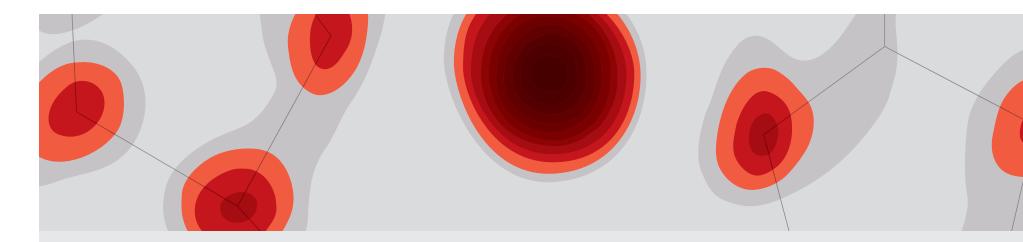
#### Part of **SPRINGER NATURE**



#### Making data publication a first class research output

Andrew L. Hufton

Managing Editor, Scientific Data

https://www.nature.com/sdata/

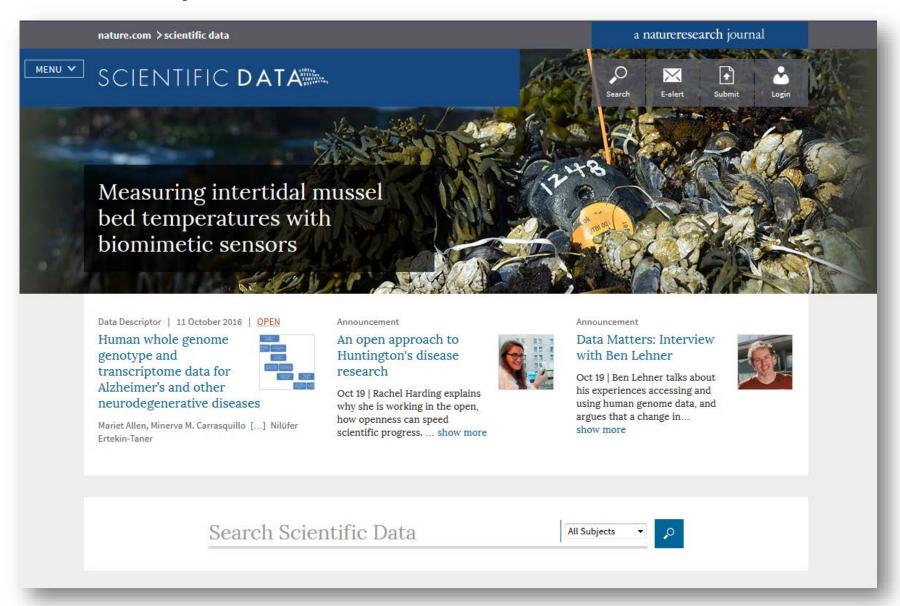
Helping Researchers Publish,

University of Cambridge,

Oct 2017

natureresearch

#### Launched in May 2014



#### the data paper





- A clear, peer reviewed description of data, to maximize usage
- Citable publications that give credit for reusable data
- Visit our journal homepage to learn more: <a href="https://www.nature.com/sdata/">https://www.nature.com/sdata/</a>



#### **Get Credit for Sharing Your Data**

Publications will be indexed and citeable.



#### **Open-access**

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#### **Focused on Data Reuse**

All the information others need to reuse the data; no interpretative analysis, or hypothesis testing



#### Peer-reviewed

Rigorous peer-review focused on technical data quality and reuse value



#### **Promoting Community Data Repositories**

Not a new data repository; data stored in community data repositories



## The Data Descriptor article-type

1

## Data Descriptor Focus on data reuse

- Detailed descriptions of the methods and technical analyses supporting the quality of the measurements.
- Does not contain tests of new scientific hypotheses

#### **Sections:**

- Title
- Abstract
- Background & Summary
- Methods
- Data Records



- Usage Notes
- Figures & Tables
- References
- Data Citations



All the samples used in this study are summarized in Table 1. Consistent identifiers are used in Tables 2 and 3 to allow mapping between the proteomic and transcriptomic data outputs.

#### Data Record 1

The raw data, peaklists (.mgf), ProteomeDiscoverer result files (.msf) and ProteomeDiscoverer workflow files (.xml) have been uploaded to ProteomeXchange (http://www.proteomexchange.org/) with the following accession number PXD000134 (ref. 67; Table 2).

#### Data Record 2

Microarray data are available at the NCBI Gene Expression Omnibus (GEO) database under the accession numbers GSE26451 (ref. 68) and GSE26453 (ref. 69; Table 3).

#### Data Record 3

The peptide and protein identification data sets have been annotated by The Global Proteome Machine at http://gpmdb.thegpm.org/

#### Data Record 4

The peptide and protein identification data sets have been annotated by the StemCellOmicsRepository (SCOR) at http://scor.chem.wisc.edu/

#### **Data Citations**

67. Low, T. Y. et al. ProteomeXchange: PXD000134 (2013).

68. Chin, A. et al. Gene Expression Omnibus: GSE26451 (2011).

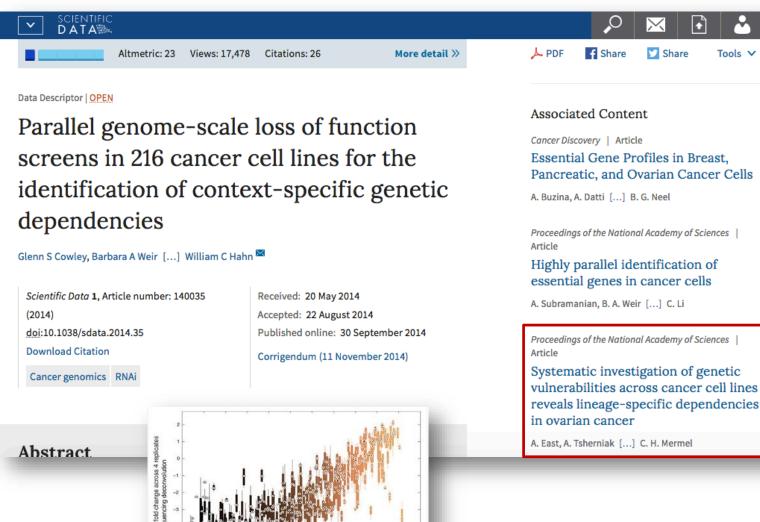
69. Chin, A. et al. Gene Expression Omnibus: GSE26453 (2011).



#### High-throughput screening

#### SCIENTIFIC DATA

Tools V



- Screen results and in-depth analysis published in 2011 at PNAS
- Full screen data published at Scientific Data in 2014
- Data at **figshare**
- Data Descriptor cited 117 times according to Google Scholar!

#### Neuroscience

#### SCIENTIFIC DATA (1011) (1011) (1011) (1011) (1011)





Altmetric: 57

Citations: 27

More detail >>

Data Descriptor | OPEN

A high-resolution 7-Tesla fMRI dataset from complex natural stimulation with an audio movie

Michael Hanke <sup>™</sup>, Florian J. Baumgartner, Pierre Ibe, Falko R. Kaule, Stefan Pollmann, Oliver Speck, Wolf Zinke & Jörg Stadler

Scientific Data 1, Article number: 140003 (2014)

doi:10.1038/sdata.2014.3

Received: 04 November 2013

Accepted: 22 January 2014

Published online: 27 May 2014

#### Previously unpublished dataset

- Data in OpenfMRI
- Cited 42 times according to Google Scholar
- Authors ran an analysis challenge after publication



**Code in GitHub** 

#### **Data Citations**

Abstract • Background & Summary • Methods • Data Records • Technical Validation • Usage Notes • Additional information • References • Data Citations • Acknowledgements • Author information • Supplementary information

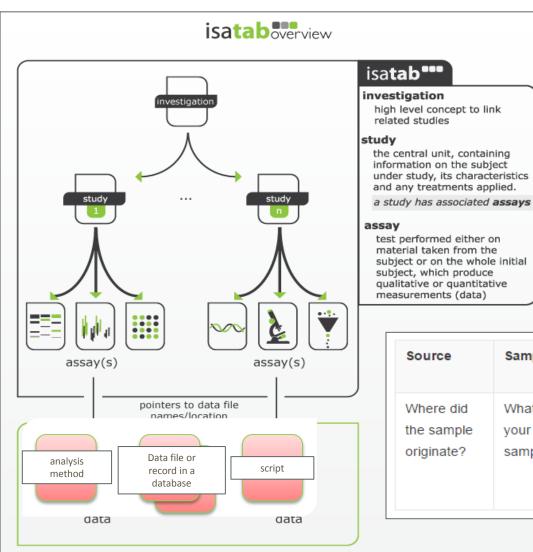
 Hanke, M., Baumgartner, F. J., Ibe, P., Kaule, F. R., Pollmann, S., Speck, O., Zinke, W., & Stadler, J. OpenfMRI ds000113 (2014).

#### Additional resources

- More information and updates available at: <a href="http://www.studyforrest.org">http://www.studyforrest.org</a>
- Source code repository: http://github.com/hanke/gumpdata

### Data Descriptor

Structured metadata



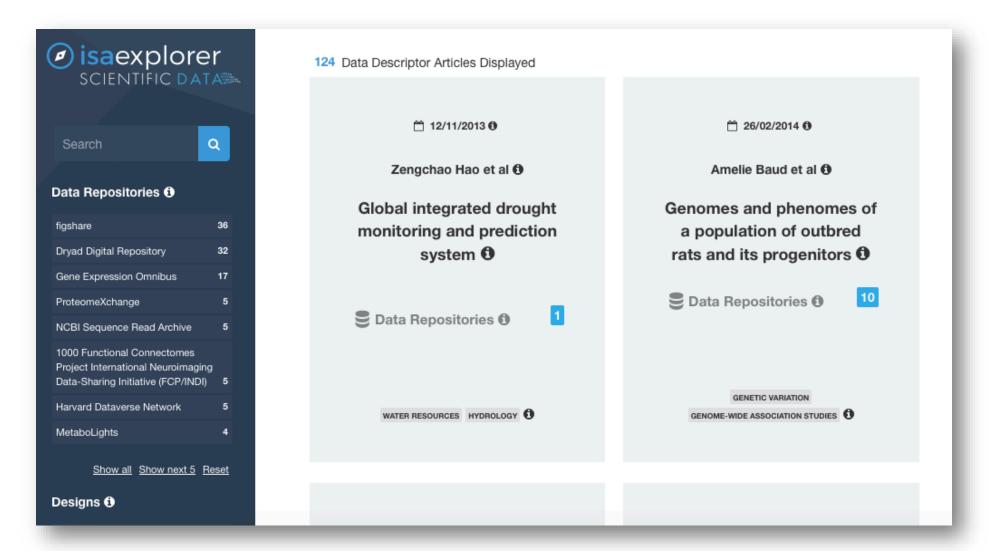
- assists users to submit the structured content via simple templates and an internal authoring tool
- performs value-added semantic annotation of the experimental metadata

natureresearch

Source	Sample	Characteristics	Protocol	Data
Where did the sample originate?	What is your sample?	What do users need to know about your source and samples to fully understand your work?	How did the samples become data?	Where is the data?

#### In-house curation team:

#### Explore this structured metadata with ISA-explorer



## Open data is about more than disclosure – it must be "FAIR"

- Findable
- Accessible
- Interoperable
- Re-usable

Wilkinson et al. Sci. Data doi:10.1038/sdata.2016.18 (2016) https://www.nature.com/articles/sdata201618

## Helping authors find the right repository for their data

2

#### Find the right repository for your data



http://www.nature.com/sdata/policies/repositories

Browse our recommended data repository online.

- We currently list more than 90 repositories, across the biological, physical and social sciences
- We advise authors on the best place to store their data
- Support the use of institutional repositories, including the University of Cambridge data repository (<a href="https://www.data.cam.ac.uk/repository">https://www.data.cam.ac.uk/repository</a>)

# A rigorous, community-based editorial process

3

## Criteria for publication

https://www.nature.com/sdata/policies/for-referees

- Acceptance for publication is based on the
  - technical rigour of the procedures used to generate the data
  - the reuse value of the data
  - the completeness of the data description
- Evaluation is not be based on the perceived impact or novelty of the findings associated with the datasets.

## Data policies

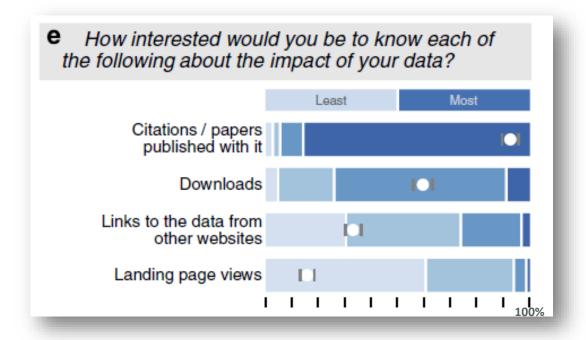
https://www.nature.com/sdata/policies/data-policies

- Editors and peer-reviewers have the right to view the data during their evaluation
  - Before peer-view we check:
    - Links to the datasets work (see first page of PDF)
    - Data are hosted in a repository that meets our policies
- At time of publication, authors must publicly release the data under terms that permit wide reuse (i.e. as "open data")
  - Restrictions on commercial use or redistribution are not permitted

# Promoting a culture of credit for data sharing

4

## Researchers want to know when their data is used in published works

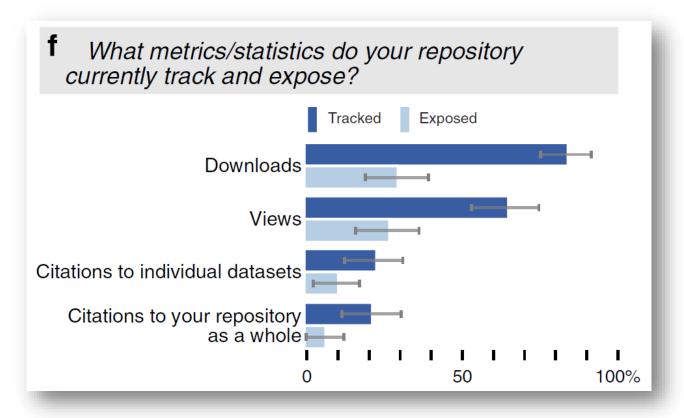


From

#### Making data count

John E. Kratz & Carly Strasser. *Sci. Data* doi:10.1038/sdata.2015.39 (2015)

#### But repositories have a hard time tracking actual data use



From

#### Making data count

John E. Kratz & Carly Strasser. *Sci. Data* doi:10.1038/sdata.2015.39 (2015)

#### In-article data citation

SCIENTIFIC DATA | DATA DESCRIPTOR OPEN



Plant traits, productivity, biomass and soil properties from forest sites in the Pacific Northwest, 1999–2014

The dataset (NACP TERRA-PNW: Forest Plant Traits, NPP, Biomass, and Soil Properties, 1999–2014) is hosted with other contributions from the North American Carbon Program (NACP) by the Oak Ridge National Laboratory Distributed Active Archive Center for Biogeochemical Dynamics (Data Citation 1) Oak Ridge National Laboratory Distributed Active Archive Center

2016



#### **Data Citations**

Abstract • Background & Summary • Methods • Data Records • Technical Validation • Additional Information • References • Data Citations • Acknowledgements • Author information

 Law, B. E., & Berner, L. T. Oak Ridge National Laboratory Distributed Active Archive Center http://dx.doi.org/10.3334/ORNLDAAC/1292 (2015).

Plant trait measurements are needed for evaluating ecological responses to environmental

ACCINITIONS repulgives earthers Publish, Cambridge 2017/alanmont parameterization and testing. We



### Get the most from your data

Preserve it
Encourage reuse
Get credit

Encourage others to do the same

## SCIENTIFIC DATA 11011010 Thanks!

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