

VIVO

connect
share
discover

THE VIVO PROJECT



Webinar 1: Data Information Models for Scientific Research
June 23, 2022



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Research & Innovation Manager

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Index:

- 1) Introducing the VIVO community
- 2) The VIVO software and the Ontology
- 3) VIVO use cases
- 4) VIVO examples
- 5) Innovation in the VIVO community
- 6) Conclusions

Open Source Community Supported program

- ✓ Software built by, for and with communities
- ✓ Identifying common needs
- ✓ Affordable



<https://www.lyrasis.org/Pages/Main.aspx>



VIVO Core Values



OPEN SOURCE

VIVO, and all VIVO components are provided as open source. Download at GitHub.

VIVO and all components of VIVO are open source. **Download from GitHub.**

USE

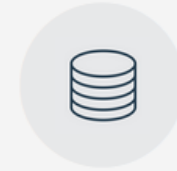


OPEN COMMUNITY

The VIVO community is open to everyone. You can follow the work of VIVO at the VIVO wiki.

The VIVO community is open to everyone. You can follow VIVO's work on our wiki.

JOIN



OPEN DATA

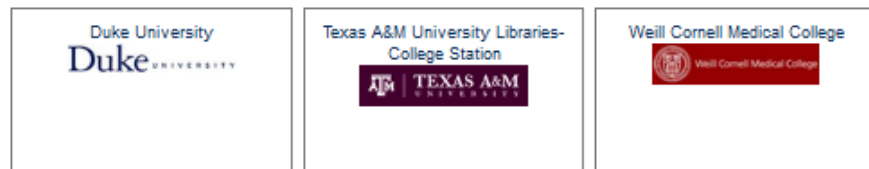
VIVO produces Linked Open Data which is easily shared and combined across VIVO sites.

VIVO produces linked open data that can be easily shared and combined across all VIVO sites.

SHARE

Member-Supported Community

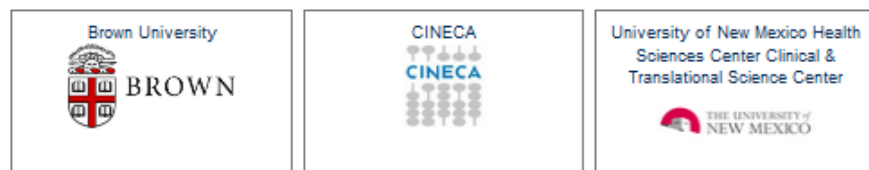
Platinum Members,



Gold Members,



Silver Members,



Bronze Members,



VIVO Service providers:



VIVO Strategic partner (MOU):



Other partners:



Community Organisation

Leadership Group

Define the strategic direction

Committers Group

Developers in charge of the maintenance and evolution of the VIVO base code.

Technical Lead

Users Groups

Community-created groups with common interests, grouped by region or zone:

- North American User Group
- German User Group
- Iberoamerica User Group

Interest Groups

Groups created by the community to support initiatives but without limited time

Task Forces

Groups created by the community with a specific and finite objective in time

Access the
interactive
VIVO map!

+60
institutions
and agencies

+20
countries
implementing
VIVO



A Look Back at 2021-2022

- Partnering with Lyrasis
- Strengthened governance
- Release of VIVO 1.11/1.12
- Pandemic caused financial stress
- New members. Berlin Alliance at Germany

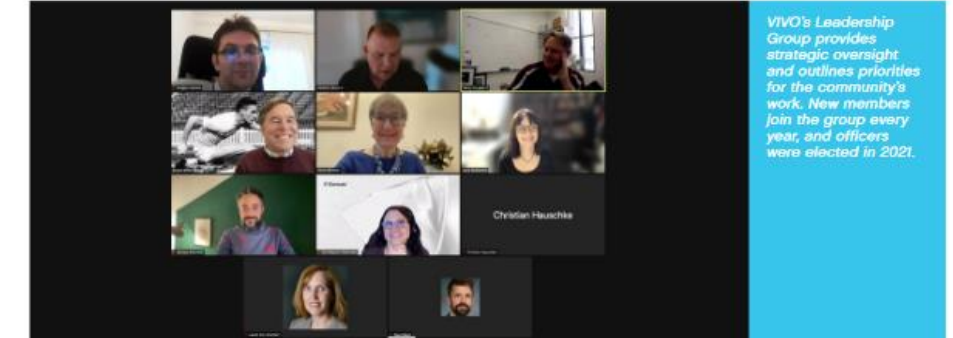
Focus on Community, Members, and the VIVO System

VIVO | connect share discover Annual Report 2021

VIVO is in its thirteenth year as an open-source, member-supported platform for representing scholarship. This Annual Report highlights accomplishments in the fiscal year from July 1, 2020 through June 30, 2021.

There's lots of good news about VIVO. Despite the pandemic, the community made outstanding contributions to VIVO, including a major new release, new ontologies, new members, and four well-attended events.

In countries all over the world, interest, energy and enthusiasm for VIVO grew substantially during this time, presenting opportunities for VIVO leadership and the community.



Metrics: From July 2020 to June 2021

Countries with Active Sites

22

Active Sites

64

New Software Releases

1 1.12.0, released in June 2021

Development Sprints

3

Financials:

Revenue

\$167,010

Expenses

\$128,758

Net Income

\$38,252

Net Assets

\$101,266

2021-22 Community Events

- VIVO 2021 conference (Virtual) – 252 attendants, most international, from 32 countries worldwide
- North American User Group Meeting
- Launch of the Spanish-speaking User Group Meeting with more than 600 attendants
- VIVO track at the CRIS2022
- New German User Group Meeting in progress
- VIVO 2022 conference in progress



VIVO connect share discover

Lanzamiento de la Comunidad
VIVO en español

27 de mayo
14:00 GMT

Seminario Web
VIVO: Un sistema de gestión de la investigación en el contexto de la Ciencia Abierta

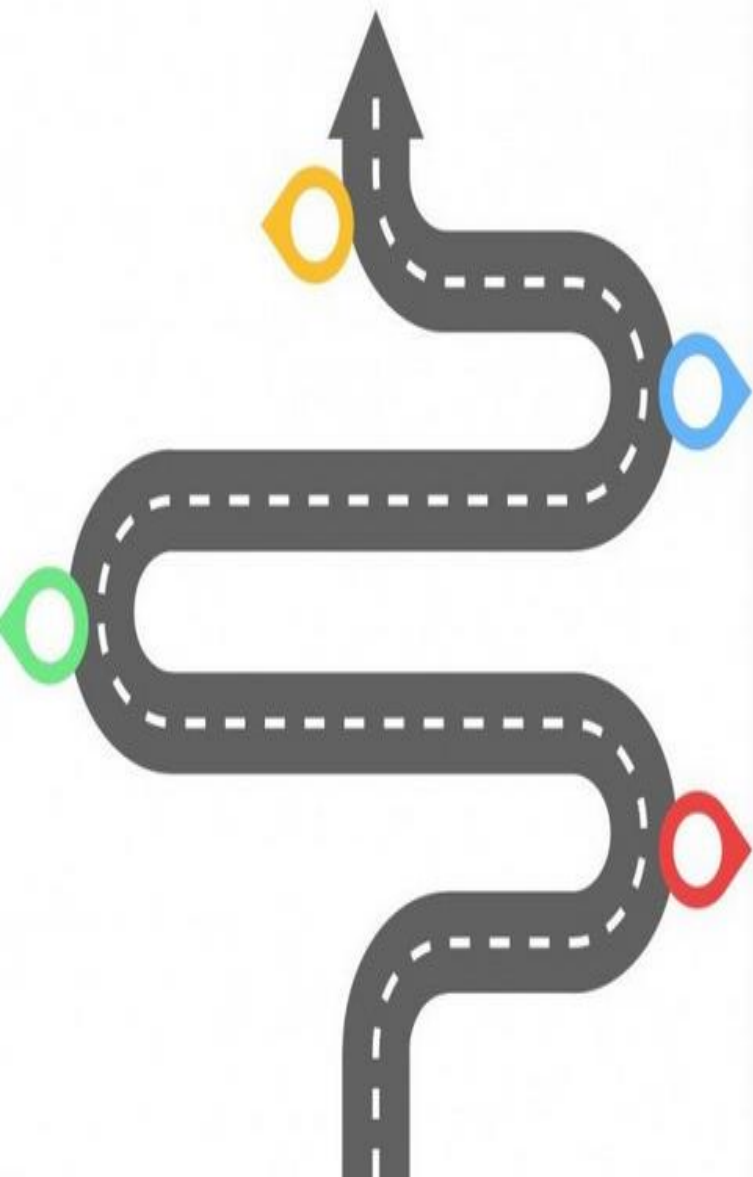
Conozca las posibilidades que ofrece VIVO, su adopción y aclare sus dudas acerca de este sistema.

El seminario web es **gratuito, en español y está abierto a todas las personas interesadas en asistir.**

Inscripción <https://bit.ly/vivoespanol>

LYRISIS SIGMA LA Referencia SENACYT

ICA UNA uc3m Universidad Carlos III de Madrid



Usability and Utility

- Dynamic API

Software evolution

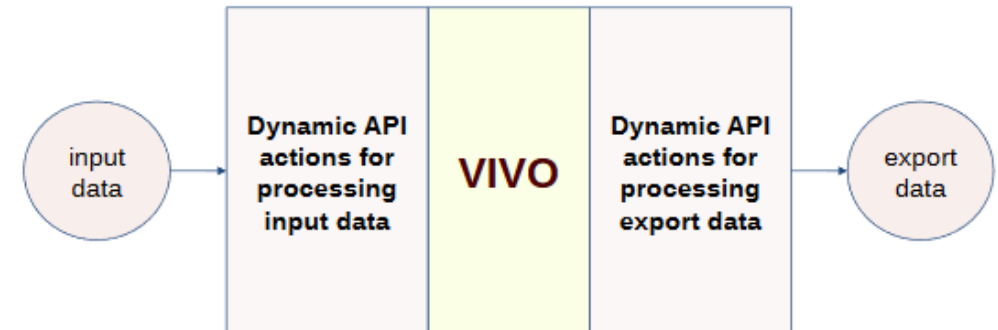
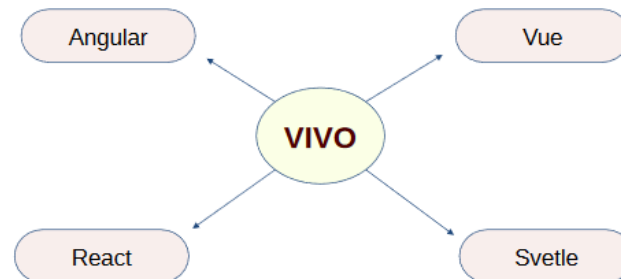
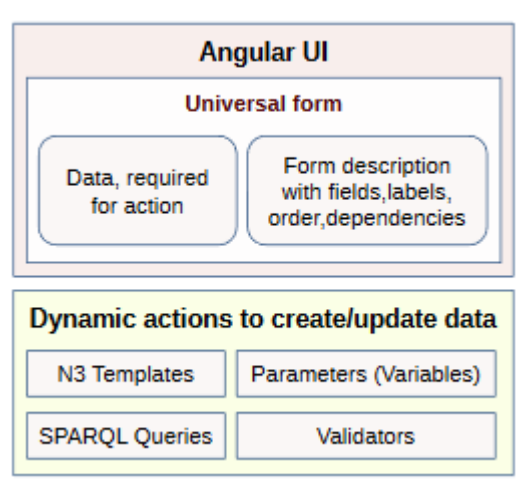
- VIVO 1.13.0 release.

Interoperability

- CERIF2VIVO mapping - Collaboration with EuroCris to align CERIF model to VIVO ontology. (ongoing)
<https://wiki.lyrasis.org/display/VIVO/Ontology+Interest+Group>
- Integrating Dspace and VIVO: (ongoing)
<https://wiki.lyrasis.org/display/VIVO/DSpace-VIVO+integration+task+force>
- Enhance interoperability with Fedora, & ORCID

Goal: Dynamic API would lead to decoupling frontend and backend, would enable easier customization of VIVO.

- Dynamic custom entry forms
- New web interfaces
- Better integration with external application.



A new **VIVO 1.13.0** has already been released for the community test

What's new:

- A new functionality to upload a file and link it with an individual in VIVO (for example, a book).
- Password authentication on external smtp servers
- New features for sparql update API
- Online translator editor, also enhancing Slavic languages.
- Deletion of individuals and related information
- Other minor improvements



Memorandum of Understanding

The purpose of this Memorandum of Understanding (MoU) is to establish and promote a strategic and cooperative partnership between VIVO and euroCRIS.

euroCRIS is a not-for-profit, statutory association (<https://www.eurocris.org>) established in 2002, governed by Dutch law and dedicated to the development and implementation of efficient and effective institutional, national and international research information systems and their interoperability, based on CERIF (Common European Research Information Format). One of euroCRIS's main objectives is the promotion of cooperation and exchange of expertise between stakeholders in the research information domain, in particular by setting up Strategic Partnerships with international organisations in the field of research information.

The VIVO community (<https://www.lyrasis.org/programs/Pages/VIVO.aspx>) is an open-source software community focused on the development of an open-source software platform of the same name and an ontology for representing scholarship. The VIVO community is one of eight community-supported software programs supported by LYRASIS that serve a wide-range of organization types, sizes and disciplines across the globe. LYRASIS is a 501 c 3 (US law) non-profit membership organization whose mission is to support enduring access to the world's shared academic, scientific and cultural heritage through leadership in open technologies, content services, digital solutions and collaboration with archives, libraries, museums and knowledge communities.

Motivation

Both VIVO and euroCRIS share the same vision of realising an optimal availability of and access to information on research through research information systems, for the benefit and support of all stakeholders involved in research. In this respect, standards and ontologies are key to recording information on research and scholarly activities in research information systems in an unambiguous manner. In addition, standards allow for the exchange of interlinked metadata on research in a

MOU with EuroCRIS

- **Align CERIF framework and VIVO ontology**
- **Collaborate on interoperability**
- **Communications between the communities**
- **Attend mutual conferences**

Takes advantage of mutual interests

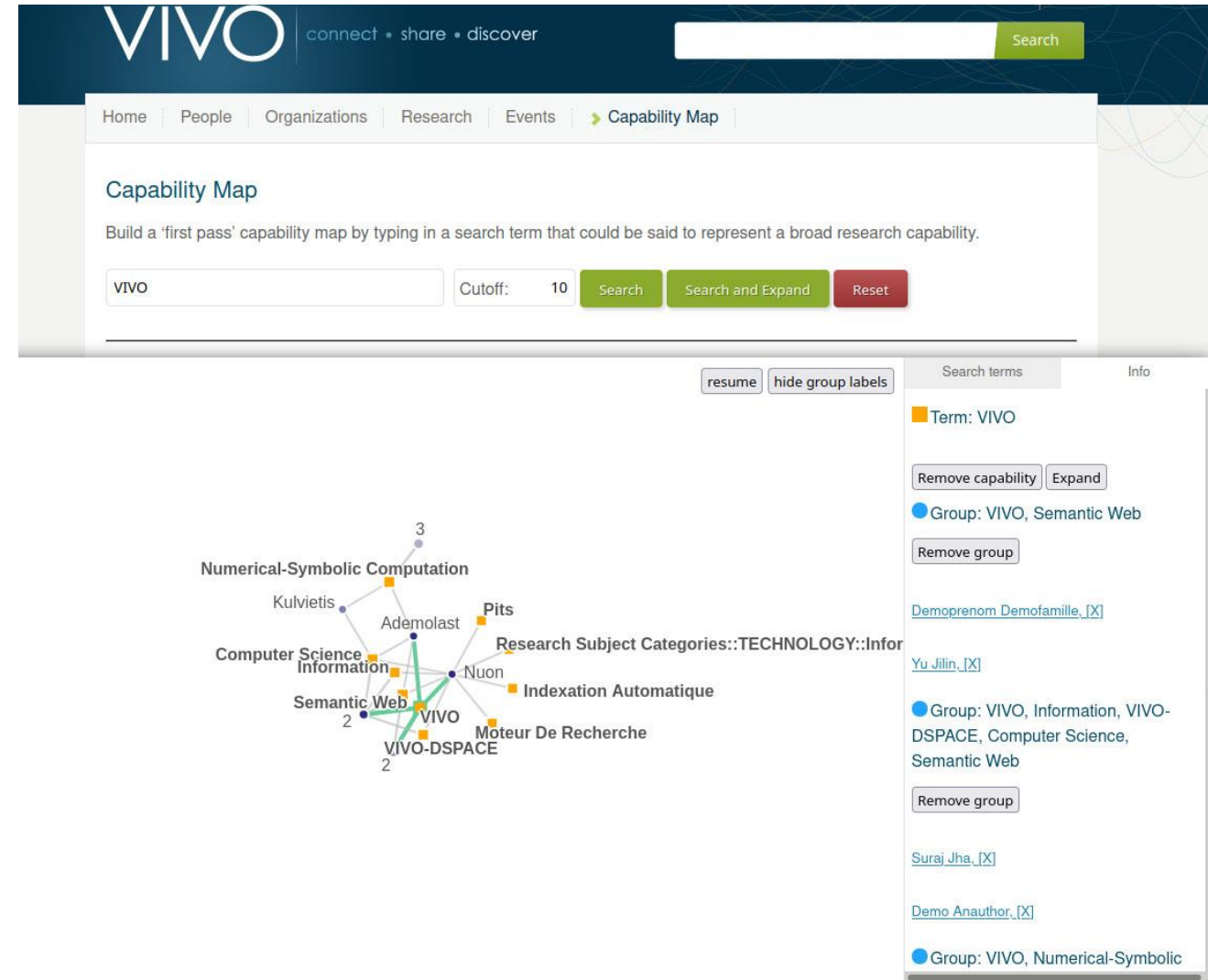
Benefits of the mapping:

- ✓ Interoperability between VIVO platforms and CERIF-compatible CRIS systems
- ✓ Knowledge transfer
- ✓ Improvement of CERIF model and VIVO ontology by analyzing the other side
- ✓ Extensions of the data models
- ✓ Addition of descriptions and annotations
- ✓ Machine-executable mapping for various purposes and in various notations, for example, for a CERIF-compliant data export from VIVO

VIVO Roadmap. Integrating DSpace and VIVO

Goal: consider using VIVO as a frontend for one or multiple DSpace instances at the institution

- A new presentation of DSpace items and semantic web aspect to existing DSpace repositories
- DSpace-VIVO migration assigns a unique ID to the researchers and subjects (keywords)
- The 'Capability Map' allows an expertise mapping across data sources



Project information:

<https://github.com/vivo-community/DSpace-VIVO>

<https://wiki.lyrasis.org/display/VIVO/DSpace-VIVO+Technical+Documentation>

The VIVO software



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CONNECT

VIVO creates a connected, integrated record of the scholarly work of your institution, ready for reporting, visualization, and analysis

DISCOVER

VIVO is developing world standards for big data representation of scholarly work

SHARE

VIVO provides a single shared vocabulary for data regarding scholarship

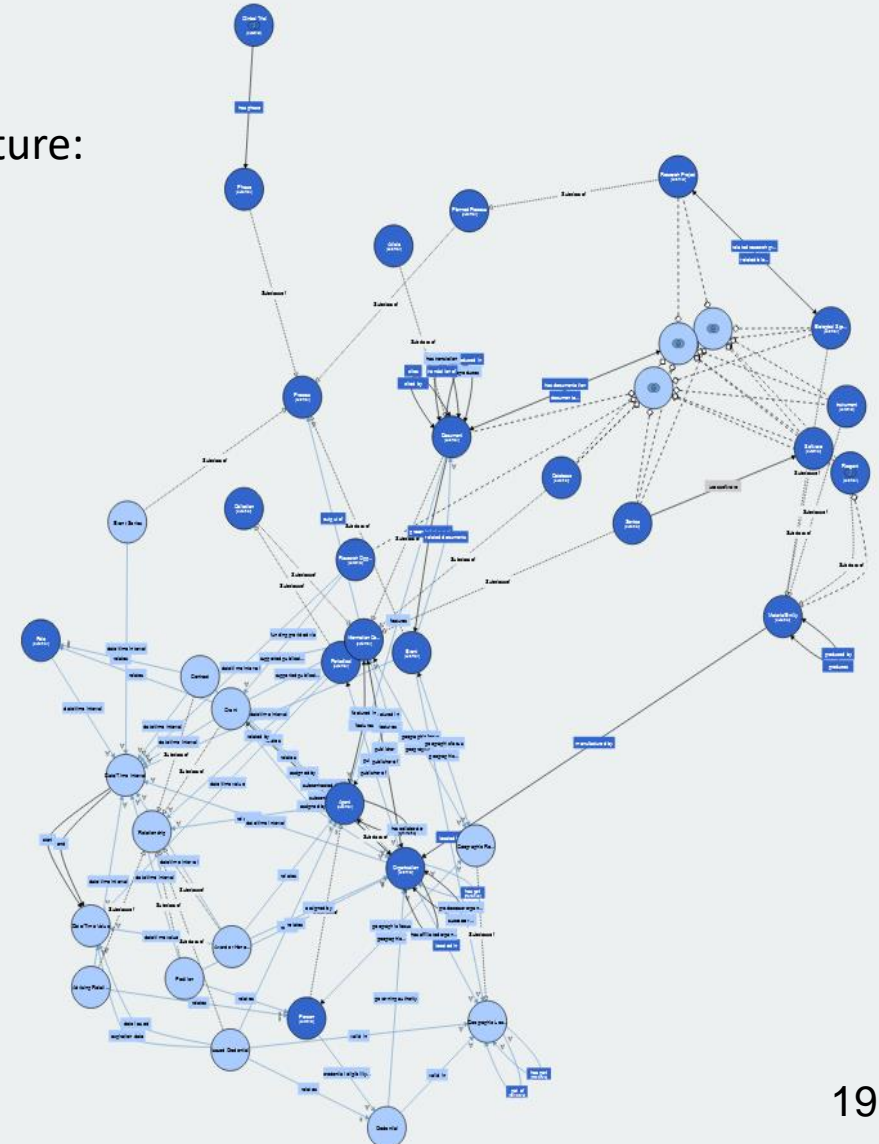
VIVO is an example of an application built entirely with [Semantic Web](#) technologies promoted by the [World Wide Web Consortium](#).

- Implements an Ontology based on standard international ontologies
- Stores data as [RDF](#) expressed in terms of [vocabularies called ontologies](#)
- provides [persistent URIs for data](#).
- Represents the expertise of people engaged in the creation, transmission, and preservation of knowledge and creative works.
- Contains FAIR data, complying with Linked Open Data Standards
- System requirements: VIVO may be hosted on one or more physical servers, on virtual servers, or in the cloud. Components:
 - Recommended installation (*): 4 cores x64 (min 2), 32 Gb RAM (min 2GB), 500 GB SSD (min 100 GB HDD)
 - OS Linux
 - TomCat Web application
 - MySQL database (with the default Jena SDB triple store)
 - Apache Solr search index.

Ontologies Used in the VIVO Ontology

The VIVO Ontology leverages the following ontologies in a unified, semantic structure:

- eagle-i Resource Ontology (ERO) – <http://www.obofoundry.org/ontology/ero.html>
- Basic Formal Ontology (BFO) – <http://www.obofoundry.org/ontology/bfo.html>
- Bibliographic Ontology (BIBO) – <http://bibliontology.com/>
- **Event** Ontology – <http://motools.sourceforge.net/event/event.html>
- Friend of a Friend (FOAF) – <http://www.foaf-project.org/>
- Gene Ontology (GO) – <http://obofoundry.org/ontology/go.html>
- [Geopolitical.owl](#), from the U.N. Food and Agriculture Organization
- Information Artifact Ontology (IAO) – <http://www.obofoundry.org/ontology/iao.html>
- Ontology for Biomedical Investigations (OBI) – <http://www.obofoundry.org/ontology/obi.html>
- Ontology of Clinical Research (OCRe) – <http://code.google.com/p/ontology-of-clinical-research/>
- Relations Ontology (RO) – <http://www.obofoundry.org/ontology/ro.html>
- Software Ontology (SWO) – <http://www.obofoundry.org/ontology/swo.html>
- **SKOS** (Simple Knowledge Organization System) – <http://www.w3.org/2004/02/skos/>
- **vCard** – <http://www.w3.org/TR/vcard-rdf/>
- SPAR ontologies, including FABIO, CiTO, and C4O: <https://purl.org/spar/fabio>



Ontology model. Main entities in VIVO

Fundamental:

- foaf:person
- Foaf:organization
- iao: informationContentEntity/vivo:document
- Skos: concept
- Event

With (important parts of linking them together):

- vivo:Relationship/vivo:Position
- bfo:Role
- bfo:Process

Access to the VIVO [Ontology](#) diagram

Some examples

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Weill Cornell Medical College

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Soo Jung Cho | Assistant Professor of Medicine

[Publications](#)
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[Contact](#)
[Other](#)

- Assistant Professor of Medicine, Medicine , Weill Cornell Medical College 2018 -
- Instructor in Medicine, Medicine , Weill Cornell Medical College 2016 - 2017
- Postdoctoral Associate in Medicine, Medicine , Weill Cornell Medical College 2014 - 2016

Phone

+1 646 962 2916

PUBLICATIONS

selected publications

Sort by Newest Co-Author Network

- Identification of Robust Protein Associations With COVID-19 Disease Based on Five Clinical Studies. *Frontiers in immunology*. 2022 Academic Article GET IT
- Antiviral Gene Expression in Young and Aged Murine Lung during H1N1 and H3N2. *International journal of molecular sciences*. 2021 Academic Article GET IT
- Association of circulating cell-free double-stranded DNA and metabolic derangements in idiopathic pulmonary fibrosis. *Thorax*. 2021 Academic Article GET IT
- Cytokine signatures of end organ injury in COVID-19. *Scientific reports*. 2021 Academic Article GET IT
- Decreased IDO1-dependent tryptophan metabolism in aged lung during influenza. *The European respiratory journal*. 2021 Academic Article GET IT

Times cited: 3

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James Abbruzzese

D. C. I. Distinguished Professor of Medical Oncology

My research interests include the clinical study and treatment of pancreatic cancer.

[Manage This Profile](#)
[Add Data to my Website](#)

Current Appointments & Affiliations

- D. C. I. Distinguished Professor of Medical Oncology, [Medicine, Medical Oncology, Medicine](#) 2018
- Professor of Medicine, [Medicine, Medical Oncology, Medicine](#) 2015
- Chief, Division of Medical Oncology, [Medicine, Medical Oncology, Medicine](#) 2013
- Member of the Duke Cancer Institute, [Duke Cancer Institute, Institutes and Centers](#) 2013

Contact Information

440 Mudd Building, Box 3406, Durham, NC 27710

440 Mudd Building, Box 3406, Durham, NC 27710

✉ james.abbruzzese@duke.edu

Background

- Education, Training, & Certifications
- Previous Appointments & Affiliations

Recognition

- In the News

Expertise


- Subject Headings

Research

Researchers@Brown

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Overview

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Curriculum Vitae [PDF]


Lynn Rothschild

Adjunct Professor of Molecular Biology, Cell Biology and Biochemistry

Overview

Prof. Rothschild is an astrobiologist/ synthetic biologist at NASA Ames specializing in molecular approaches to evolution, particularly in microbes, and the application of synthetic biology to NASA's mission. With a foundation in protistology and evolution, research interests include the early evolution of life, life in extreme environments and the search for life in the universe. In 2008 she established a program in synthetic biology for NASA and represented the Agency on the OSTP synthetic biology working group. Flight experience includes high altitude ballooning for astrobiology, the PI on the PowerCell payload on DLR's Eu:CROPIS satellite (launched December 2018), and Co-I on ESA's BIOMEX experiment on ISS. Extensive outreach including lectures worldwide, documentaries and a TEDx talk. Teaching experience includes "Astrobiology and Space Exploration", Stanford, 2004-13 (astrobiology.stanford.edu), directing theses (current Ph.D. students from Columbia, TU Delft, and UC Santa Cruz,), and the faculty advisor of the award-winning Brown-Stanford iGEM team. iGEM projects included synthetic biology for Mars Exploration (2011), Synthetic biology for astrobiology, including biomining (2012), Synthetic biocommunication (2013), Towards a Biodegradable UAS (2014), BiOrigami (2015) for which the team won "Best Manufacturing", and "BioBallooning" (2016) for which the team won "Best Measurement" and runner up for "Best Manufacturing", "Mars: getting there and staying there" (2017), Stanford-Brown-RISD iGEM team, "Mycos for Mars" (2018) for which the team was the runner up for the best new composite part and for best in manufacturing. In 2019 we joined forces with Princeton to form the Brown-Stanford-Princeton team, which won the iGEMers Prize. During the pandemic, Prof. Rothschild is supervising students remotely.

Brown Affiliations

 Molecular Biology, Cell Biology and Biochemistry

Research Areas

astrobiology | evolution | microbiology | protistology | space exploration | synthetic biology

On the Web

Profile, Motherboard/VICE Spring 2017

TED talk 2019, The living tech to support life on other planets

Isaac Asimov Award Lecture

NASA 360 podcast, Urban Biomining

NASA 360 podcast, Mycotecture

Wikipedia page

<https://vivo.brown.edu/>

TIB VIVO

Home

People

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Research

Events

Capability Map

Search for an Expert...

Search



Cahill, Brian

Positions

Mitarbeiter, Nachwuchsforschungsgruppe Learning and Skill Analytics , Programmbereich C - Forschung und Entwicklung 2020 -

Publications

Contact

Identity


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selected publications


academic article

Give and take on the MSCA: Programme for early career researchers aims to be more inclusive with less money. Research Europe. 2021

Researcher Mental Health and Well-being Manifesto 2021


The Impact of the COVID-19 Pandemic on the Working Conditions, Employment, Career Development and Well-Being of Refugee Researchers. Societies. 11:71. 2021  7

Four ways to fight science-funding cuts across Europe. Nature. 2020  159

Non-linearity and dynamics of low-voltage electrowetting and dewetting. Physical Chemistry Chemical Physics. 21:18290-18299. 2019  1

... more

article

Creating Research Environments that foster Mental Health and Wellbeing  8

Increasing Awareness of Researcher Mental Health

Who is responsible for transferable skills and how can RRI and Open Science help?

blog posting

How MSCA is changing under Horizon Europe. Research Professional Europe. 2021

chapter

Electrical Sensing in Segmented Flow Microfluidics. Micro-Segmented Flow. 73-100. 2014

Electrical Switching of Droplets and Fluid Segments. Micro-Segmented Flow. 31-54. 2014


Introduction. Micro-Segmented Flow. 1-3. 2014

Publications in VIVO

Co-author Network

Map of Science

Contact Info

 brian.cahill@tib.eu

<https://vivo.tib.eu/fis/>

23

■ Enhancing the research impact

Display the detailed information of a publication with bibliometric indicators for information and evaluation purposes.

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SANZ, FERRAN

Publications in VIVO
69 in the last 10 full years

Co-author Network

Co-investigator Network

Map of Science

Publications (256)

Book chapters (13)

Data Integration and Sharing Supporting Drug R&D. 436-443.

Educación asistida por ordenador en el ámbito sanitario

Evaluación de soluciones informáticas en biomedicina

Initial results on knowledge discovery and decision support for intracranial aneurysms. 265-272.

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DisGeNET: a comprehensive platform integrating information on human disease-associated genes and variants Journal articles

Authors Piñero J, Bravo A, Queralt-Rosinach N, Gutiérrez-Secristán, Deu-Pons J, Centeno E, García-García J, Sanz F, Furlong LI

Related people BRAVO SERRANO, ALEX Authorship
FURLONG, LAURA INÉS Authorship
SANZ, FERRAN Authorship

Subtype Investigation article

Journal title Nucleic Acids Research

Year of publication 2017

Volume 45

Number D1

Pages 833-839

ISSN 0305-1048

Abstract The information about the genetic basis of human diseases lies at the heart of precision medicine and drug discovery. However, to realize its full potential to support these goals, several problems, such as fragmentation, heterogeneity, availability and different conceptualization of the data must be overcome. To provide the community with a resource free of these hurdles, we have developed DisGeNET (<http://www.disgenet.org>), one of the largest available collections of genes and variants involved in human diseases. DisGeNET integrates data from expert curated repositories, GWAS catalogues, animal models and the scientific literature. DisGeNET data are homogeneously annotated with controlled vocabularies and community-driven ontologies (... more)

Bibliometric Indicators

Scopus 789 times cited

WEB OF SCIENCE 769 times cited

Scopus Sources Category, Rank and Percentile

SJR Index Scimago 9,025(2017)

Index H SJR 537(2020)

Quartile SJR Q1(2018)

Field SJR Genetics (Q1)(2018)

MIAR Link to Miar

Altmetric See more details
Picked up by 2 news outlets
Tweeted by 19
655 readers on Mendeley
3 readers on CiteULike

doi Publication link

Full text

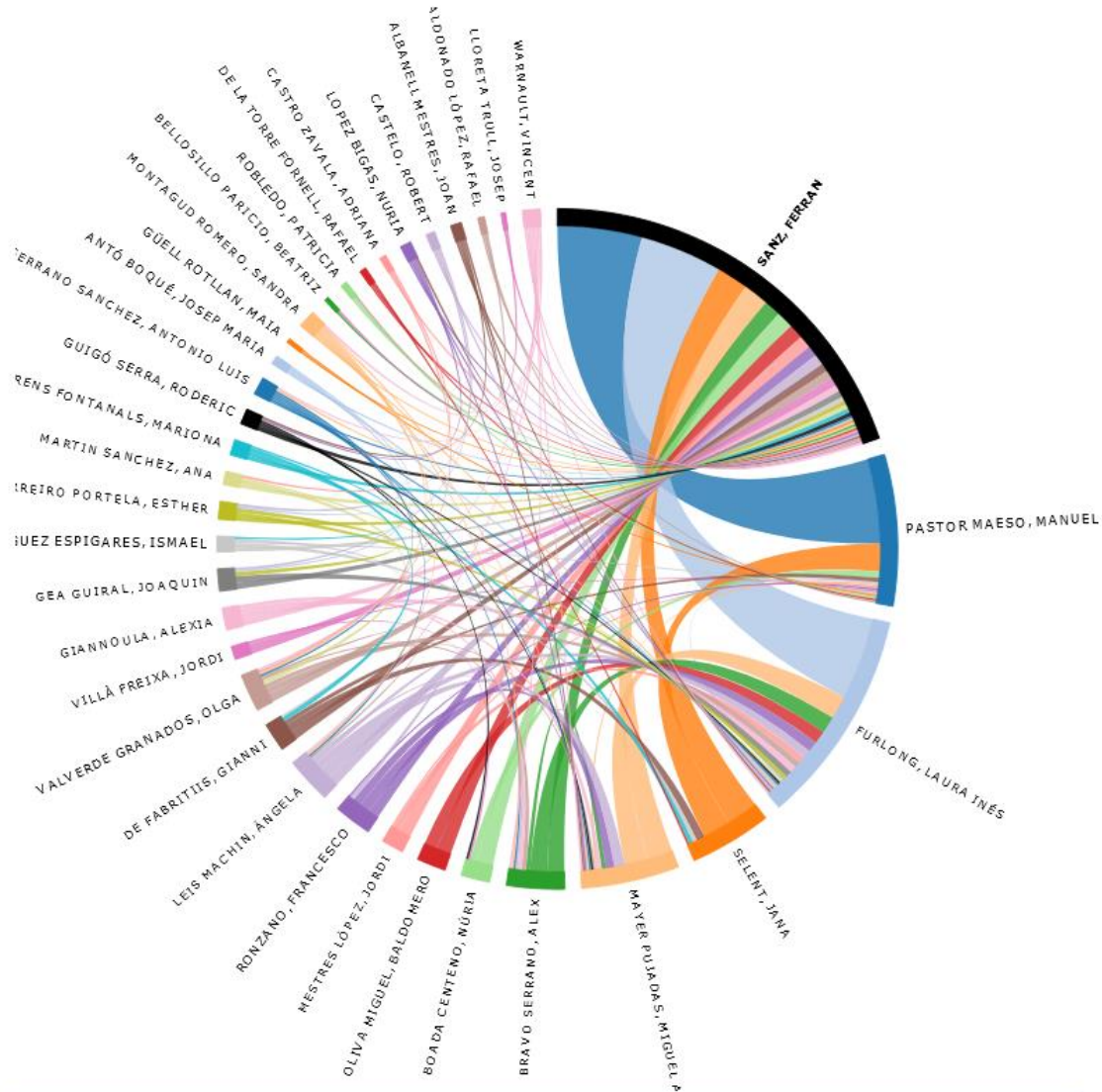
Citation Piñero J, Bravo A, Queralt-Rosinach N, Gutiérrez-Secristán, Deu-Pons J, Centeno E, García-García J, Sanz F, Furlong LI. DisGeNET: a comprehensive platform integrating information on human disease-associated genes and variants. Nucleic Acids Research 2017; 45(D1): 833 - 839.

Citation format Vancouver

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Co-author Network (GraphML file)



To boost collaborations



Tables

Publications per year (.CSV File)

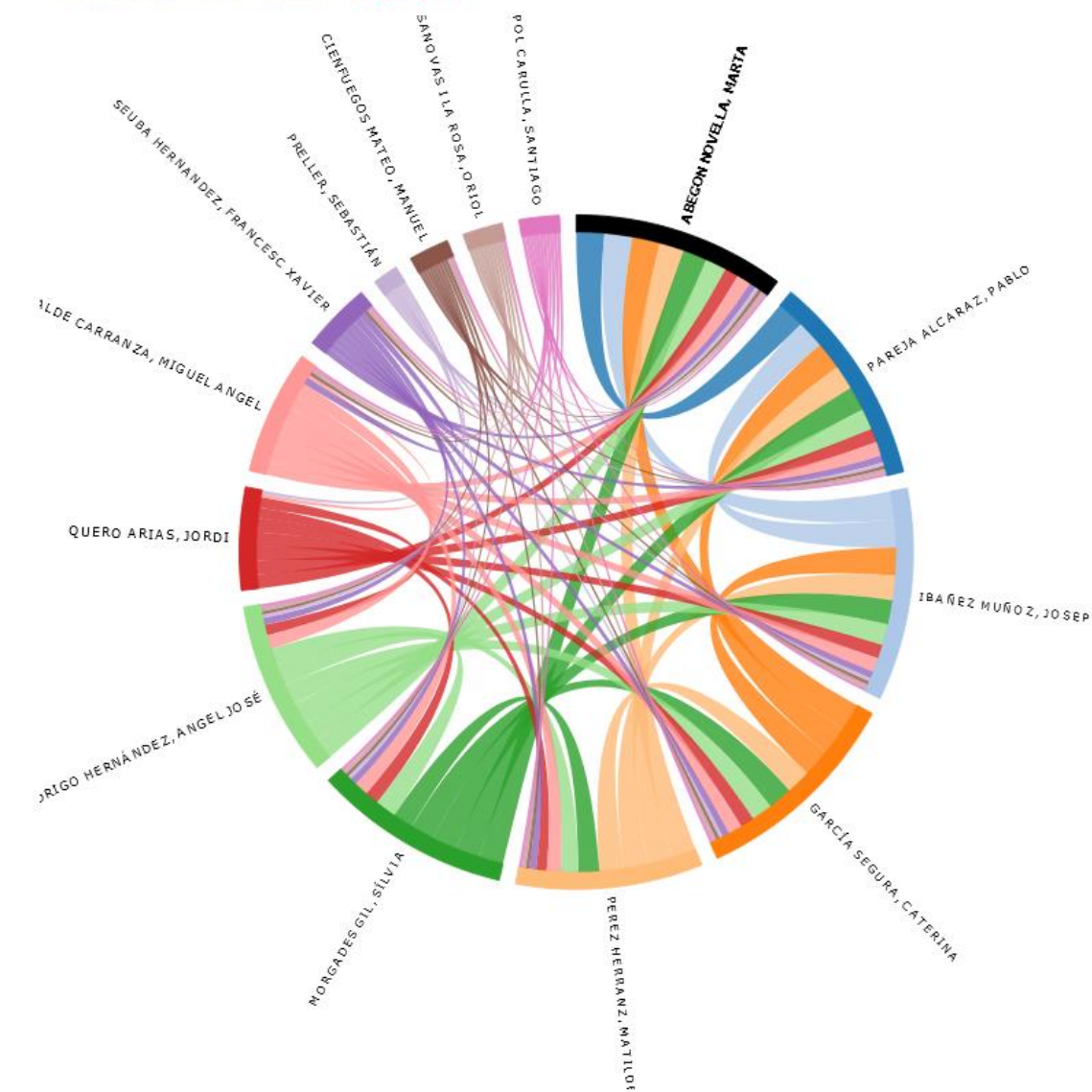
Year	Publications
1975	1
1977	2
1978	2
1979	1
1982	2
1983	4
1984	2
1986	4
1987	1
1988	8
1989	6
1990	5
1991	6
1992	4
1993	6
1994	8
1995	5
1996	8
1997	6
1998	4
1999	5

Co-authors (.CSV File)

Author	Publications with
PASTOR MAESO, MANUEL	40
FURLONG, LAURA INÉS	39
SELENT, JANA	15
MAYER PUJADAS, MIGUEL ANGEL	12
BRAVO SERRANO, ALEX	8
BOADA CENTENO, NÚRIA	8
OLIVA MIGUEL, BALDOMERO	7
MESTRES LÓPEZ, JORDI	6
RONZANO, FRANCESCO	5
LEIS MACHIN, ANGELA	5
VALVERDE GRANADOS, OLGA	4
DE FABRITIIS, GIANNI	4
VILLÀ PREIXA, JORDI	4
GEA GUIRAL, JOAQUIN	3
GIANNOULA, ALEXIA	3
GUIGÓ SERRA, RODERIC	2
BARREIRO PORTELA, ESTHER	2
RODRIGUEZ ESPIGARES, ISMAEL	2

Co-investigator Network

(GraphML File)



to boost collaborations



8 grants
from 2009 through 2021
(.CSV File)



13 co-investigators
from 2009 through 2021
(.CSV File)

Tables

The information in the following tables is for all years. [?](#)

Grants per year (.CSV File)

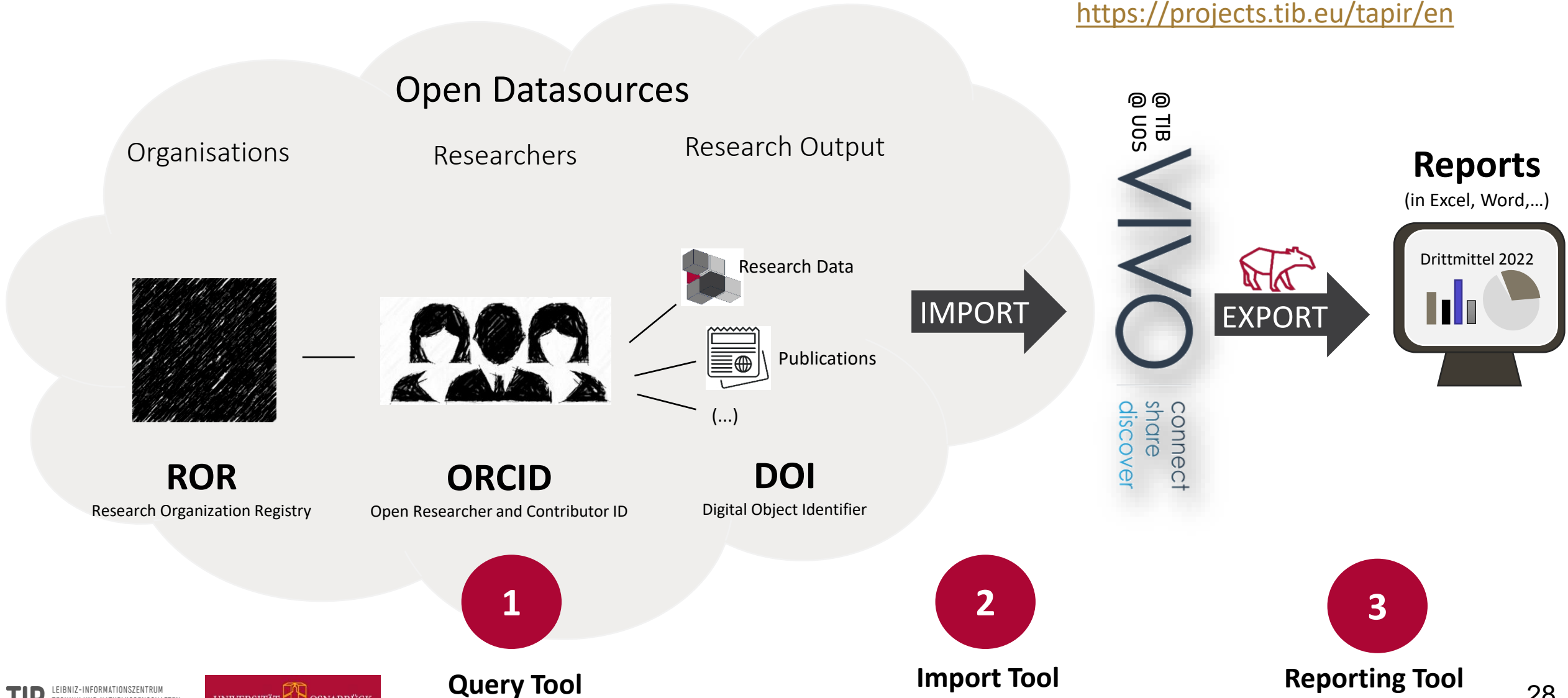
Year	Grants
2009	1
2012	2
2014	1
2015	1
2016	1
2017	1
2018	1

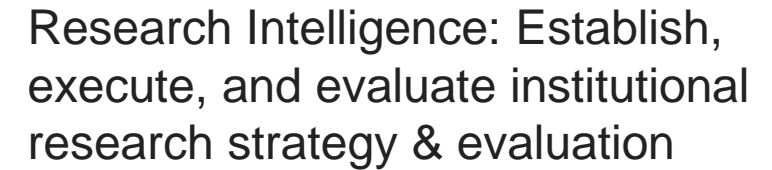
Co-investigator(s) (.CSV File)

Investigator	Grants with
PAREJA ALCARAZ, PABLO	8
IBAÑEZ MUÑOZ, JOSEP	8
GARCÍA SEGURA, CATERINA	8
PEREZ HERRANZ, MATILDE	7
MORGADES GIL, SÍLVIA	7
RODRIGO HERNÁNDEZ, ANGEL JOSÉ	6
QUERO ARIAS, JORDI	4
ELIZALDE CARRANZA, MIGUEL ANGEL	4
SEUBA HERNANDEZ, FRANCESC XAVIER	2
PRELLER, SEBASTIÁN	1
CIENFUEGOS MATEO, MANUEL	1
CASANOVAS I LA ROSA, ORIOL	1
RÍPOL CARULLA, SANTIAGO	1

Innovation in the VIVO Community

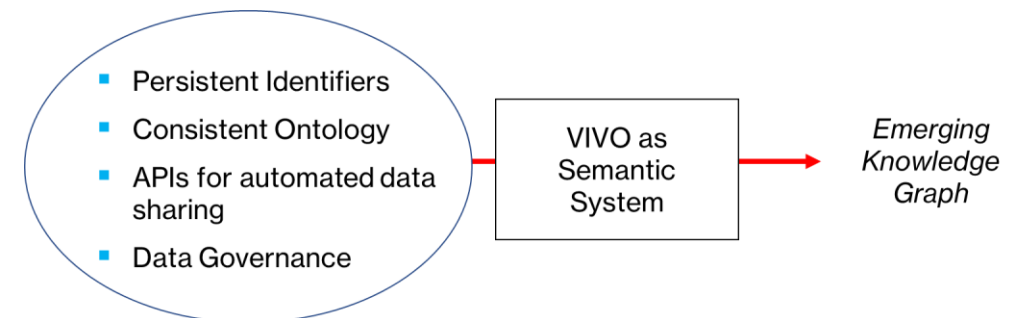
<https://projects.tib.eu/tapir/en>



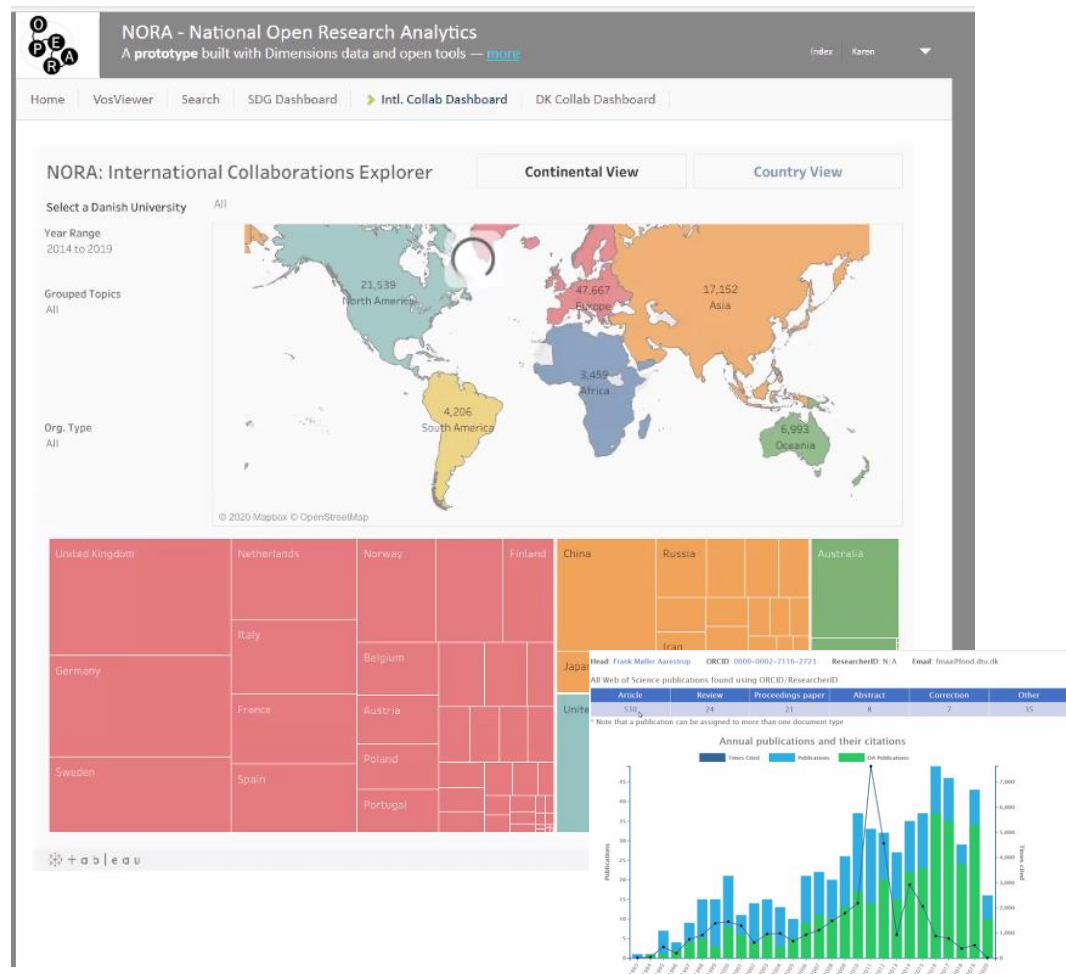


- Principle Investigators
- College and Department Program Reviews
- Vice President of Research

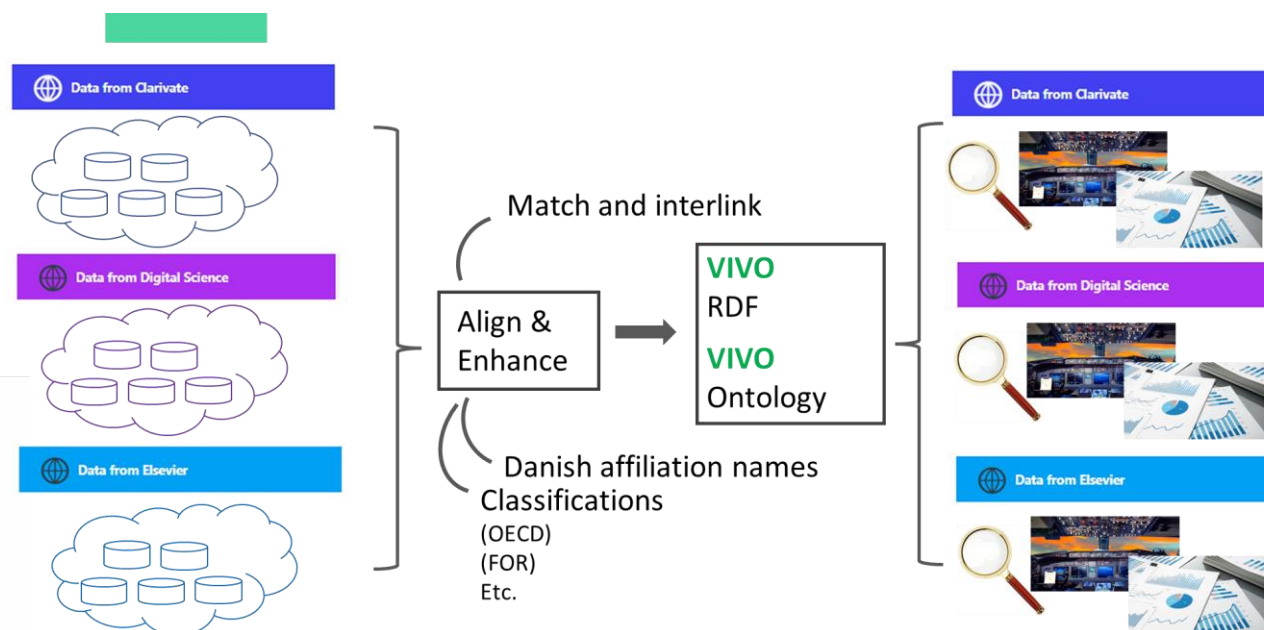
Takes advantage of linked data!



Research intelligence as emerging use case



Nora project from DTU – Denmark
Using the VIVO ontology to find research insights
(from Global datasources & Danish CRIS)



NORA, National Open Research Analytics, is a national initiative to enable robust and open insights and analytics of Danish research. NORA is focused on national level insights, and thus NORA supplements rather than replaces existing institutional systems, offering deep and detailed insights at various levels inside the institution, and existing global databases and research intelligence systems, offering insights and advanced analytics at the global level.

National/regional Research Portals. BrCRIS

- **BrCris** - Information System on the Ecosystem of the Brazilian Scientific Research
- Aggregation of different national and international data bases.
- Lattes Platform (research profiles of more than 1.5 million Brazilian masters and doctors) and for Oasisbr (aggregation node of the LA Referencia Network).
- Data aggregation software used is LA Referencia Platform(*) (exports to VIVO, APIS and visualizations).
- Entities and relationships recommend by the OpenAIRE Guidelines for CRIS Managers (CERIF-based)

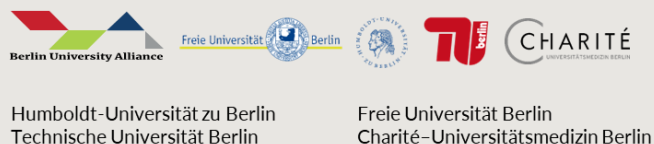



(*) Latin American open access science repository network.


National/regional Research Portals. BUA platform


- One research information platform for 3 universities + Charité with VIVO


VIVO-based Research Information Platform for the Berlin University Alliance



 Current Runtime: 05/21 - 12/23

 Current Data Sources:
3 Clusters of Excellence of the BUA

 Fadwa Alshawaf, Ph.D.
Claudia Adam, Florian Kotschka,
Rolf Guescini, Malte Dreyer

 Funded by the Federal Ministry of Education and Research (BMBF)
and the state of Berlin under the Excellence Strategy of the
Federal Government and the Länder.

GOAL

Creating a platform for a structured and transparent presentation of research information using semantic web technology

CHANCES

- Connect researchers to their work across disciplines and institutions
- Improve visibility and discoverability of expertise
- Facilitate new research collaborations across disciplines
- Quick overview through analyses and visualization

CHALLENGES

- Protection of personal data (GDPR)
- Quality of available data
- Lack of awareness of the importance of Metadata
- Creation of representative taxonomies & ontologies
- Using the correct semantics for all partners

How interoperable or aggregated CRIS can impact the country's policies for research and innovation?

- 1 - Mapping investment in Science & Technology versus Innovation results
- 2 - Creating a fairer Science evaluation system, in view of Open Science precepts
- 3 - Identifying over- or under-funded strategic areas
- 4 - Identifying and stimulating collaborative networks and the creation of research groups in thematic areas
- 5 - Generating access to open scientific information, scientific publications, research data and innovative products
- 6 - Connecting the entire scientific ecosystem, allowing quick visualisation of complex variables, generating information for decision-makers

- VIVO has a great community behind it, has strengthened its governance and is working on a roadmap that will allow it to evolve in line with new trends, focused on open Science and data sharing.
- It is working on fostering partnerships with relevant organizations with which important collaborations can be made.
- The versatility and adaptability of the tool and the advantages offered by an ontology based on international standards that provides linked open data, are highlighted.
- It has important ongoing projects led by a great group of developers, coordinated by a technical leader.
- There are innovative projects in the community that offer VIVO-based solutions focused on research intelligence and knowledge graphs.
- There is a clear trend to use VIVO as a research portal at local or national level, as an aggregator of data from different RIM/CRIS systems, to provide relevant information to governments, for decision making or policy definition.

Interested in learning more?

Get involved in the VIVO community!

- Visit vivoweb.org
- Read more at wiki.duraspace.org
- Follow @vivocollab on Twitter
- Join VIVO mailing lists
- Email anna.guillaumet@sigmaaie.org or beherbert@tamu.edu



The top half of the image features a dark blue background with a blurred image of a laptop screen. The word "VIVO" is written in large, white, sans-serif capital letters. To the right of "VIVO", separated by a thin vertical white line, are the words "connect", "share", and "discover" stacked vertically in a smaller, white, sans-serif font.

VIVO

connect
share
discover

Thank you very much!