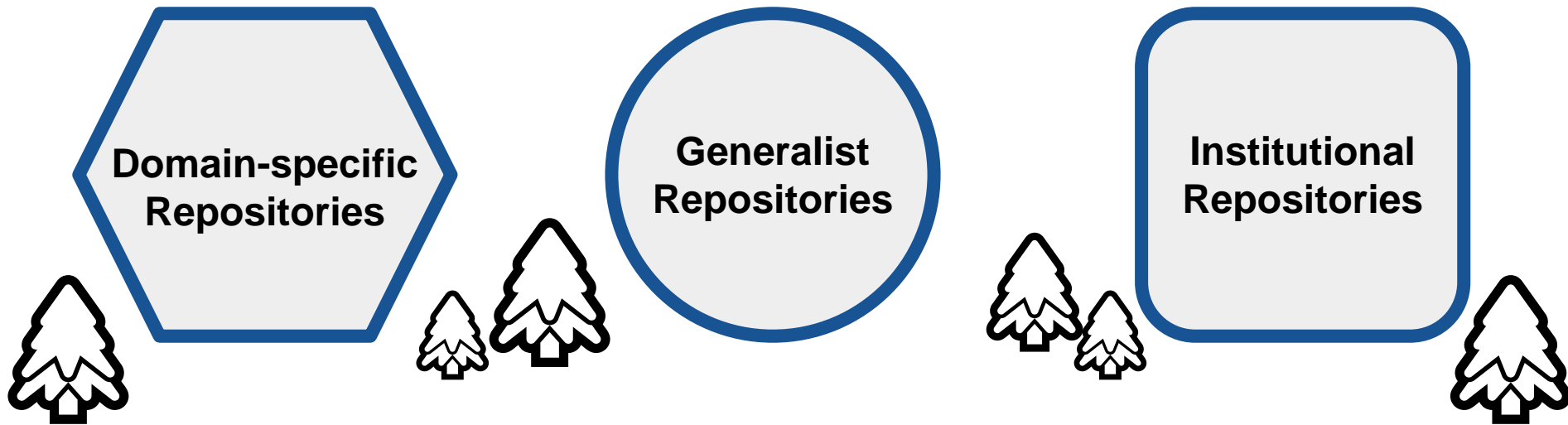
GREI@nih.gov • <https://bit.ly/3en3rRy>



Generalist Repository Features



NIH Research Data Ecosystem



Desirable Characteristics of Data Repositories

When choosing a repository to manage and share data resulting from Federally funded research, here are some desirable characteristics to look for:

- **Unique Persistent Identifiers**
- **Long-Term Sustainability**
- **Metadata**
- **Curation and Quality Assurance**
- **Free and Easy Access**
- **Broad and Measured Reuse**
- **Clear User Guidance**
- **Security and Integrity**
- **Confidentiality**
- **Common Format**
- **Provenance**
- **Retention Policy**

[Guidance set forth by NIH](#)

And by [The National Science and Technology Council](#),
cited in OSTP guidance



Generalist Repository Comparison Chart

doi: 10.5281/zenodo.3946720

This chart is designed to assist researchers in finding a generalist repository should no domain repository be available to preserve their research data. Generalist repositories accept data regardless of data type, format, content, or disciplinary focus. For this chart, we included a repository available to all researchers specific to clinical trials (Vivli) to bring awareness to those in this field.

<https://fairsharing.org/collection/GeneralRepositoryComparison>

TOPIC	HARVARD DATAVERSE	DRYAD	FIGSHARE	MENDELEY DATA	OSF	VIVLI	ZENODO
Brief Description	Harvard Dataverse is a free data repository open to all researchers from any discipline, both inside and outside of the Harvard community, where you can share, archive, cite, access, and explore research data.	Open-source, community-led data curation, publishing, and preservation platform for CCO publicly available research data Dryad is an independent non-profit that works directly with: <ul style="list-style-type: none"> researchers to publish datasets utilizing best practices for discovery and reuse publishers to support the integration of data availability statements and data citations into their workflows institutions to enable scalable campus support for research data management best practices at low cost 	A free, open access, data repository where users can make all outputs of their research available in a discoverable, reusable, and citable manner. Users can upload files of any type and are able to share diverse research products including datasets, code, multimedia files, workflows, posters, presentations, and more. With discoverable metadata supporting FAIR principles, file visualizations, and integrations, researchers can make their work more impactful and move research further faster.	Mendeley Data is a free repository specialized for research data. Search more than 20+ million datasets indexed from 1000s of data repositories and collect and share datasets with the research community following the FAIR data principles.	OSF is a free and open source project management tool that supports researchers throughout their entire project lifecycle in open science best practices.	Vivli is an independent, non-profit organization that has developed a global data-sharing and analytics platform. Our focus is on sharing individual participant-level data from completed clinical trials to serve the international research community.	Powering Open Science, built on Open Source. Built by researchers for researchers. Run from the CERN data centre, whose purpose is long term preservation for the High Energy Physics discipline, one of the largest scientific datasets in the world
Size limits	No byte size limit per dataset. Harvard Dataverse currently sets a file size limit of 2.5GB.	300GB/dataset	Soft limit of 20GB/file for free accounts. System limit of 5000GB/file. Unlimited storage of public data but 20GB storage for private data for free accounts. Email info@figshare.com to have upload and storage limits raised.	10GB per dataset	Projects currently have not storage limit. There is a 5GB/file upload limit for native OSF Storage. There is no limit imposed by OSF for the amount of storage used across add-ons connected to a given project.	If more than 10GB per study data, reach out to us	50GB per dataset, contact us via https://zenodo.org/support for higher limits
Storage space per researcher	1 TB per researcher	No limit	No limit	No limit	No limit	No limit	No limit
Persistent, Unique Identifier Support	DOI, Handle	DOI	DOI	DOI	DOI	DOI	DOI

Common features and unique features

Common:
 Core Metadata
 Persistent Identifiers
 Discoverable
 Flexibility
 Open access, FAIR
 Metrics

Unique:
 Output types
 Storage, size limits
 Licenses
 Review
 Controlled Access
 Visualization
 Costs

<https://doi.org/10.5281/zenodo.3946719>

(Updated version 2!)





A few questions for you: A Zoom Poll



1. What is your role in NIH Data Sharing?

1. Have you published data in a repository previously?

1. What do you feel are the biggest barriers to data sharing for you?



Let's meet our GREI repositories



Dataverse

The
Dataverse[®]
Project 



Dataverse World-wide Community (dataverse.org)

- Open-source software
- Two bi-weekly Community calls
- Annual Community meetings (since 2015)
- Global Dataverse Community Consortium (GDCC)

DATVERSE REPOSITORIES - A WORLD VIEW



Open source research data repository software



Researchers

Enjoy full control over your data. Receive *web visibility, academic credit, and increased citation counts*. A personal Dataverse collection is easy to set up, allows you to display your data on your personal website, can be branded uniquely as your research program, makes your data more discoverable to the research community, and satisfies data management plans. *Want to set up your personal Dataverse collection?*



Journals

Seamlessly manage the submission, review, and publication of data associated with published articles. Establish an *unbreakable link* between *articles in your journal and associated data*. Participate in the open data movement by using a Dataverse collection as part of your journal data policy or list of repository recommendations. *Want to find out more about journal Dataverse collections?*



Institutions

Establish a research data management solution for your community. Federate with a growing list of Dataverse repositories worldwide for increased discoverability of your community's data. Participate in the drive to set norms for sharing, preserving, citing, exploring, and analyzing research data. *Want to install a Dataverse repository?*



Developers

Participate in a vibrant and growing community that is helping to drive the norms for sharing, preserving, citing, exploring, and analyzing research data. Contribute code extensions, documentation, testing, and/or standards. *Integrate research analysis, visualization and exploration tools, or other research and data archival systems with the Dataverse Project. Want to contribute?*



Current Features



Dataverse Collections

- Own administration
- Own branding (and can be embedded anywhere)

Datasets

- Citation
- Metadata
- Versioning
- Private URL
- Custom Terms/Multiple License/Permissions
- Guestbooks
- Publishing Workflows



Files

- Citation
- Ingest
- Preview/Explore
- Metadata
- Versioning
- Permissions/Embargo/Restrictions



Log In

Log in or sign up with your institutional account — more information about account creation. Leaving your institution? Please contact Harvard Dataverse Support for assistance.

Your Institution



Harvard University

Please select...

Continue

Allow me to type the name of my institution

Other options

Username/Email

GitHub

Google

ORCID

Sign up for a Dataverse account.



Brain Genomics Superstruct Project (GSP) Dataverse (Harvard University)
GSP

General Information
Theme + Widgets
Permissions
Groups
Dataset Templates
Dataset Guestbooks
Featured Dataverses





Current Features



Installations



Collections



Datasets



Files

NIH OTA Adds...



Datasets

- New Workflows for **Large Data Support** ([Globus](#), [demo](#))
- **Remote** Data Support ([TRSA](#))
- Flexible **Biomedical Metadata** Support through External Vocabularies and Data Dictionaries (UMLS, CEDAR, MeSH)



Files

- Additional Metadata for **Code files** ([Codemeta](#))
 - Support for **Replication Packages** through Workflows and Containers
 - **Sensitive Data Support** through Differential Privacy ([OpenDP](#), [DataTags](#), [PSIprivacy](#))
 - Encryption
- Usage Metrics
 - UX/UI Enhancements
 - New APIs
 - Interoperability
 - Metadata Harvesting
 - Curation Services
 - [Training and Outreach](#)





Harvard Dataverse



Current Features



Dataverse Collections

- Own administration
- Own branding (and can be embedded anywhere)

Datasets

- Citation
- Metadata
- Versioning
- Private URL/*Anonymous Peer Review
- Custom Terms/*Multiple Licenses/Permissions
- Guestbooks
- Publishing Workflows



Files

- Citation
- Ingest
- Preview/Explore
- Metadata/Provenance
- Versioning
- Permissions/Embargo/Restrictions



Dataset Metrics
13,076 Downloads

version	downloads	year	total	used	size	version
1	2	USA	1816	0	0	3.2
2	2	USA	1817	0	0	3.2
3	2	USA	1818	0	0	3.2
4	2	USA	1819	0	0	3.2
5	2	USA	1820	0	0	3.2
6	2	USA	1821	0	0	3.2
7	2	USA	1822	0	0	3.2
8	2	USA	1823	0	0	3.2
9	2	USA	1824	0	0	3.2
10	2	USA	1825	0	0	3.2
11	2	USA	1826	0	0	3.2
12	2	USA	1827	0	0	3.2
13	2	USA	1828	0	0	3.2
14	2	USA	1829	0	0	3.2
15	2	USA	1830	0	0	3.2
16	2	USA	1831	1	0	3.2
17	2	USA	1832	1	0	3.2
18	2	USA	1833	1	0	3.2
19	2	USA	1834	1	0	3.2
20	2	USA	1835	1	0	3.2
21	2	USA	1836	1	0	3.2
22	2	USA	1837	1	0	3.2
23	2	USA	1838	1	0	3.2
24	2	USA	1839	1	0	3.2
25	2	USA	1840	1	0	3.2
26	2	USA	1841	1	0	3.2
27	2	USA	1842	1	0	3.2





Harvard Dataverse



- Support for FAIR Data Principles
- Data citation for datasets and files
- OAI-PMH (Harvesting)
- APIs for interoperability and custom integrations
- API client libraries
- DataCite integration
- Login via Shibboleth
- Login via ORCID, Google, GitHub, or Microsoft
- Login via OpenID Connect (OIDC)
- Internationalization
- Versioning
- Restricted files
- Embargo
- Custom licenses
- Custom terms of use
- Publishing workflow support
- File hierarchy
- File previews
- Preview and analysis of tabular files
- Usage statistics and metrics
- Guestbook
- Fixity checks for files
- File download in R and TSV format
- Faceted search
- Customization of collections
- Private URL
- Widgets
- Notifications
- [Schema.org](https://schema.org) JSON-LD
- External tools
- External vocabulary
- Dropbox integration
- GitHub integration
- Integration with Jupyter notebooks
- User management
- Curation status labels
- Branding
- Backend storage on S3 or Swift
- Direct upload and download for S3
- Export data in BagIt format
- Post-publication automation (workflows)
- Pull header metadata from Astronomy (FITS) files
- Provenance
- Support for rsync
- Auxiliary files for data files

Dryad





An open data publishing platform & community committed to the open availability and routine re-use of all research data



- Not for profit
- Started by researchers
- Connected to the publishing workflow
- Supported by institutions



DRYAD

Not-for-profit and online since
2008. Learn more: datadryad.org



- Serving all research domains
- Leader in research data
- Interconnected
- Fully curated



DRYAD

Not-for-profit and online since
2008. Learn more: datadryad.org



Dryad is best for data that:

1. Doesn't have a home in a specialist repo
2. Can be released under a CCo license
3. Is intended for reuse



DRYAD

Not-for-profit and online since
2008. Learn more: datadryad.org



Focus on reuse

<https://bit.ly/dryad-reuse>



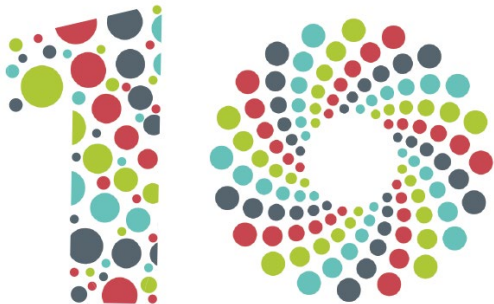
DRYAD

Not-for-profit and online since
2008. Learn more: datadryad.org



Figshare





10 years of figshare

4 million+ research outputs

500,000+ users

100s of TBs of data stored

100,000+ citations

80+ Research Organizations

A screenshot of the Figshare website homepage. The background is a vibrant, multi-colored molecular structure. At the top, the Figshare logo is on the left, followed by a search bar with the text "Search on figshare..." and a magnifying glass icon. To the right of the search bar are links for "Browse", "Log in", and "Sign up". In the center, a white box contains the text "store, share, discover research" where "share" is green, "discover" is purple, and "research" is white on a dark purple background. Below this, it says "get more citations for all of the outputs of your academic research over 30,000 citations of figshare content to date" and "ALSO FOR INSTITUTIONS & PUBLISHERS". At the bottom of the white box, there is a quote: "figshare wants to open scientific data to the world" followed by the WIRED logo. Below the white box, there is a caption: "The background figure: Comparative model of novel coronavirus 2019-nCoV... by Christian Gruber in Virology".

simplify your research workflow

[Upload](#) > [Manage](#) > [Share](#) > [Publish](#)













a freely available generalist repository for all research disciplines and outputs

figshare.com


Flexibility

-  Share any research output or file type
-  Files up to 20GB, Support for big datasets
-  Preview files in the browser
-  Collections





Researcher Workflows

-  Open API and FTP
-  GitHub, GitLab, BitBucket Integrations
-  Collaborative spaces
-  Restricted Access





Persistent Metadata

-  Unique DOI for each output, reservable
-  ORCID integration
-  Link to publications
-  Link Funding via Dimensions

Open Access

-  Open Access to all public files and metadata
-  CC0 and CC-BY Licenses
-  Discoverable across search engines, indexes
-  FAIR commitment






Impact

-  Public Author Profile
-  Views, Downloads, Citations, Altmetrics
-  Citations from full text literature
-  Faceted Search



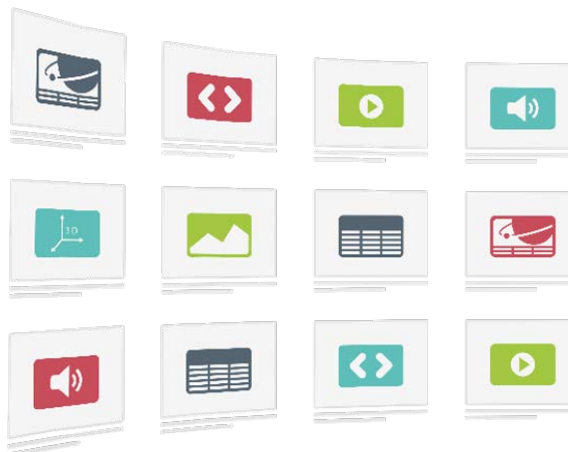
a FAIR repository for big data

plus.figshare.com

-  Publish datasets over 20GB+ to 5TB or more
-  File sizes up to 5TB
-  Expert deposit support
-  Dataset review
-  One-time data publishing charge



Many item types, any file, group as items & collections



Suppl. Movies for Scherer et al 'R_h mechanisms for host-mediated spread

Meets posted on 31.01.2020, 12:32 by Roman Seman, Joshua Kalley

Suppl. Movies for Scherer et al 'R_h mechanisms for host-mediated spread'

Movie S1. As described in Fig. 2A, larvae were infected with yeast-*lacZ* phagocytes and dissemination of *Y. enterocolitica* from the surface of 3rd instar

Index	Group	Size	Download	View	Download	Embed	Share	Weight	Size	Year	Strength	File	Download Progress	Space SP	Quota SP	Disk Space
1	NULP	1	26.1	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
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3	NULP	1	42.39	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
4	NULP	1	4.49	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
5	NULP	1	34.62	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
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48	NULP	1	42.29	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
49	NULP	1	32.39	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
50	NULP	1	27.7	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%

Download all (372.74 MB)

Index	Group	Size	Download	View	Download	Embed	Share	Weight	Size	Year	Strength	File	Download Progress	Space SP	Quota SP	Disk Space
1	NULP	1	10.44	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%
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3	NULP	1	32.58	0	0	1	0	0	20	20	2.0	1	100%	100%	100%	100%

SBCapSeq Calcul... .xlsx (10.44 KB)

SBCapSeq DT... .xlsx (10.4 KB)

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SBCapSeq Protocol manuscript files for 'Quantifying tumor heterogeneity, clonal dynamics, and cancer driver gene evolution from Sleeping Beauty transposon mutagenesis models using SBCapSeq'

Dataset posted on 26.12.2019, 01:59 by Michael Mann, Karen M. Mann, Ana M. Corbano-Sandoval, Liliana Guzman-Rojas, Justin Newberg, Amanda L. Meshey, Devin J. Jones, Felipe Amaya-Manzaneres, Neal G. Copeland, Nancy A. Jenkins

330 views 10 downloads 0 citations

item
single file, single DOI, item-level metadata

item
multiple files, single DOI, item-level metadata

collection
single DOI, collection-level metadata
contains multiple items, each with their own DOI and item-level metadata



Describe your work

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Select categories

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Keyword(s)

Add keywords for easy discovery. Hit enter after each

Description

Describe your data as well as you can. Formatting is preserved when pasting from other sources and counts towards character limits

Funding

Search grant by name/number or add your own

+ Add another grant

References

Link to references or related content

Licence (what's this?)

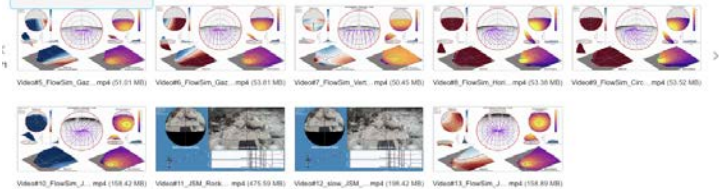
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Tips

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Retinal Optic Flow During Natural Locomotion - Videos

Media posted on 27.01.2022, 12:20 authored by Jonathan Mathis, Karl S. Miller, Kathryn L. Bornn, Mary M. Hayhoe

Store of the videos associated with the publication titled "Retinal Optic Flow during Natural Locomotion" published in *PLoS Computational Biology* <https://doi.org/10.1371/journal.pcbi.1009575>

See related dataset at: <https://doi.org/10.25420/figshare.plus.17113883>

Author Summary

We recorded the full body kinematics and binocular gaze of humans walking through real-world natural environment and estimated visual motion (optic flow) using both computational video analysis and geometric simulation. Contrary to the established theories of the role of optic flow in the control of locomotion, we found that eye-movement-free, head-centric optic flow is highly unstable due to the complex phasic trajectory of the head during natural locomotion, rendering it an unlikely candidate for heading perception. In contrast, retina-centered optic flow consisted of a regular pattern of outflowing motion centered on the fovea. Retinal optic flow contained highly consistent patterns that specified the walker's trajectory relative to the point of fixation, which may provide powerful, retinotopic cues that may be used for the visual control of locomotion in natural environments. This examination of optic flow in real-world contexts suggest a need to re-evaluate existing theories of the role of optic flow in the visual control of action during natural behavior.

Abstract

We examine the structure of the visual motion projected on the retina during natural locomotion in real world environments. Bipedal gait generates a complex, rhythmic pattern of head translation and rotation in space, so without gaze stabilization mechanisms such as the vestibular-ocular-reflex (VOR) a walker's visually specified heading would vary dramatically throughout the gait cycle. The act of fixation on stable points in the environment nulls image motion at the fovea, resulting in stable patterns of outflow on the retinae centered on the point of fixation. These outflowing patterns retain a higher order structure that is informative about the stabilized trajectory of the eye through space. We measure this structure by applying the curl and divergence operations on the retinal flow velocity vector fields and found features that may be valuable for the control of locomotion. In particular, the sign and magnitude of foveal curl in retinal flow specifies the body's trajectory relative to the gaze point, while the point of maximum divergence in the retinal flow field specifies the walker's instantaneous overground velocity/momentary vector in retinotopic coordinates. Assuming that walkers can determine the body position relative to gaze direction, these time-varying retinotopic cues for the body's momentum could provide a visual control signal for locomotion over complex terrain. In contrast, the temporal variation of the eye-movement-free, head-centered flow fields is large enough to be problematic for use in steering towards a goal. Consideration of optic flow in the context of real-world locomotion therefore suggests a re-evaluation of the role of optic flow in the control of action during natural behavior.

FUNDING

CPS Training Grant
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GAZE AND THE VISUAL CONTROL OF FOOT PLACEMENT WHEN WALKING OVER ROUGH TERRAIN
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Draft (Version 3)

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Luca Belletti X + Add contributor

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Published version: Version 2

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SOURCES

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Artificial_Voice_Assistant_for_COVID_19_Suspects
Artificial Voice Assistant for COVID-19 Suspects Artificial Voice Assistant for COVID-19 Suspects
Published 29 December 2021 | Mendeley Data
COVID-19... Covid-19 Survey without 0 or 1.csv
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SMAP run
Wouter Haak, Anita de Waard
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Higher airborne pollen concentrations correlated with increased SARS-CoV-2 infection rates, as evidenced from 31 countries across the globe
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COVID-19

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Artificial_Voice_Assistant_for_COVID_19_Suspects

Published: 29 December 2021 | Version 1 | DOI: 10.17632/jnmxmgk7mk.1
Contributor: Artificial Voice Assistant for COVID-19 Suspects Artificial Voice Assistant for COVID-19 Suspects

Description

COVID-19 outbreak occurred from China which spreads between people through close contact of the infected person. In this pandemic, managing such a high number of patients is difficult. Already, 5.3 million people have died. For this disease, people from all over the world became interested in telemedicine. Till now lots of people solved many problems using telemedicine that's why we are proposing Artificial intelligence voice assistant that can help whether a particular person is COVID-19 suspected or not. Our artificial voice Assistant to help people deal with this type of circumstance. We design some questionnaires that will be asked by the machine and user will answer accordingly. Then based on their answer machine will analysis and predict, whether that particular user might have COVID or not. We took a survey and collected data from different 513 peoples. We applied several machine learning algorithms like Gini Index, Random Forest, Entropy, KNN, Decision Tree. Out of that random forest provide us the highest accuracy. Throughout the research, using random forest algorithm, we had a 92.85% prediction accuracy, which was reasonable. So, our ultimate goal is during this kind of epidemic, medical voice assistant, assists people in overcoming any problems they may be experiencing.

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Institutions

American International University Bangladesh

Categories

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Version 1
Published: 29 Dec 2021
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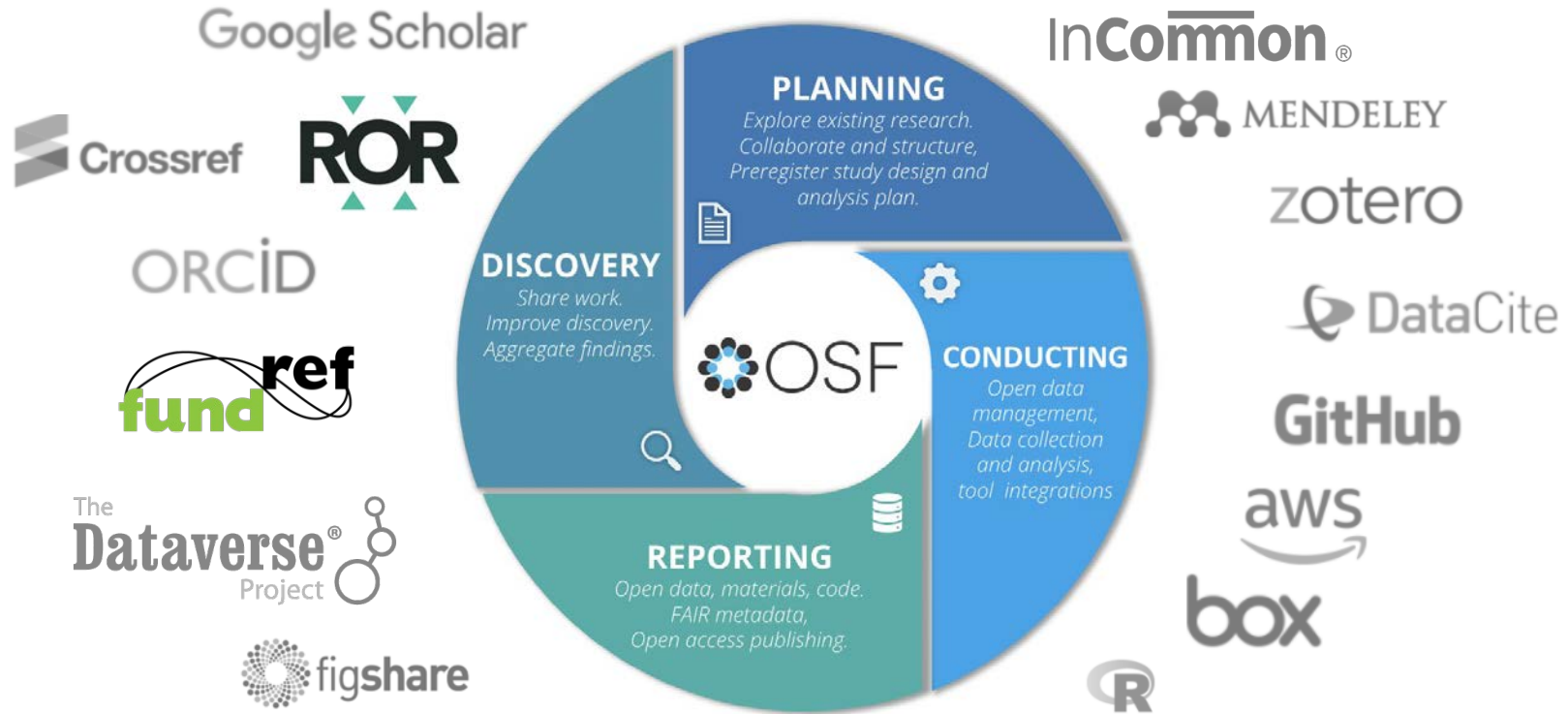
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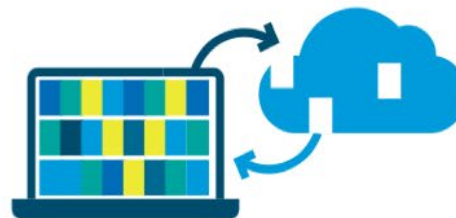


Vivli – The Organization

- Non-profit membership organization



- Sharing individual participant-level data from 6,800 studies



- 43 members from industry, academia, and NIH (BioLINCC, ImmPORT, etc.)



Vivli – The Platform

- ✓ A user-friendly, secure, state-of-the art platform that meets all NIH Desirable characteristics
-

- ✓ We make it as easy as possible to share and analyze NIH-funded human subjects data

- Submitted data must be anonymized
- Managed access: data requesters submit request and sign DUA



- ✓ In 2023, we will be implementing CRediT for research teams who share their data on Vivli



Contributor Roles Taxonomy

Vivli's NIH Data Sharing Model

Current NIH model

1) Vivli supports discoverability of NIH domain repositories



Promotes discoverability of NIH domain repository studies (NCT-IDs) for those datasets available for cross-listing on Vivli (BioLINCC, ImmPort, NIAID, etc.)

2) NIH grantees can directly share their data on Vivli



If no domain repository exists, Vivli will support NIH grantees to share their anonymized data to meet their grant commitments.

3) Data Aggregation



NIH-funded datasets may be aggregated with industry and other datasets on Vivli through either a secure environment or via download



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- FAIR data

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Multi-disciplinary - custom metadata

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December 31, 2018

Fig. 7 in A mountain of millipedes VI. New records, new species, a new genus and a general discussion of Odontopygidae from the Udzungwa Mts, Tanzania (Diplopoda, Spirostreptida, Odontopygidae)

Enghoff, Henrik

Fig. 7. *Hoffmanides dissutus* (Hoffman, 1963), ♂, from Udzungwa Mts National Park. Photograph by A. Illum. Scale bar = 5 mm.

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Publication date: December 31, 2018

DOI: 10.5281/zenodo.1146170

Keyword(s): Biodiversity, Taxonomy, Animals, Diplopoda, Spirostreptida, Odontopygidae

Published in: European Journal of Taxonomy

Related identifiers: Cited by: <http://treatment.plazi.org/id/03DF87B7FF86FFBF6F68FF340F5BFE10> (Taxonomic treatment)

Part of: 10.5852/fig.2018.394, <https://doi.org/10.5852/fig.2018.394>, <https://doi.org/10.5852/fig.2018.394>, <https://doi.org/10.5852/fig.2018.394>

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Pidapalooza 2021, 27-28.01.2021 (Session Block 2, track 2)

Related identifiers:

Cites

10.11646/zootaxa.4193.3.7 (Publication)
10.1653/0015-4040(2005)88[502:KTTFOC]2.0.CO;2 (Publication)
10.11646/zootaxa.4747.2.10 (Publication)

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<http://treatment.plazi.org/id/03DF87B7FF86FFBF6F68FF340F5BFE10> (Taxonomic treatment)
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10.5281/zenodo.5497141 (Figure)
10.5281/zenodo.5497144 (Figure)

Custom keywords:

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Order [Paucituberculata](#)
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R / V Alis, EBISCO Expedition, st. DW 2613, Plateau des Chesterfield (-19.616667, 158.7) NEW CALEDONIA: 42.5 × 19.4 mm, R/V Alis, EBISCO Expedition, st. DW 2613, Plateau des Chesterfield, 19°37' S, 158°42' E, 519-522 m (MNHN IM-2007-30854; Fig. 9A; GenBank accession number (cox1 sequence): KJ550158).
R / V Alis, EBISCO Expedition, st. DW 2610, Plateau des Chesterfield (-19.566668, 158.683333) 16.3 × 13.0 mm, R/V Alis, EBISCO Expedition, st. DW 2610, Plateau des Chesterfield, 19°34' S, 158°41' E, 486-494 m (MNHN IM-2000-30789; paratype 2; Fig. 9D; fragment of the spire, well preserved).
Coral Sea (-19.616667, 158.7) NEW CALEDONIA: Coral Sea, Plateau des Chesterfield, 19°37' S, 158°42' E, 519-522 m (EBISCO st. DW 2613).



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17 January 2019

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https://docs.github.com/en/repositories/archiving_a_github_repository/referencing_and_citing_content

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GitHub / Releases

Release	Status	Time
1.0.14 zenodo-testing/my-project: test	Published	1 year, 11 months ago
1.0.12 zenodo-testing/my-project: test	Published	1 year, 11 months ago
1.0.0 zenodo-testing/my-project: Test	Published	2 years ago
11 zenodo-testing/my-project: test2	Published	2 years ago

November 13, 2019

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Sebastian Troitzsch, Tommaso Mori, Anthony Vautrin

New features

- Demand side flexibility evaluation
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Available in **GitHub**

Publication date: November 13, 2019

DOI: 10.5281/zenodo.3540372

Related identifiers: Supplement to <https://github.com/TUMCREATE/ESTL/cobmo/tree/0.3.0>

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Versions

Version	Published
Version 0.3.0	Nov 13, 2019
Version 0.2.0	Oct 31, 2019
Version 0.1.0	Mar 5, 2019

Files (2.4 MB)

Name	Size
TUMCREATE-ESTL/cobmo-0.3.0.zip	2.4 MB
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