OpenAIRE Explore

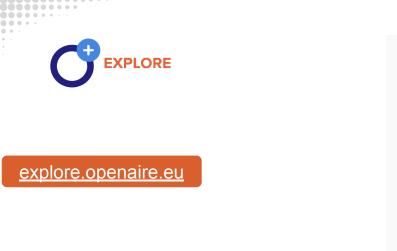
Konstantina Galouni

Service Manager kgalouni@athenarc.gr

October 5th 2023 | webinar



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Search in OpenAIRE

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A comprehensive and open dataset of research information covering 179m publications, 59m research data, 380k research software items, from 129k data sources, linked to 3m grants and 195k organizations. All linked together through citations and semantics.

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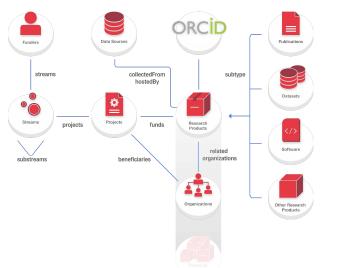
Access and "explore" the OpenAIRE Graph, its entities and their relationships





A graph is made up of nodes which are connected by edges







A massive collection of **metadata** and **links** between scientific products such as **articles**, **datasets**, **software**, and other research products, entities like **organisations**, **funders**, **funding streams**, **projects**, **communities**, **and data sources**.

The most unique benefit to the semantic graph database approach is the ability to infer or understand the meaning of information.



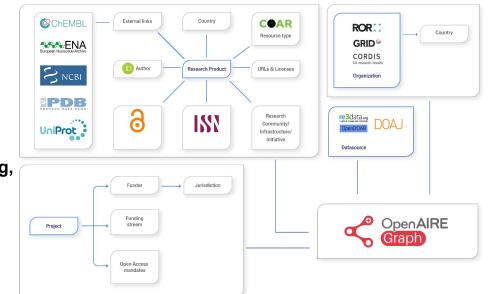
The OpenAIRE Graph **aggregates** millions of metadata records collected from **trusted data sources**, including:

- Open Access journals registered in DOAJ
- Crossref
- Unpaywall
- ORCID
- Microsoft Academic Graph
- Datacite

And **repositories** registered in **OpenDOAR**, **re3data.org**, **FAIRSharing.org**, **and the EOSC Service Catalogue**. Among these, prominent repositories such as:

- UKPubMed
- ArXiv
- HAL
- Zenodo
- Figshare
- Dryad
- Repec







Why OpenAIRE Graph?

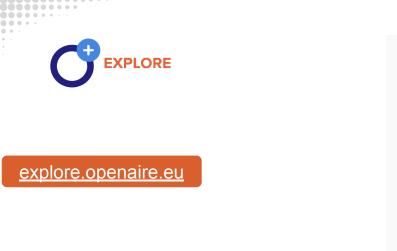
graph.openaire.eu

- Open and transparent
- Decentralized and interoperable
- Intelligent linking
- Embedded metrics

- **181M** publications
- 62M research data
- 348K research software items
- from **134K** data sources
- linked to **3M** grants and
- 200K organizations

All linked together through citations and semantics.





| Discover open |
|-----------------|
| linked research |

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Access and "explore" the OpenAIRE Graph, its entities and their relationships





Explore basic functionalities

Search

Link

Deposit

Search & filter

Download search results & reports

Link OpenAIRE entities

Link with external results

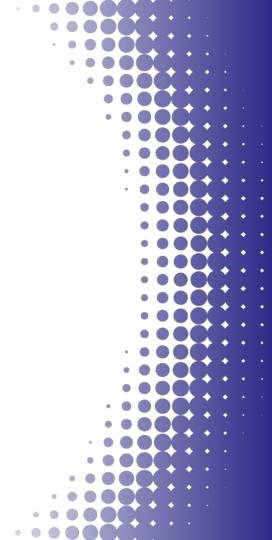
Enhance the information graph

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Search

Simple and advanced

keyword search, persistent • identifiers

research products, projects, ٠ data sources, organizations

Filters

Customizable response

Sorting, number of results •

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| Other research products | | 10.5281/zenodo.51440 ¹⁰ , 10.5281/zenodo.53077 ¹⁰ , 10.5281/zenodo.1321718 ¹⁰ , |
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| | Becker, Amelie; Loeden, Aenne; 👩 Manghi, Paolo; Principe | a, Padro; +1 Authors |
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Search

Simple and advanced

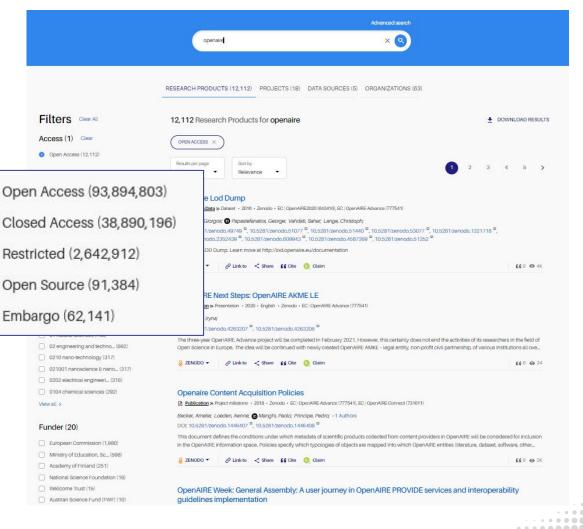
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Filters

Customizable response

• Sorting, number of results

Downloadable search results



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Search

Simple and advanced

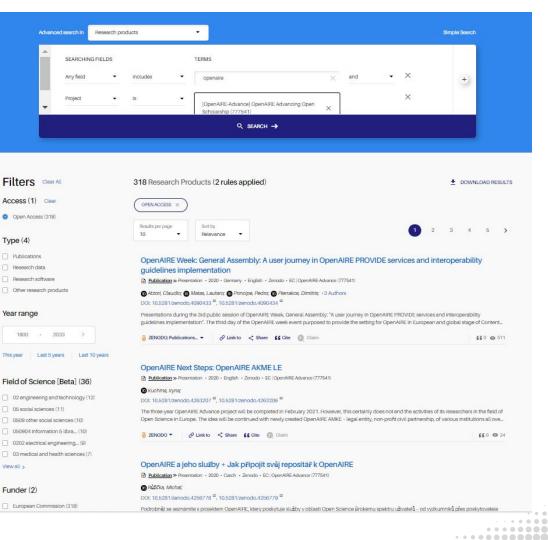
- keyword search, persistent identifiers
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Detailed pages - Research product

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| Relationships | Ouesta H Karlsson; Celine Schelcher; Elizabeth Verrall; Carlo Petosa; Alison J Sinclair; DOI: 10.1371/journal.ppat.1000005 [©] PMID: 18369464 [©] PMC: PMC:2267006 [©] | Beta Suggest SDGs Suggest Image: Suggest state stat |
| Statistics & | Summary Subjects References (26) Related research (12) External Databases (2) Metrics | Beta View all & suggest Fields of Science (5) View all & suggest medical and health sciences clinical medicine |
| Metrics & impact-based indicators | Abstract Epstein-Barr virus (EBV) causes infectious mononucleosis and is associated with various malignancies, including Burkitt's lymphoma and nasopharyngeal carcinoma. Like all herpesviruses, the EBV life cycle alternates between latency and lytic replication. During latency, the viral genome is largely silenced by host- driven methylation of CpG motifs and, in the switch to the lytic cycle, this epigenetic silencing is overturned. A key event is the activation of the viral BRLF1 gene by the immediate-early protein Zta. Zta is a bZIP transcription factor that preferentially binds to specific response elements (ZRES) in the BRLF1 promoter (Rp) when these elements are methylated. Zta's ability to trigger lytic cycle activation is severely compromised when a cysteine residue in its bZIP domain is mutated to serine (C189S), but the molecular basis for this effect is unknown. Here we show that the C189S mutant is defective for activating Rp in a Burkitt's lymphoma cell line. The mutant is compromised both in vitro and in vitro and in vitro and in vitro for binding two methylated ZREs in Rn (ZRE2 and ZRE3) although the effect is striking only for ZRE3. Molecular C Powered by the OpenAIRE Graph | Funded by WT |

Actions

Detailed pages - Research product

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| University of Sussex University of Wisconsin-Madison University of Wisconsin-Madison | | ner, Elizabeth Verrall; Carlo Petosa; Alison J Sinclair; 15 [©] PMID: 18369464 [©] PMC: PMC2267006 [©] | | Beta SUggest SDGs Suggest Image: Control of the second se |
| Nated Organizations University of Wisconsin-Oshkosh Urnhed States | Effect of PCMBS on 1998 • Inferred by OpenAIRE • Is | CO2permeability of Xenopus oocytes expressing aquaporin 1 or its C189S AmongTopNSimilarDocuments <u>aino Acid Residue in the Basic Region of the Epstein Barr Virus (FRV) 1 vt</u> 18 • PLoS Pathogens, volume 4, page e 1000005 (eissn: 1553 <i>7</i> /14, Copyright policy ⁽²⁾) • Public Libra | tic Cycle Switch Protein 7ta (BZI E1) | Citations 34 Popularity AVERAGE Influence AVERAGE Influence AVERAGE Impulse AVERAGE Downloads 22 |
| | 12 Research Products Replacing C189 in t 2018 • Inferred by OpenAIRE • Is | he bZIP domain of Zta with S, T, V, or A changes DNA binding specificity to 1 | four types of double-stranded DNA. | ۹ الا |

Detailed pages - Data source

Tailored for each entity

Overview

Relationships

Statistics & Metrics & impact-based indicators

Actions

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| To update or not to update the ESCMID guidelines for the diagnosis and treatment of biofilm infections That is the qu | uestion! The opinion of the ESGB board | | Detailed information @ OpenDOAR [®] |
| Authors: Holby, N; Moser, C; Oliver, A; Williams, C; +4 Authors HANDLE: 182285451 ⁽²⁾ | RepositoriUM | | |
| Background The work on the ESGB guidelines for diagnosis and treatment of biofilm infections began in 2012 and the result was published in 2014. The guidelines have for people working with biofilm infections. At the ESGB Biofilm conference in Mailorca 2022 (Eurobiofilms2022) the board of the ESGB decided to evaluate the 2014 guidelines in the second s | - been and | arch products (547) Related Data sources Metrics | i, Statistics |
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| Optimization of a hybrid bacterial/Arabidopsis thaliana fatty acid synthase system II in Saccharomyces cerevisiae Debilication > Other literature type - 3203 - Forugal - English - Esewire B.V. | +Only top 100 Data sources that host Research products which are also | available via the Federated Research Data Repository are shown. | |
| Authors: Pozdniakova, T. A.; Cruz, Josio P.; Silva, Paulo César, Azevedo, Flávio; +4 Authors HANDLE: 1822/85220 [©] | 98 Data Sources, Page 1 of 10 | | |
| Faity acids are produced by eukanotes like baker's yeast Saccharomyces cerevisiae mainly using a large multifunctional type I faity acid synthase (FASI) where seven cat | DATA SOURCE | | RESEARCH PRODUCTS IN EXPLORE |
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Detailed pages - Project

Tailored for each entity

Overview

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Actions

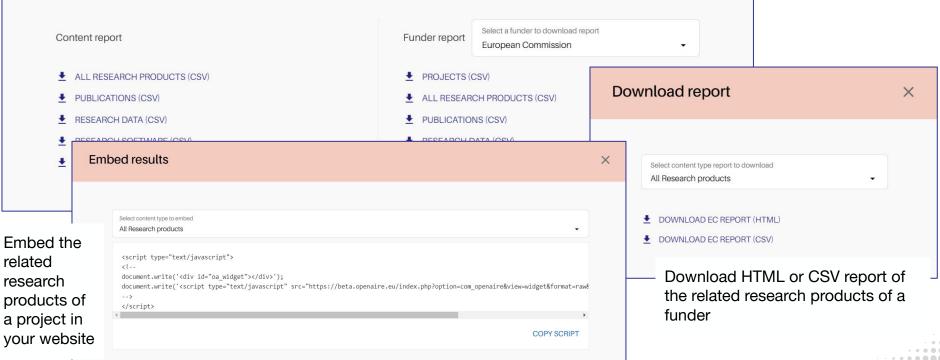
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| OpenAIRE-Advance OpenAIRE Advancing Open Scholarship | n 2018 (Started) - 28 Feb 2021 (Ended) + Closed |
| Funder: European Commission Project code: 777541 Call for propose | L: H2020-EINFRA 2017 |
| Funded under: H2020 RIA Overall Budget: 10,000,000 EUR Funder 0 | iontribution: 10,000,000 EUR |
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| Summary Publications (252) Research data (20) Research sof | Summary Publications (252) Research data (20) Research software (9) Other research products (51) DMPs (2) Metrics |
| | DATA MANAGEMENT PLANS |
| Description OpenAIRE Advance continues the mission of OpenAIRE to supp infrastructure, comprised of a human network and nobust techni | VIEW ALL IN SEARCH PAGE \rightarrow \mathfrak{S}^{2} start a New DMP in Argos $^{\boxtimes} \rightarrow$ |
| Its communities to Open Science, aiming to be a trusted e-In OpenAIRE-Advance strives to empower its National Open Ar infrastructures, positioningOA and open science onto national a | Title of DMP - demo |
| thematic areas(open policies, RDM, legal issues, TDM), promotin with support and training toolkits, training resources and w | Papadopoulou, Elli; |
| publishing and next generation repositories, to develop essent focuses on the operation and maintenance of the OpenAIRE | DOI: 10.5281/zenodo.4244637 ¹² , 10.5281/zenodo.4244636 ¹² |
| optimizing their performance and scalability, b) refining their | description |
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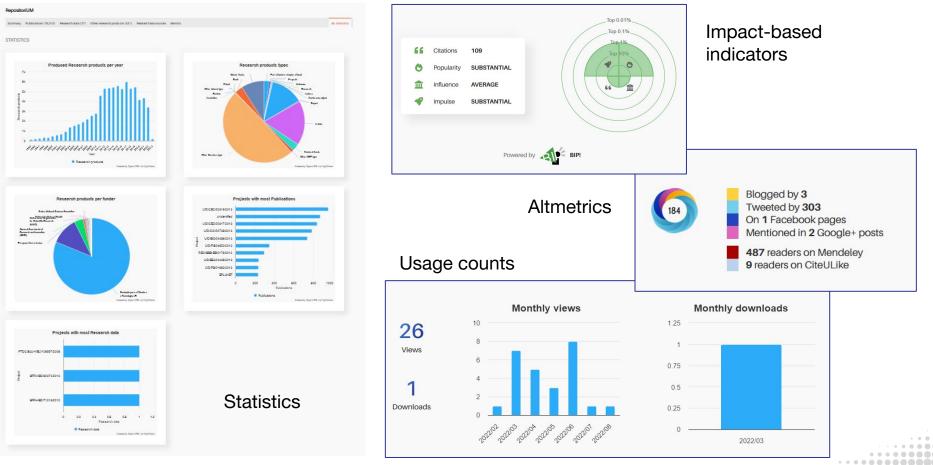
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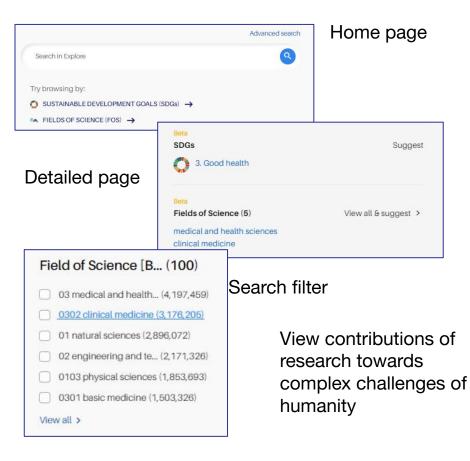


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Statistics & Metrics & Impact-based indicators



Fields of Science (FoS)



Beta Fields of Science .

Home A Fields Of Science

01 Natural Sciences

Sciences

02 Engineering And Technology

03 Medical And Health Sciences 04 Agricultural And Veterinary

OB Humanities And The Arts

We have integrated a held of Science (FoS) taxonomy into our dataset to organize and discover research more effectively. Using the full capabilities of the OpenAIRE Research Graph (full texts, citations, references, venues) we apply AI and bring forward any multidiscipiinarity potential.

Our work is based on the work from our partner Athena Research Center: SciNoBo A Hierarchical Multi-Label Classifier of Scientific Publications https://doi.org/10.1145/3487553.3524677



| 01 natural sciences | | |
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| 0101 mathematics 010101 Applied Mathematics 010102 General Mathematics 010103 Numerical & Computational Mathematics 010104 Statistics & Probability | 0102 computer and information sciences 010201 Computation Theory & Mathematics | O103 physical sciences 010301 Acoustics 010302 Applied Physics 010303 Attronom § Astrophysic 010306 Fluids & Plasma 010306 Granet Bhysics 010307 Mathematical Physics 010309 Mathematical Physics 010309 Copies |
| 0104 chemical sciences 010401 Analytical Chemistry | 0105 earth and related environmental sciences | 0106 biological sciences |
| 010402 General Chemistry | 010501 Environmental Sciences | 010602 Entomology |
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| | SDGs | Suggest |
| Detailed page | 3. Good health | |
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Linking

Users enhance the graph with relationships

- projects/funding
- communities
- research products

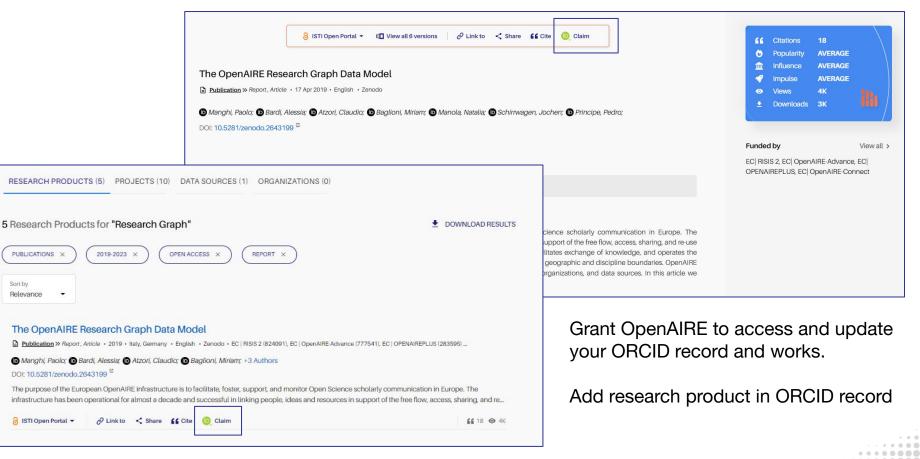
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4 Research Products, Page 1 of 1

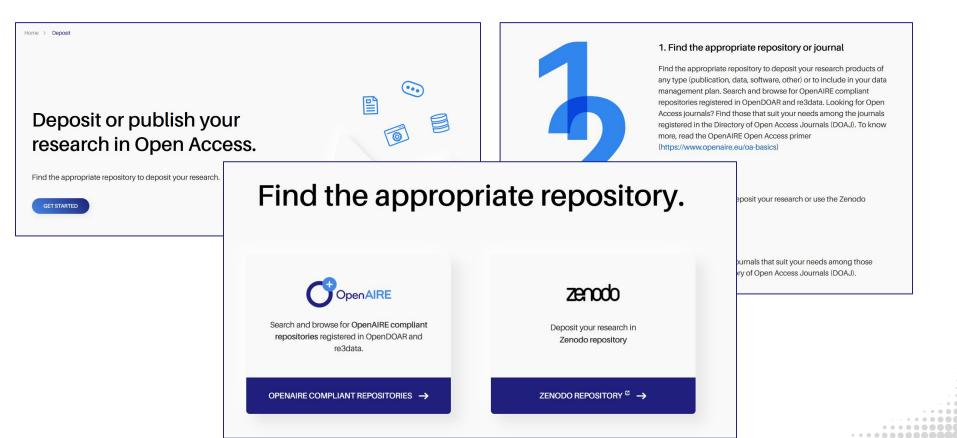
| Solvated protein-DNA docking using HADDOCK | ORCID (D ACTIONS |
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| Publication » Article · 2013 · Netherlands · Springer Science and Business Media LLC · EC EGHNSPIRE (261323), NWO Predicting, modelling and (2300135688), EC WENMR (261572) | VIEW ORCID WORK |
| Visscher K.M.; Bonvin A.M.J.J.; Kastritis P.L.; Van Dijk M.; | |
| DOI: 10.1007/s10858-013-9734-x [@] PMID: 23625455 [@] HANDLE: 1874/290382 [@] | |
| Interfacial water molecules play an important role in many aspects of protein-DNA specificity and recognition. Yet | C UPDATE ORCID WORK |
| they have been mostly neglected in the computational modeling of these complexes. We present here a solvated | |
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|) Journal (116,068) | Results per page 1 2 3 4 5 ≯ 10 |
| Institutional Repository (5,666) | |
| Data Repository (4,353) | Mammalian Transcriptomic Database |
|) Thematic Repository (402) | Data Repository • China (People's Republic of) • Not yet registered • NGDC, CNCB / BIG, CAS • Transcriptomics, Expression data, Gene expression |
| Journal Aggregator/Publisher (296) | re3data: r3d100012214 ^{E2} |
| Publication Repository (204) | The Mammalian Transcriptomic Database (MTD) was created to store information on mammalian transcriptomes. The MTD allows browsing of genes based on |
| /iew all > | their neighboring genomic coordinates or joint KEGG pathway and provides expression information on exons, transcripts and geness via a geneme browser. The |
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