

Forschungsdaten-Repositorien

Typen, Herausforderungen und Perspektiven

Heinz Pampel, Deutsches GeoForschungsZentrum GFZ

Open-Access-Tage 2012, Wien, 26.09.2012

Agenda

- Hintergrund
- Forschungsdaten-Repositorien
- re3data.org
- Vernetzung

The Royal Society

Recommendation 1

Scientists should communicate the data they collect and the models they create, via methods that allow free and open access, and in ways that are intelligible, assessable and usable for other specialists in the same or linked fields wherever they are in the world. Where the data justifies it, scientists should make them available in an appropriate data repository.

Where possible, communication with a wider public audience should be made a priority, and particularly so in areas where openness is in the public interest.

The Royal Society. (2012). Science as an open enterprise. The Royal Society Science Policy Centre report 02/12. Retrieved from http://royalsociety.org/uploadedFiles/Royal_Society_Content/policy/projects/sape/2012-06-20-SAOE.pdf

The Royal Society

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EC: Empfehlung an die Mitgliedstaaten

E-infrastructures

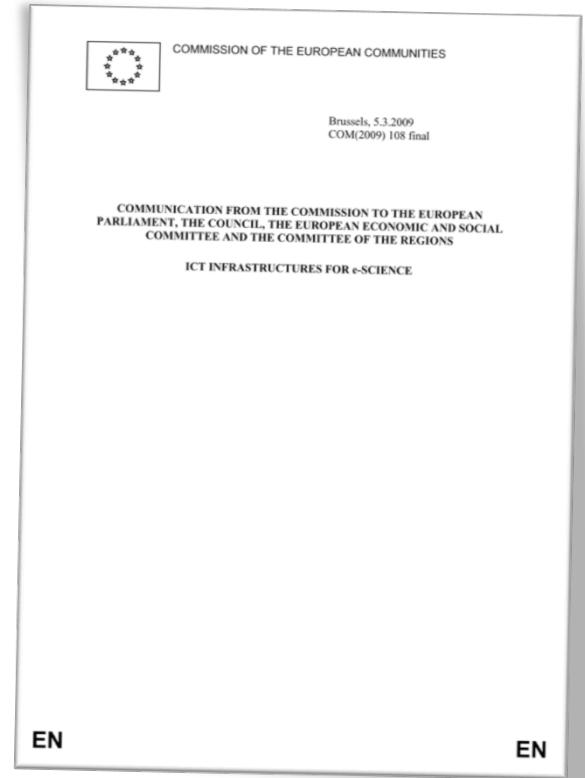
5. Further develop e-infrastructures underpinning the system for disseminating scientific information by:
 - Supporting scientific data infrastructures for dissemination of knowledge, research institutions and funding entities to address all stages of the data life cycle. These stages should include acquisition, curation, metadata, provenance, persistent identifiers, authorisation, authentication and data integrity. Approaches need to be developed to provide a common look and feel to data discovery across disciplines, thus reducing the learning curve required to achieve productivity;

European Commission. (2012). Commission Recommendation on access to and preservation of scientific information. C(2012) 4890 final. Retrieved from http://ec.europa.eu/research/science-society/document_library/pdf_06/recommendation-access-and-preservation-scientific-information_en.pdf

Forschungsdaten-Repositorien

- EC: ICT infrastructures for e-science

„The landscape of data repositories across Europe is fairly heterogeneous, but there is a solid basis to develop a coherent strategy to overcome the fragmentation and enable research communities to better manage, use, share and preserve data.“



European Commission. (2009). ICT infrastructures for e-science. Communication from the Commission to the European Parliament, the Council, the European Economic and Social Committee and the Committee of the Regions. COM(2009) 108 final. Retrieved from <http://eur-lex.europa.eu/LexUriServ/LexUriServ.do?uri=COM:2009:0108:FIN:EN:PDF>

Forschungsdaten-Repositorien

The screenshot shows the NCBI GEO homepage. At the top left is the NCBI logo. In the center is the GEO logo with the text "Gene Expression Omnibus". Below the logo are links for "GEO Publications", "FAQ", "MIAME", and "Email GEO". A "Login" link is at the bottom right. The main content area has a heading "Gene Expression Omnibus: a public functional genomics data repository supporting submissions. Array- and sequence-based data are accepted. Tools are provided to analyze experiments and curated gene expression profiles. More information »". Below this is a "GEO navigation" section with two main categories: "QUERY" and "BROWSE". The "QUERY" category contains links for "DataSets", "Gene profiles", "GEO accession", and "GEO BLAST", each with a "GO" button. The "BROWSE" category contains links for "DataSets" and "GEO accessions", which further lead to "Platforms", "Samples", and "Series".

The screenshot shows the PANGAEA website. At the top right is the PANGAEA logo with the tagline "Data Publisher for Earth & Environmental Science". Below the logo is a search bar with dropdown menus for "All", "Water", "Sediment", "Ice", and "Atmosphere". Underneath the search bar are links for "Help", "Advanced Search", "Preferences", and "more...". At the bottom of the page are links for "About", "Submit Data", "Projects", "Software", "WDC-MARE", and "Contact". A note at the bottom states "This work is licensed under a Creative Commons License".

GEO, <http://www.ncbi.nlm.nih.gov/geo/>

PANGAEA, <http://www.pangaea.de>

Forschungsdaten-Repositorien

Scientific Drilling Database

Data from Deep Earth Sampling and Monitoring

- + Home
- + About SDDB
- + News
- + Data Publications
- + Catalogue
- + Authors
- + Dataset
- + Research Programs
- + Sampling Gear
- + Analytical Parameters
- + Publication Process
- + Admin

GFZ
POTS DAM

icdp

SDDB, <http://www.scientificdrilling.org>

Dataset Description

Citation: Demory, Francois; Nowaczyk, Norbert; Witt, An
Intercorrelation of down-core variations of the hig
for CON01-603 and CON01-605. *Scientific Driller*
/GFZ.SDDB.1003

[Download Citation \(EndNote\)](#)

Title: Intercorrelation of down-core variations of the hig
for CON01-603 and CON01-605

Abstract: Down-core variations of the high-resolution mag
been obtained for kasten and pilot cores from sit
allowed the transfer of AMS 14C dating performe
on kasten cores to the pilot cores subjected to g
discrepancies in the magnetic susceptibility cur
01-605 (Vydriño Shoulder) measured in 2001 and
large peaks visible in the first measurement from
after a 2-year-long storage. This is a first hint for
chemically unstable greigite.

[Show in Google Earth](#)

Related Publications:

- N. Fagel, L. Y. Alleman, L. Granina, F. Ha
André, Vivianite formation and distribution
Planet. Change 46 (2005), pp. 315-336. d

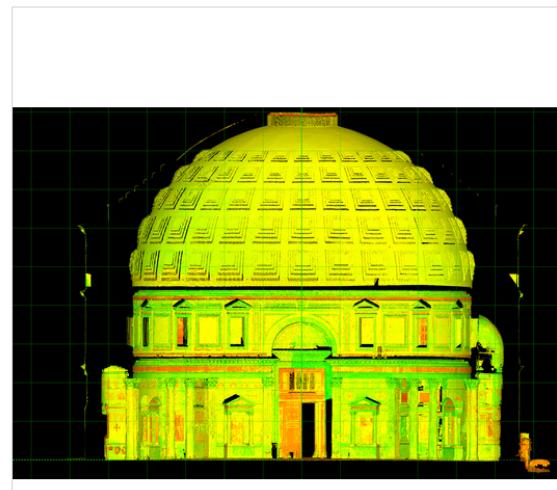
The Bern Digital Pantheon Project

Home The Building Research Publications The Team Repository Contact Search

You are here: Home > Repository > BDPP0019

Repository

first previous Record 198 of 776 next last



The Bern Digital Pantheon Project: Transverse cross-section, facing north

The Bern Digital Pantheon Project: Transverse cross-section, facing north. In: Gerd Graßhoff, Markus Wälfle, Jon Albers and Christian Berndt (ed.); Digital Repository of the Bern Digital Pantheon Project. Bern 2009, BDPP0019 (Published online at: <http://www.digitalpantheon.ch/BDPP0019>).

BibTeX:

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@other{BDPP0019,
  author = {The Bern Digital Pantheon Project},
  title = {(Transverse cross-section, facing north)},
  year = {2009},
  keywords = {Medium Visualisation Building Interior},
  owner = {berndt},
  timestamp = {2009.05.28},
  editor = {Graßhoff, Gerd and Wälfle, Markus and Albers, Jon and Berndt, Christian},
  address = {Bern},
  repository = {Digital Repository of the Bern Digital Pantheon Project}
}
```

Usage of the repository

Forschungsdaten-Repositorien

PURDUE UNIVERSITY

Purdue University Research Repository | PURR

Home Browse Content Projects Get Started Contact Us Search

You are here: Publications > Datasets > Phosphorus and Potassium Influence on Alfalfa ... > About

Phosphorus and Potassium Influence on Alfalfa Nutrition

By Jeffrey J Volenec

Data from several studies on the influence of phosphorus and potassium nutrition on alfalfa. It includes numeric data such as yield, plant mass, plant counts, and tissue concentration of various nutrients

Listed in Datasets

About Supporting Docs Versions Reviews Questions Wish List

Download (XLS)
Additional materials available

Version 1.0 - published on Feb 2
doi:10.5072/FK2NG4S08 - cite this

Suche: Home | Browse | Erweiterte Suche | Hilfe

English Anmelden | Registrieren

Zitation: Open Data LMU: Dohrmann, Martin; Göcke, Christian; Janussen, Dorte; Reitner, Joachim; Lüter, Carsten und Wörheide, Gert (2011): Additional data to "Systematics and spicule evolution in dictyonal sponges (Hexactinellida: Sceptrulophora) with description of two new species", doi:10.5282/ubm/data.36

Abstract This is a master file of merged excel files with data from several studies the influence of phosphorus and potassium nutrition on alfalfa. It includes numeric data such as yield, plant mass, plant counts, and tissue concentration of various nutrients. Conventional wisdom states that plants that are poorly fertilized and do not have adequate phosphorus and especially low potassium will become winterkill. This study showed that plants did not necessarily die in winter with low potassium stress.

Anderes (Sequence data, structure & tree file)
11Kb

Anderes (Additional movie M1_Dive4)
67Mb

Anderes (Additional movie M2_Dive2)
74Mb

Anderes (Additional movie M3_Dive6)
249Mb

DOI: <http://dx.doi.org/10.5282/ubm/data.36>

Beschreibung

In this paper we report on recently collected specimens of glass sponges belonging to Farreidae Gray, 1872 and Tretodictyidae Schulze, 1886 (Porifera: Hexactinellida: Hexactinosida). All specimens represent new geographical records for their genera: Coral Sea for *Acanthocionula* Reiswijk, 2002 (Farreidae) and *Pellicularv* Tiima, 1927 (Tretodictyidae); W. Atlantic for *Sarostegia* Tonigkeit, 1904.

Open Data LMU, <http://data.ub.uni-muenchen.de>

Forschungsdaten-Repositorien

The screenshot shows a research dataset titled "Herring Hepatic Transcriptome 34300 contigs.fa" on the figshare platform. The page includes a navigation bar with "My data", "Browse", "Upload", and a user profile for "H. Pampel". The main content area displays the dataset's content, which is a FASTA file containing genomic sequences. Below the content are statistics: 3677 views, 7 shares, and 0 cites coming soon. The dataset was published on Jan 19, 2012, at 16:37 (GMT) and has a filesize of 22.40 MB. Categories listed include Molecular biology and Marine Biology. Authors are Steven Roberts. Tags include contigs, herring, fasta, and ngs. Export options are available, and social sharing buttons for Facebook, Twitter, and Google+ are present. A preview and download button are also visible.

Herring Hepatic Transcriptome 34300 contigs.fa

```
>ConsensusfromContig1 Average coverage: 2.00
GTGTGTGTGTGTCTACTGCTACTCGCTGTCACTGTCT
CGACGGGGTGTCTAGGCATTTGGGTGTGCTCTCTTGGGGTTCAAGCTGGACAGC
TTGGGGCTGCTGATTGGATGGCTGTGGCGCTGGGCTCTCCATTGGTCCCTGGCC
TCAGGGTCCGCCCTCTGGGCTTGCGGGAGCCTCAGTCTCTGCACTCGCTGCTGCC
TCTGGCTCAGGCTGGCCTCGGCTACTGTGCTCCCGCACCTGACGGAGGAGGG
AGCAACTGCTGCTGCTGTCTTGCTGTGCACTATGTGATGTACATGTTG
TACACCAAGTTGGCATTTCGGCATTTGGCATTTGGCTGGCTGGGAGGAGGAGG
TCTCCAATAGGCTGGGTGACTACTATTAGTCCGTATGTGATGCTAGTCATAAGGGCGCG
GTGAGC
>ConsensusfromContig2 Average coverage: 1.64
TCGAGGAGCTGACGGCATGGCTCTCGCAGAACGACGGACTGGTCAGACCCCTGCGCTG
CAGCTCGAGATCGGGCAGCAAGCAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG
AAGGAGACCCCTCAGCAGCCACCCAGAGGGGGCTGTTGCTGGGGTAAGGGAGGAGG
CCCCACGCTGTAACCCGGCAGCAGCTTACCACTGCTGCCCTCCCCGGCGTGGAGGCCATC
CGCCTCACCGTAAAAGAGGAAGCAGCAGGGAGAGGAGGAGGAGGAGGAGGAGGAGAG
GAGATGGACGAGCTGATGGCGAGCAGAACGACTGGATGTGCG
>ConsensusfromContig3 Average coverage: 3.34
CTGGCTATCAGGTAAATTCCAGGACTCAGTTCACTGGCTTGCTCCAGGTTGTC
TTAGTCTGTCAGGCTCAGGTTCACTGGCTTGCTCCAGGTTGTC
```

Preview Download

Share this: [f Share](#) 0 [Tweet](#) 6 [+1](#)

Cite this: Herring Hepatic Transcriptome 34300 contigs.fa. Steven Roberts. Figshare. Retrieved 10:28, May 11, 2012 (GMT). hdl.handle.net/10779/084d34370bda29bbc67b3c5ecb02575

3677 views 7 shares 0 cites coming soon

Published on 19 Jan 2012 - 16:37 (GMT)
Filesize is 22.40 MB

Categories

- [Molecular biology](#)
- [Marine Biology](#)

Authors

[Steven Roberts](#)

Tags

#contigs #herring #fasta
#ngs

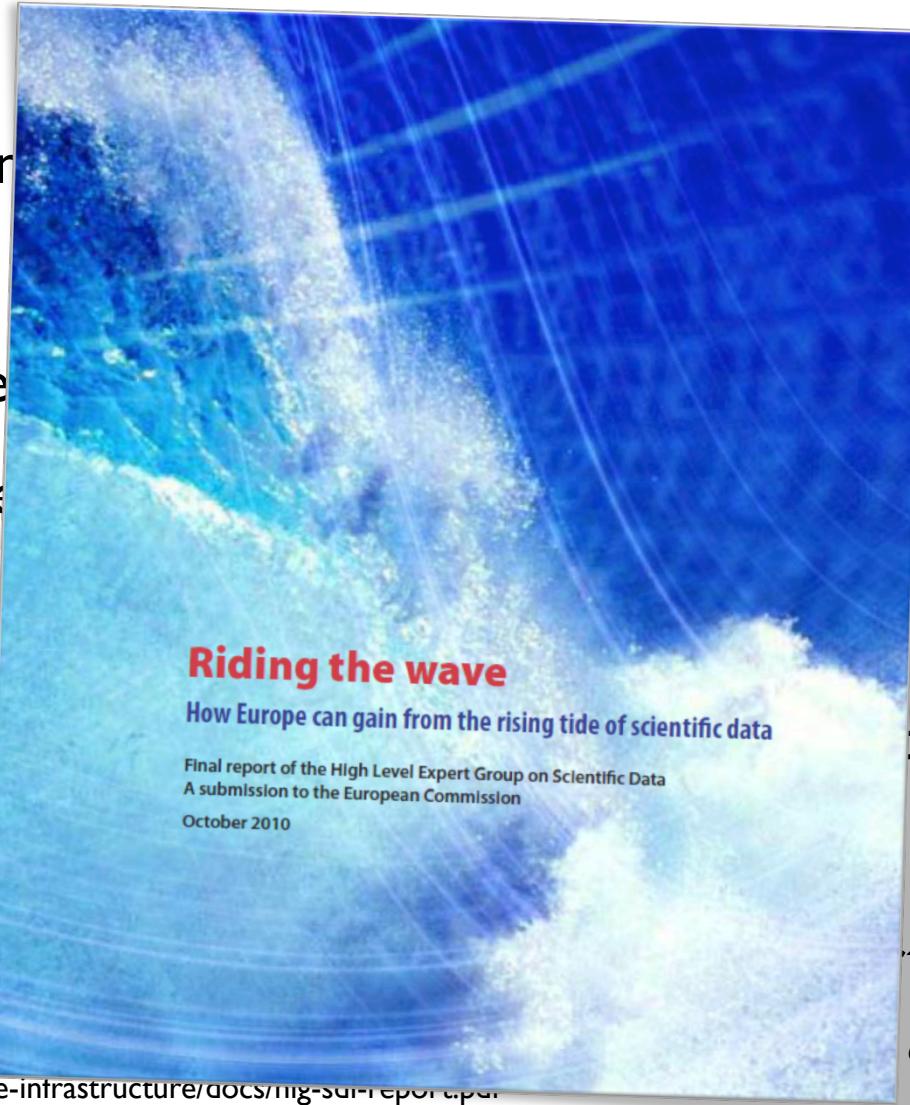
Export

- [Export to Ref. Manager](#)

Figshare, <http://figshare.com>

Vision 2030

- „Researchers are able to find, access and evaluate the data they need and they can evaluate the quality of the data.“
- “Producers of data prefer to deposit their data in international standard repositories. A



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and they
usted.“
access, and
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of scientific data. Retrieved

High Level Expert Group on Scientific Data. (2010). Report on the future of scientific data management in Europe. Brussels: European Commission.
from <http://cordis.europa.eu/fp7/ict/e-infrastructure/docs/mig-sud-report.pdf>.

Vision 2030

- „Researchers and practitioners from any discipline are able to find, access and process the data they need. They can be confident in their ability to use and understand data, and they can evaluate the degree to which that data can be trusted.“
- “Producers of data benefit from opening it to broad access, and prefer to deposit their data with confidence in reliable repositories. A framework of repositories is guided by international standards, to ensure they are trustworthy.”

High Level Expert Group on Scientific Data. (2010). *Riding the wave. How Europe can gain from the rising tide of scientific data.* Retrieved from <http://cordis.europa.eu/fp7/ict/e-infrastructure/docs/hlg-sdi-report.pdf>

Beispiel: Biomedizin

- 200 Repositorien
- 100 Institutionen
- 350 Personen
- > 10.000 Nutzerinnen und Nutzer
- 60 Millionen Zugriffe per Monat
- 35 Millionen EUR jährliche Kosten

Published online 5 December 2011

Nucleic Acids Research, 2012, Vol. 40, Database issue DI-D8
doi:10.1093/nar/gkr1196

The 2012 Nucleic Acids Research Database Issue and the online Molecular Biology Database Collection

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ABSTRACT

The 19th annual Database Issue of *Nucleic Acids Research* features descriptions of 92 new online databases covering various areas of molecular biology and 100 papers describing recent updates to the databases previously described in *NAR* and other journals. The highlights of this issue include, among others, a description of neXtProt, a knowledgebase on human proteins; a detailed explanation of the principles behind the NCBI Taxonomy Database; NCBI and EBI papers on the recently launched BioSample databases that store sample information for a variety of database resources; descriptions of the recent developments in the Gene Ontology and UniProt Gene Ontology Annotation projects; updates on Pfam, SMART and InterPro domain databases; update papers on KEGG and TAIR, two universally acclaimed databases that face an uncertain future; and a separate section with 10 wiki-based databases, introduced in an accompanying editorial. The *NAR* online Molecular Biology Database Collection, available at <http://www.oxfordjournals.org/nar/database/a/>, has been updated and now lists 1380 databases. Brief machine-readable descriptions of the databases featured in this issue, according to the BioDBcore standards, will be provided at the <http://biobanking.org/biobdbcore> web site. The full content of the Database Issue is freely available online on the *Nucleic Acids Research* web site (<http://nar.oxfordjournals.org/>).

COMMENTARY

This current, 19th annual Database Issue of *Nucleic Acids Research* (NAR) features descriptions of 92 new online databases covering a variety of molecular biology data, 77 update papers on databases that have been previously described in the NAR Database Issue and 23 papers with updates on database resources whose descriptions have previously been published in other journals (Table 1). The accompanying *NAR* online Molecular Biology Database Collection (<http://www.oxfordjournals.org/nar/database/a/>) has been revised, which resulted in updating the URLs of more than 30 databases and exclusion of more than 20 obsolete web sites. This list now includes 1380 databases sorted into 14 categories and 41 subcategories.

NEW AND UPDATED DATABASES

This issue contains an unusually high number of papers from the authors' host institutions, NCBI and EMBL-EBI, respectively. In addition to the annual papers from the International Nucleotide Sequence Database collaboration [INSDC (1)], which includes the DNA Data Bank of Japan, GenBank and the European Nucleotide Archive (2–4), Ensembl (5), UniProtKB (6) and the Protein Data Bank in Europe (7), these include two papers that describe the BioSample database project, recently launched at both institutions. The BioSample databases [<http://www.ncbi.nlm.nih.gov/biosample> and <http://www.ebi.ac.uk/biosamples/>, (8) and (9), respectively] aim at capturing essential information about each biological sample used to obtain sequence, gene expression or protein expression data, as well as the relationship between different samples and their sources. The sample

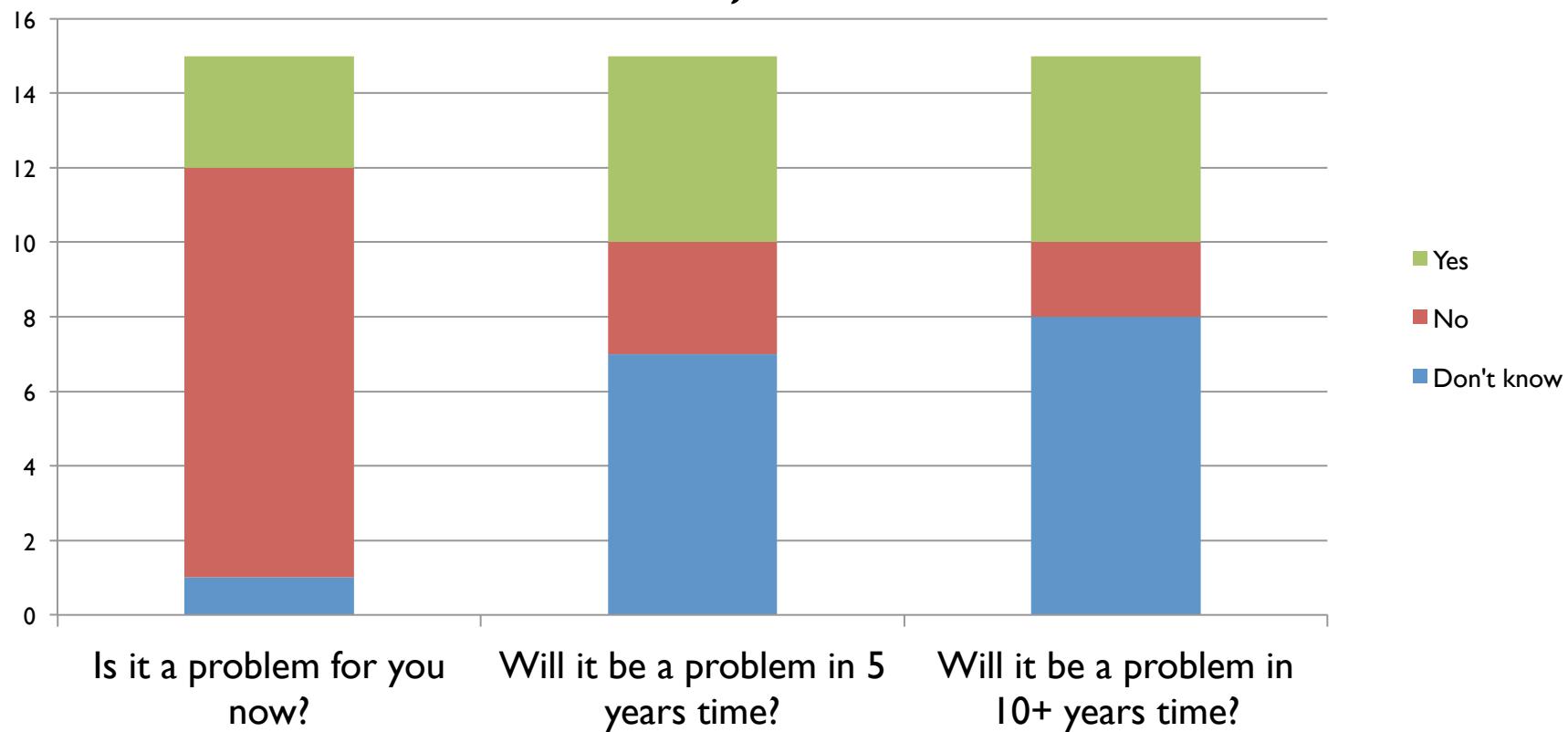
*To whom correspondence should be addressed. Tel: +301 435 5910; Fax: +301 435 7792; Email: galperin@ncbi.nlm.nih.gov
Correspondence may also be addressed to Xosé M. Fernández-Suárez, Tel: +44 (0)1223 494 591; Fax: +44 (0)1223 494 468; Email: xms@ebi.ac.uk

The authors wish it to be known that, in their opinion, the first two authors should be regarded as joint First Authors.

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ELIXIR. (n.d.). *The ELIXIR Strategy for Data Resources. Draft Report from Workpackage 2. The ELIXIR Preparatory Phase*. Retrieved from <http://www.elixir-europe.org/bcms/elixir/Documents/reports/ELIXIR%20Strategy%20for%20Data%20Resources%20report.pdf>

Funding of the repositories - now and in the future, n = 15



Dauerhafte Finanzierung

IN FOCUS NEWS

INFORMATICS

Databases fight funding cuts

Online tools are becoming ever more important to biology, but financial support is unstable.

BY MONYA BAKER

In the era of 'big data', it is a bitter blow for scientists to lose access to the online tools they use to analyse and share terabytes of information. Yet funding cuts by the US National Library of Medicine (NLM) are threatening five widely used biological databases, and user communities are now rallying to save them. "The idea that this resource could just disappear is a serious problem for everyone who relies on it," says Mark Musen, a bioinformatician at Stanford University in California, and manager of Protégé, which provides open-source software to organize and interrelate biological data.

Protégé has 200,000 registered users, and the NLM, part of the National Institutes of Health (NIH) in Bethesda, Maryland, has contributed millions of dollars to maintain it. But

ENDANGERED DATABASES

The US National Library of Medicine (NLM) is cutting resources that biologists say are vital to their research.

Resource	NLM-funded since	Function	Usage	Last NLM award
Protégé	1990	Creating tools to organize and analyse data	200,000 registered users	\$956,625
BioMagResBank	1990	Holds spectroscopy data for biomolecules	500–1,000 unique users per day	\$727,129
Repbase	1994	Identifying families of non-coding DNA across species	8,000 registered users	\$551,544
REBASE	1995	Finding where enzymes bind to and cut DNA	495,844 website hits per month	\$235,911
CASP	2001	Testing techniques to predict protein structure	More than 100 research groups participate	\$515,168

a structural biologist at the University of Wisconsin-Madison, hopes to attract other federal funders to support the database.

shifting commercial priorities.

The least vulnerable databases are those directly run by government agencies, says

Baker, M. (2012). Databases fight funding cuts. *Nature*, 489(7414), 19–19. doi:10.1038/489019a

re3data.org: Mission

- Aufbau eines “Registry of Research Data Repositories”
- Ziel ist ein weltweites, disziplinübergreifendes, web-basiertes Verzeichnis von Forschungsdaten-Repositorien
- Orientierung für Wissenschaftler, Förderer, Verlage, Infrastruktur
- Vernetzung und Standardisierung von Forschungsdaten-Repositorien
- Förderung der “cultur of sharing” und der Sichtbarkeit von Daten

re3data.org: Partner

- Deutsches GeoForschungsZentrum GFZ, LIS
- Humboldt-Universität zu Berlin, IBI
- Karlsruher Institut für Technologie (KIT), KIT Library

GFZ

Helmholtz-Zentrum
POTS DAM



DFG

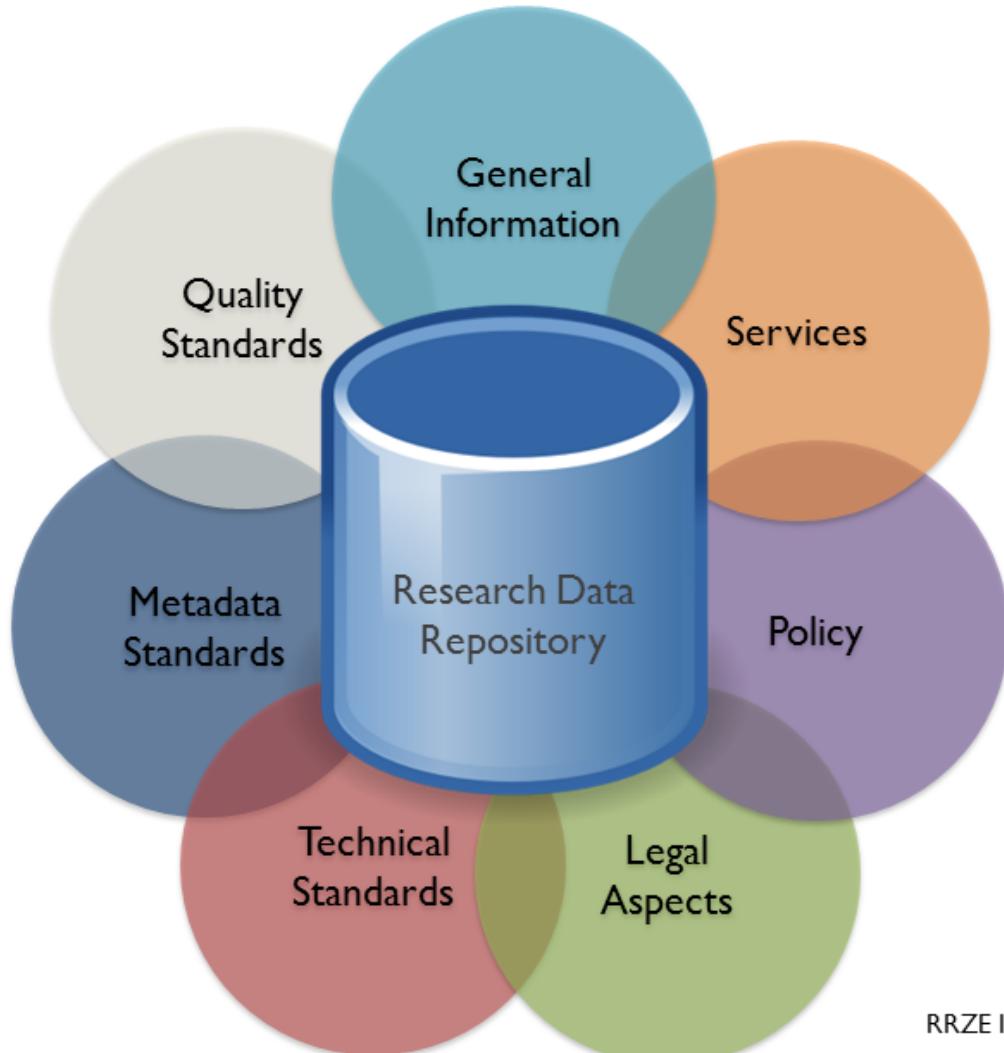


- Förderung durch die DFG (2012-2014)
- Vernetzung im Rahmen von DINI
- MoU mit DataCite

re3data.org: Status quo

- Aktivitäten in der ersten Projektphase:
 - ✓ Konzeption des Registries
 - ✓ Entwicklung eines Metadaten-Schemas
 - ✓ Formulierung von Auswahlkriterien
 - ✓ Technische Entwicklung des Registries
 - ✓ Vernetzung mit relevanten Initiativen
- Onlinegang für Herbst 2012 geplant

re3data.org: Aspekte



RRZE Icon Set (CC: BY-SA)

re3data.org: Zugang/Nachnutzung

Access Icons			
Access to Repository	open	open oder restricted	closed
Access to Data	open, (restricted, closed)	restricted, (closed)	closed
Upload	open oder restricted	open oder restricted	-

Orange unterlegte Werte werden für das jeweilige Icon benötigt, die Werte in Klammern können vorkommen.

re3data.org: Vokabular



Vocabulary for the Registration and Description of Research Data Repositories

Version 1.0

July 2012

doi:10.2312/re3.001

Authors: Paul Vierkant^b, Shaked Spier^b, Jessika Rücknagel^b, Jens Gundlach^c, Maxi Kindling^b, Heinz Pampel^a, Stefka Manova^e, Hans-Jürgen Göbelbecker^e, Jens Klump^a, Roland Bertelmann^a, Peter Schirmbacher^b, Frank Scholze^c

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^c Karlsruhe Institute of Technology (KIT), KIT Library

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- Vierkant, P., Spier, S., Rücknagel, J., Gundlach, J., Kindling, M., Pampel, H., Manova, S., et al. (2012). Vocabulary for the Registration and Description of Research Data Repositories.
- Version 1.0. doi:10.2312/re3.001

re3data.org: Vernetzung

- Vernetzung mit weiteren Projekten (in Auswahl)
 - DFG: KOMFOR, LAUDATIO, RADIESCHEN, ...
 - EU: APARSEN, OpenAIREplus, ODE, EUDAT, ...
 - Helmholtz Open Access Koordinationsbüro
- DINI-Wiki zum Thema im Aufbau
 - AG Elektronisches Publizieren
 - AG Virtuelle Forschungsumgebungen
- Mailingliste: forschungsdaten@listserv.dfn.de



Vocabulary to describe Research Data Repositories online

Posted on [July 26, 2012](#) by [re3data.org team](#)

The first draft of the re3data.org vocabulary to describe research data repositories and its documentation is [online](#) (version 1.0). The vocabulary will be used to index research data repositories gathering issues such as:

- general information (e.g. subject)
- information on the provider (e.g. responsible institutions)
- information on legal aspects (e.g. licenses)
- information on technical, metadata and quality standards (e.g. software, api and certificates).



PARTNERS



Danke für Ihre Aufmerksamkeit!

re3data.org: Team

- Humboldt-Universität zu Berlin, Institut für Bibliotheks- und Informationswissenschaft:
 - Peter Schirmbacher (Leitung), Maxi Kindling, Paul Vierkant, Jessika Rücknagel, Shaked Spier
- Helmholtz-Zentrum Potsdam, Deutsches GeoForschungsZentrum GFZ, LIS
 - Roland Bertelmann (Leitung), Jens Klump, Heinz Pampel
- Karlsruher Institut für Technologie (KIT), KIT-Bibliothek
 - Frank Scholze (Leitung) Hans-Jürgen Goebelbecker, Jens Gundlach, Stefka Manova

info@re3data.org



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