



ELIXIR : Infrastructure for life science data

Niklas Blomberg

ELIXIR Greece Launch



www.elixir-europe.org

Data challenges & Opportunities

- Geographically distributed data production
- Secure access and governance of human data
- Open data mandates of National and European funders

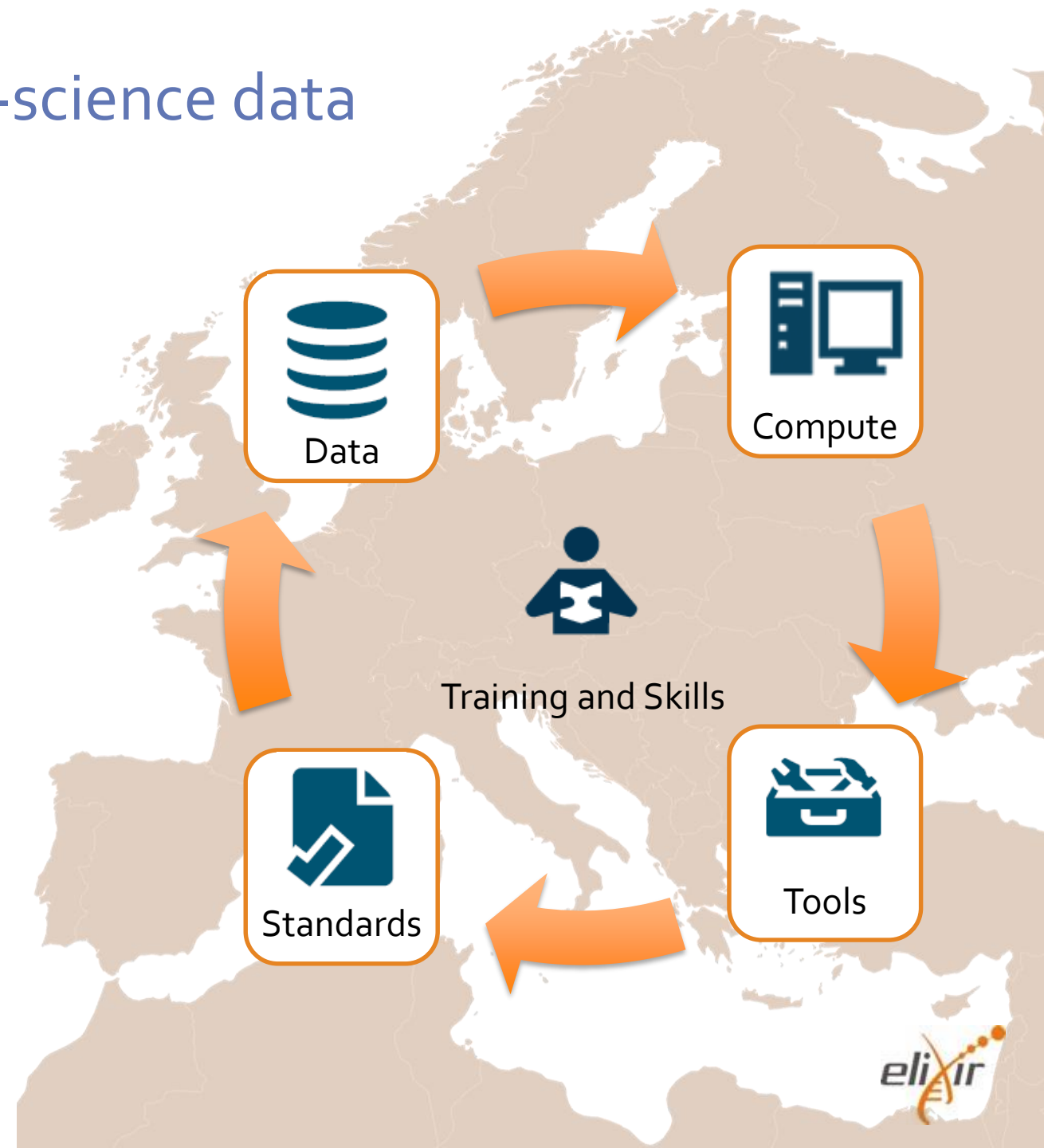


wellcome trust

European Infrastructure for life-science data

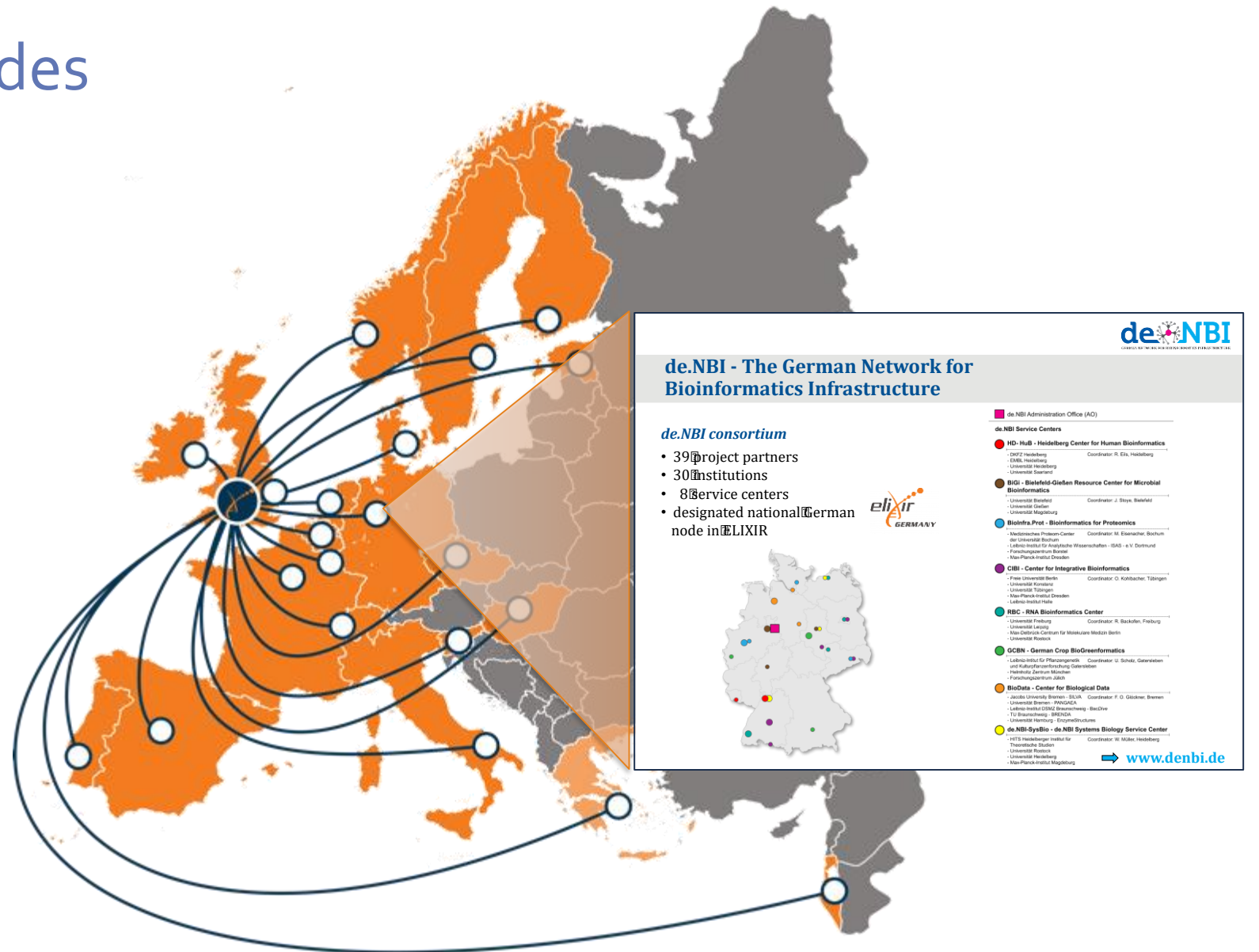


...delivered in partnership
with research communities

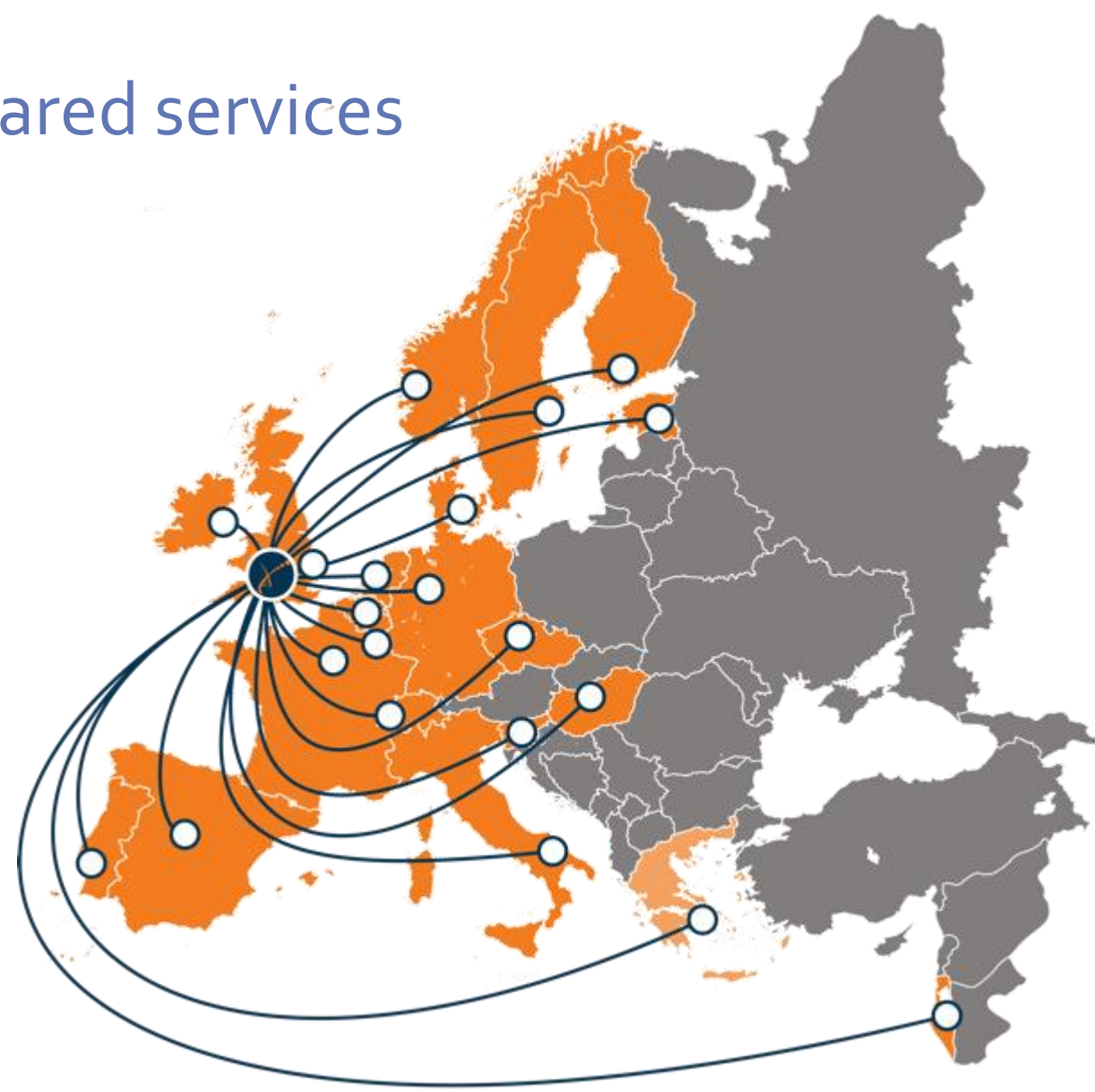


A network of data Nodes

- **ELIXIR** Nodes are funded nationally
- **ELIXIR** Nodes build on national strengths and priorities
- **ELIXIR** Nodes provides a national framework for long-term resource management



Distributed infrastructure with shared services



ELIXIR's Platforms of shared services

- Accessed internationally
- Operated by Nodes
- Funded by national and international schemes
- Connected through ELIXIR's technical programme

ELIXIR



Technical Platforms that connect **Compute, Data, Tools, Interoperability** and **Training** services between Nodes

ELIXIR Communities that connect domain experts between Nodes



Plant sciences

- Mission: facilitate genotype-phenotype analyses for crop and tree species
- Actions:
 - Develop **standards** for representation of genotypic and phenotypic data
 - Make data discoverable and interoperable through **common APIs**
 - Annotate and submit **key exemplar datasets** to relevant public archives
 - Develop reusable **modules for visualisations**
 - **Disseminate best practices** and tools to national projects



ELIXIR partners at leading plant research centres



MIAPPE

Minimum Information for

Plant phenotyping data techniques. While various phenotypes, descriptions are standardized. A basic discovery and data management development to address

MIAPPE is a Minimum necessary to fully describe all of the elements listed in the checklist, and consult characteristics, i.e. when

Please follow the links

- The current version
- The MIAPPE opinion
- A recent paper describing information required
- The current implementation
- More information at [Study-Assay file for](#)
- An [archive of previous](#)
- A mapping of MIAPPE Nucleotide Archive
- The [MIAPPE GitHub](#)
- A full list of Minimum
- Contact us as at [inf](#)

BrAPI Overview

INTRODUCTION

The Breeding API specifies a standard interface for plant phenotype/genotype databases to serve their data to crop breeding applications. It is a shared, open API, to be used by all data providers and data consumers who wish to participate. Initiated in May 2014, it is currently in an actively developing state, so now is the time for potential participants to help shape the specifications to ensure their needs are addressed. The listserv for discussions and announcements is at <http://mail2.sgn.cornell.edu/cgi-bin/mailman/listinfo/plant-breeding-api>. Additional documentation is in the [Github wiki](#).

URL structure

API requests are structured as "`<server>/brapi/v1/`", where "v1" is the version number of the API, followed by the command.

Example: `/brapi/v1/markerprofiles/2939`

To distinguish between multiple databases or crops available from the same server, include the database or crop name as part of the "`<server>`" identifier. An arbitrary number of levels can be inserted between the domain name and the crops or brapi level, if needed.

Example: `superBreedingServer.org/maize/brapi/v1/markerprofiles/2939`

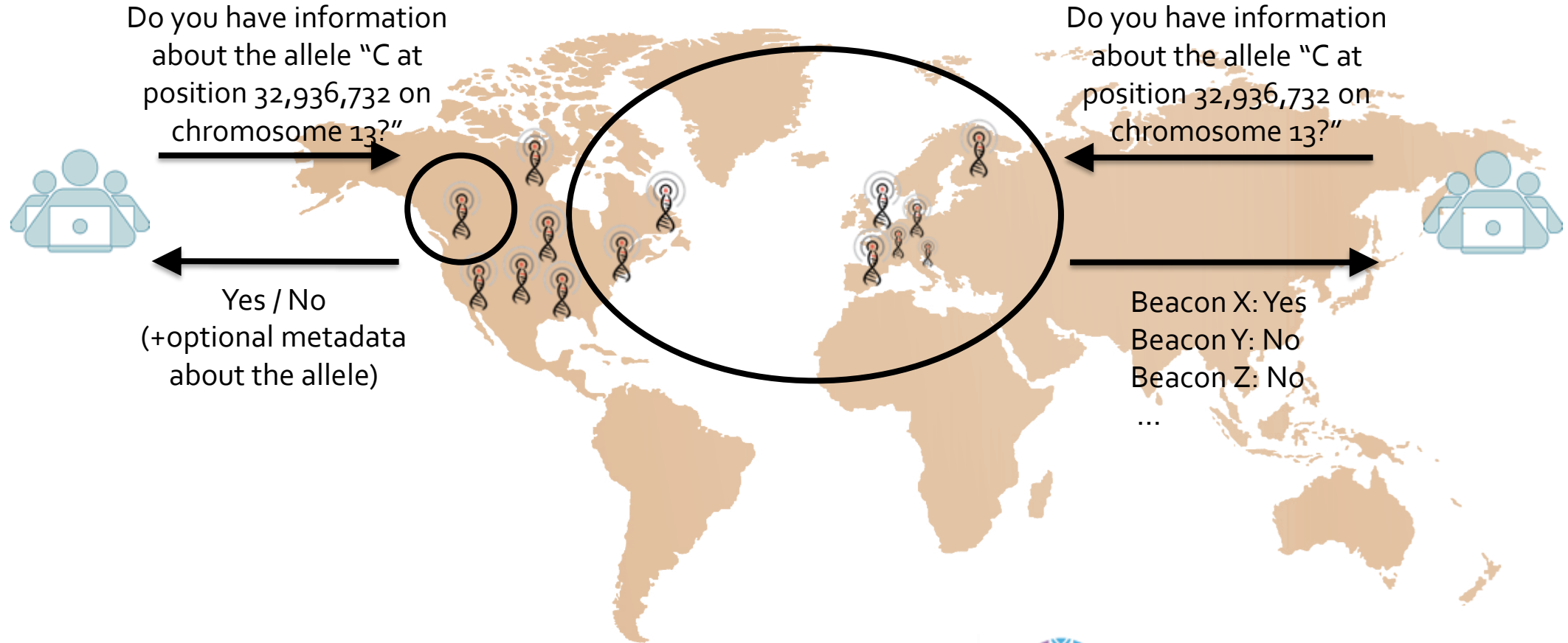
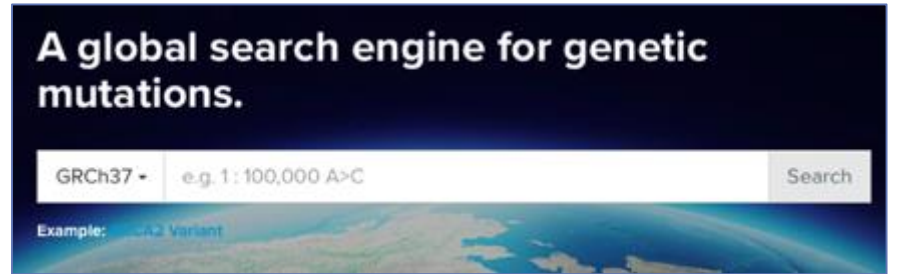
Structure of the response object:

The return objects are encoded in JSON. The response always consists of a "metadata" key that minimally contains the pagination information in a "pagination" key and the status information as well as a "datafiles" key that lists URLs to data files generated by the call.

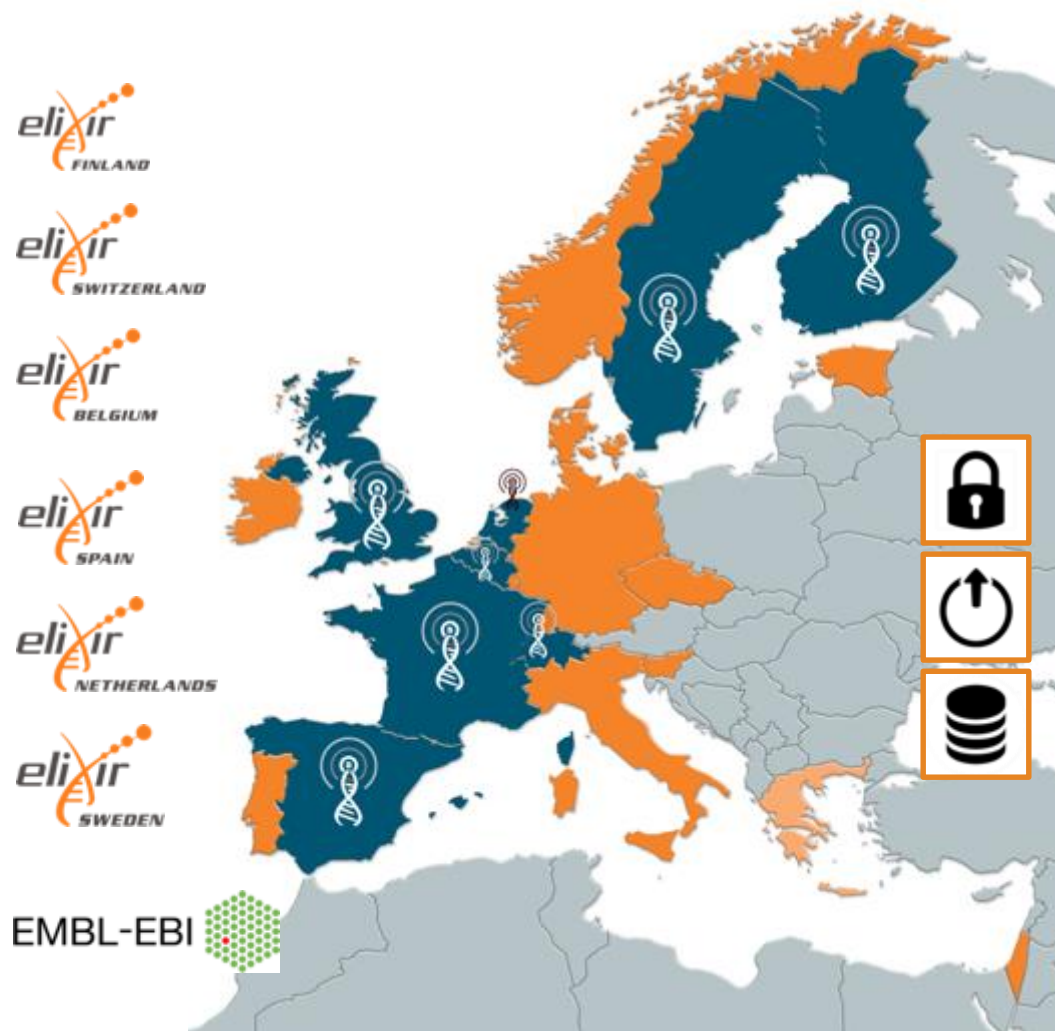
If the response is a single record that doesn't require pagination, then the value for the "pagination" key is returned with all the keys set to zero. When the results are paginated, the keys "pageSize",



Public data discovery web-service



ELIXIR and Beacons



- Funding to drive implementation of the Beacon technology within ELIXIR nodes
- ELIXIR Authentication and Authorization Infrastructure

Public	Accessible to internet users
Registered	Accessible to bona fide researchers
Controlled	Authorized – signed agreement needed

www.elixir-europe.org/beacons



Global Alliance
for Genomics & Health
Collaborate. Innovate. Accelerate.



Federation of human genome data: *localEGA*

- Many national datasets from human research participants needs to be stored locally
- ELIXIR-EXCELERATE developing “localEGA” – shared metadata (FAIR) and local data store (secure)
- Linking local EGA to national clouds – and international access (ELIXIR-AAI)



How do you find a needle in a federated haystack?





Bioschemas

*"schema.org markup
for life sciences –
minimum properties
needed for finding
data"*

<http://bioschemas.org>

The screenshot shows a Google search for "Apple pie". The search results include several recipe cards from sources like Pillsbury.com, BBC Good Food, and SimplyRecipes.com. Each card includes a small image of the pie, a title, a rating, and a brief description. The top result is from Pillsbury.com, titled "Perfect Apple Pie recipe from Pillsbury.com".

On the right side of the search results, there is a detailed "Apple pie" entry. It includes a "Nutrition Facts" table for 100 grams of the pie.

Nutrition Facts	
Amount Per 100 grams -	
Calories 237	
Total Fat 11 g	18%
Saturated fat 3.8 g	15%
Polyunsaturated fat 2.2 g	
Monounsaturated fat 4.4 g	
Cholesterol 0 mg	0%
Sodium 266 mg	11%
Potassium 65 mg	1%
Total Carbohydrate 34 g	11%
Dietary fiber 1.6 g	6%
Protein 1.9 g	3%
Vitamin A 2%	Vitamin C 5%
Calcium 1%	Iron 6%
Vitamin B-6 0%	Vitamin B-12 0%
Magnesium 1%	

Below the nutrition facts, there is a section titled "People also search for" with images and links for "Apple", "Apple cake", "Dessert", "Tart", and "Sugar".

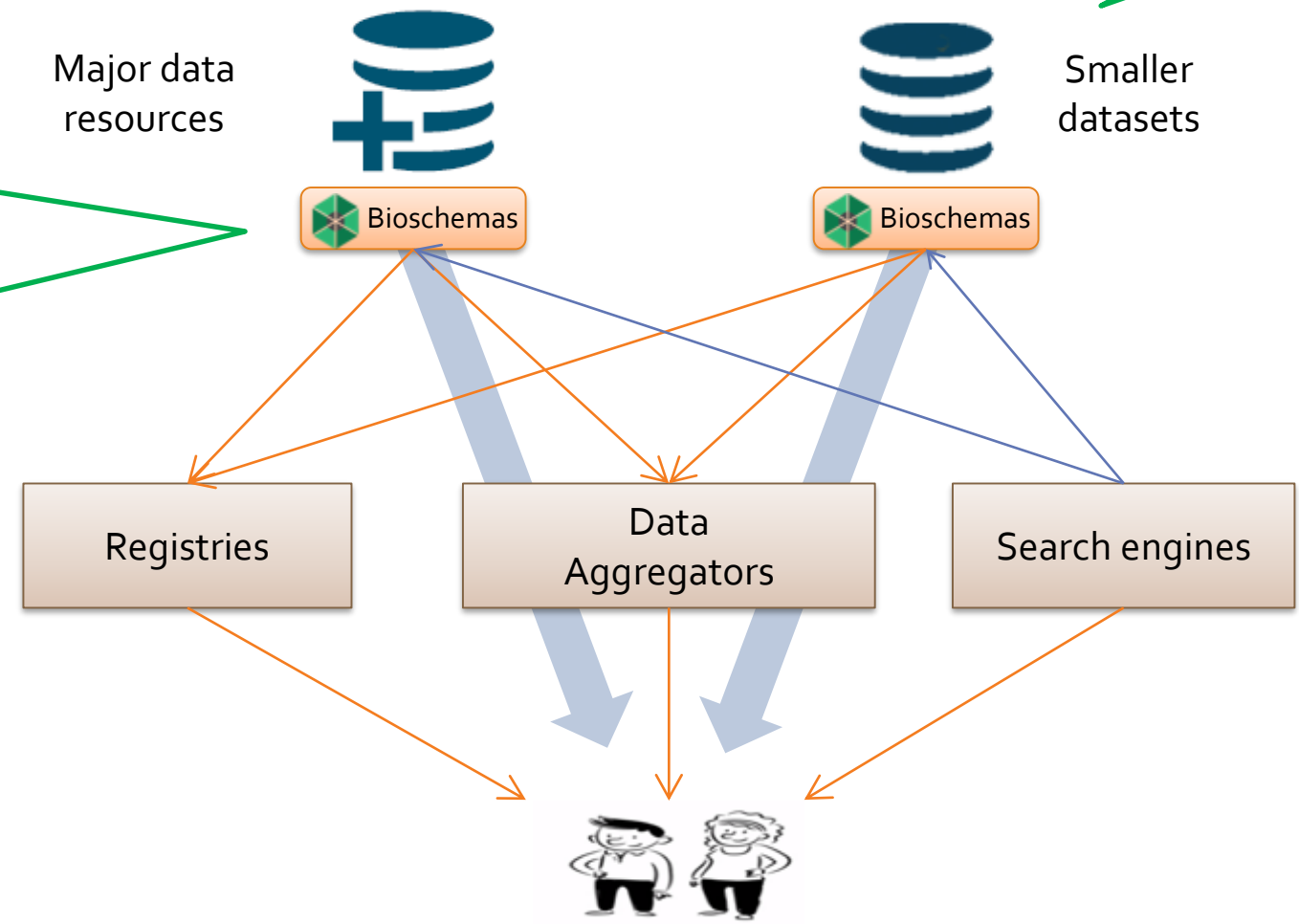


Bioschemas.org

A community initiative built on top of Schemas.org to improve **Findability** and **Accessibility** in Life Sciences

- Rapid markup
- Exposed to harvesting
- Find

- Standardised metadata
- Metadata publish and harvest without APIs or special feeds
- Feed bio registries and aggregators





Bioschemas adopted in EOSC

In November we presented early adopters...

now **16 live deployments!**

- Generic standard for data discovery
- "Research schemas" as Emerging federation architecture in EOSC (Part of EDMI)
- Created by the Bioschemas community, funded by the ELIXIR Budget

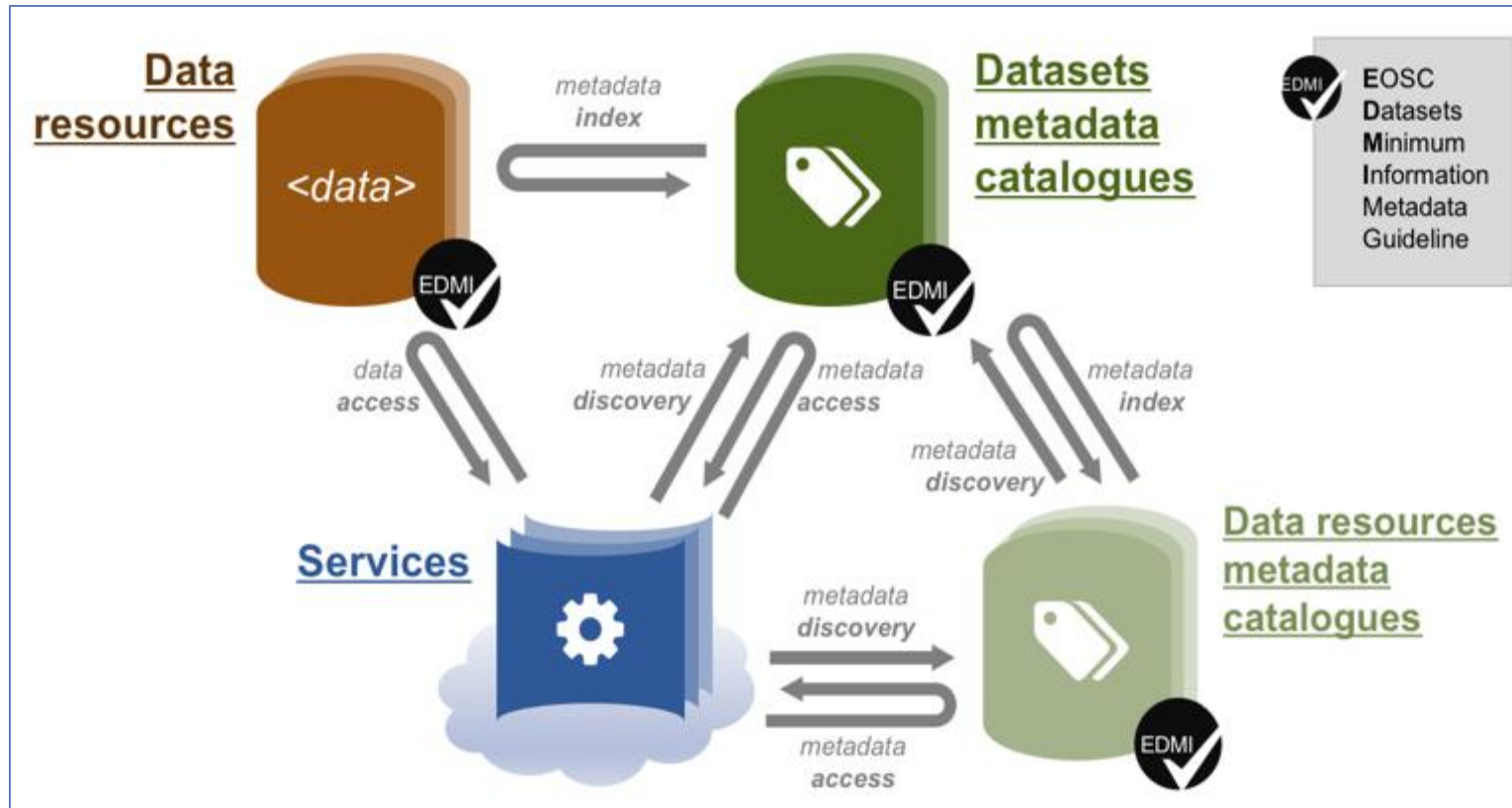


<http://bioschemas.org/liveDeploys/>

Services/sites implementing Bioschemas's markup

Name	Bioschema Profile	Profile Version	Structured Data Testing Tool
Identifiers	DataCatalog	0.1	visualise
Fairsharing	DataCatalog	0.1	visualise
Gigadb	DataCatalog	0.1	visualise
Human Protein Atlas	DataCatalog	0.1	visualise
EGA	DataCatalog	0.1	visualise
Isaexplorer	DataCatalog	0.1	visualise
IUPHAR/BPS	DataCatalog	0.1	visualise
MobiDB	DataCatalog	0.1	visualise
EGA Dataset	Dataset	0.1	visualise
MobiDB	Dataset	0.2	visualise
DataMed	Dataset	0.1	visualise
Biosamples	Sample	0.1	visualise
Pscan	Tool	0.1	visualise
PscanChIP	Tool	0.1	visualise
Cscan	Tool	0.1	visualise
BAR 3.0	Tool	0.1	visualise

European Open Science Cloud is build around domain registries



ELIXIR: Gateway for User access and mechanism for exposing life-science services (via *ELIXIR Registries*)

ELIXIR Authentication and Authorization Infrastructure AAI

Reliable electronic identification of users (ELIXIR ID) is needed to access the key services and capacities of ELIXIR.

- **You can link existing user accounts to create your ELIXIR ID today at www.elixir-europe.org**
ELIXIR AAI allows Users to continue using their federated academic, corporate or social media identity by linking it to a personal ELIXIR ID.
- The ELIXIR service providers connected to ELIXIR AAI benefit from a centralised user identity and access management services.
- Protocols SAML2, OpenIDConnect.
- <https://www.elixir-europe.org/services/compute/aai>



- 359 Home Organisation IdPs enabled for login (via eduGAIN)
- 987 ELIXIR users
- 155 groups created in ELIXIR AAI
- 61 registered Resource Providers

ELIXIR Cloud & Compute

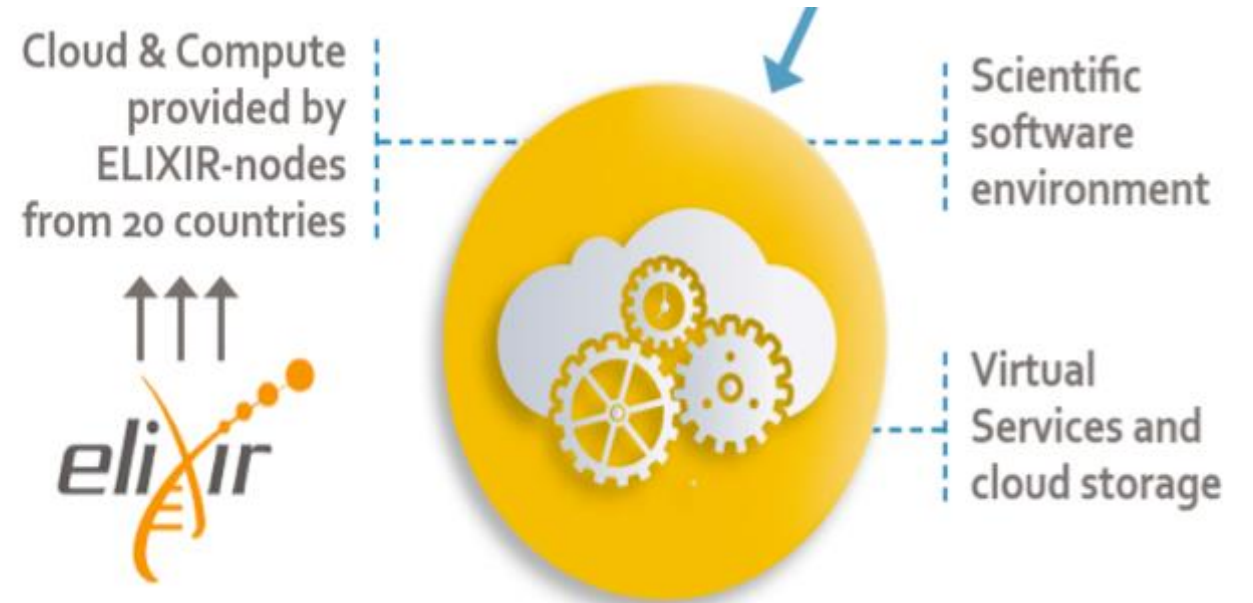
ELIXIR Cloud capacities surveyed [here](#)

DK, DE, EBI, FI, FR, CH confirmed capacity


> 60.000 compute cores

> 24.000 TB of storage

> 3.000 compute users





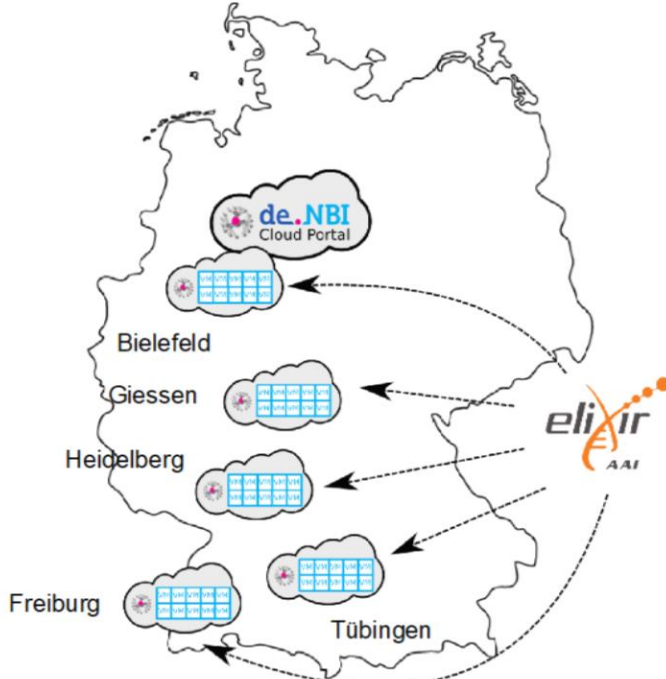
AAI as foundation for Transnational cloud access



de.NBI Cloud introduction

- Started 2016
- 5 compute centers
- >10000 Cores
- >5 PB Storage Capacity
- >100 TB RAM
- In collaboration with



Single Sign-On



23.10.2017 7

Name	Description	Elixir Node
Data Archiving	Elixir Luxembourg provides integrated storage and archiving for curated Translational Medicine data. Data is stored on transSMART servers or other suitable servers.	Elixir Luxembourg
AAI - REMS	CSC develops and hosts an open source tool REMS to assist a DAC to manage Data Access Applications and access rights to sensitive datasets. REMS is a key component in Elixir AAI.	Elixir Finland
Authentication and Authorisation Infrastructure (AAI)	The Elixir AAI (Authentication and Authorisation Infrastructure) is the Elixir service portfolio for authenticating users and helping relying services to manage users' access rights in the services.	Elixir Czech Republic, Elixir Finland
Computations	Computome is the Danish National Supercomputer for Life Sciences. It serves all life science research groups within Denmark and is also open for international collaboration.	Elixir Denmark
Consulting/Expertise	Expertise in very diverse subjects of data processing, analysis and curation as well as connections to existing infrastructure RI's like ELIOT. This expertise can be made available to research projects and organisations.	Elixir Netherlands
CSC Chipster	Chipster is a user-friendly analysis software for high-throughput data. It contains hundreds of analysis tools for next generation sequencing (NGS), microarrays, proteomics and sequence data.	Elixir Finland
CSC Cloud	The CSC Cloud service is targeted for High-Performance Computing (HPC), allowing customers to run virtual machines with exclusive access to up to 16 cores. This service includes Cloud and ePouta.	Elixir Finland
de.NBI cloud	The upcoming de.NBI cloud will provide an analytics infrastructure for bioinformatics. It will consist of computing power and storage capacity as well as flexible workflows and analysis tools.	Elixir Germany
Embassy Cloud	This is an OpenStack platform co-located with EMBL-EBI's services and data resources. Access to the Embassy Cloud is available for researchers outside EMBL if they have a collaboration with staff at EMBL.	EMBL, EBI
Help-Desk	Under Development in collaboration with SURFcon and Legature, based on prior experience deploying tools for end users / life scientists.	Elixir Netherlands
OpenConnect	Open source software underlying the technology for SURF's single sign-on and facilitating national and international online collaboration in education and research.	Elixir Netherlands
uSCORE	uSCORE provides a high-performance computing infrastructure, large-scale storage resources, scientific software and databases, server infrastructures and user support. It also provides expertise to scientific research groups.	Elixir Switzerland
Vital-IT	Vital-IT supports and collaborates with life scientists in Switzerland and beyond. It provides expertise, training and maintains a high-performance computing (HPC) and storage infrastructure.	Elixir Switzerland



FAIR Data

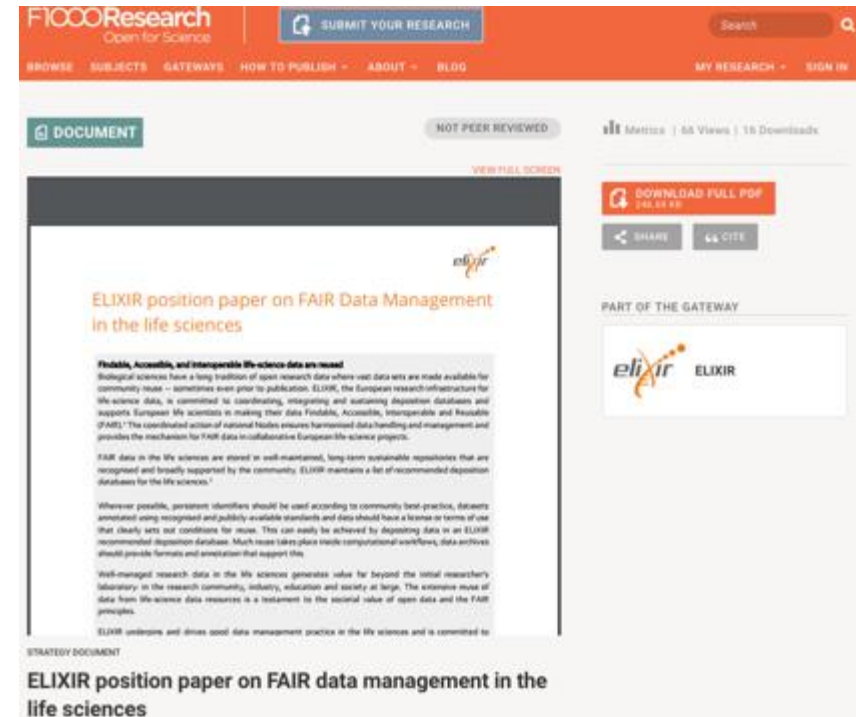
Open
data
is about
MORE
THAN
DISCLOSURE
it must be
Fair

- **F**indable
(Citable)
- **A**ccessible
(Trackable and countable)
- **I**nteroperable
(Intelligible)
- **R**eusable
(Reproducible)



ELIXIR Position Paper on FAIR data management in the life sciences

1. Open sharing of research data is a core principle
2. Data Management is crucial to science
3. Data should be submitted to deposition databases
4. All data submitted to Open Data archives should align with community-defined standards
5. ELIXIR Nodes implement FAIR for their respective nations
6. Professional skills, adequate resources and appropriate funding are needed for Data Management and infrastructure



Blomberg N and ELIXIR Consortium. ELIXIR position paper on FAIR data management in the life sciences. *F1000Research* 2017, 6(ELIXIR):1857 (document) (doi: [10.7490/f1000research.1114985.1](https://doi.org/10.7490/f1000research.1114985.1))

“Whenever possible, biological research data should be submitted to the recommended community deposition databases”

ELIXIR Deposition Database list

Deposition Database	Data type	International collaboration framework ¹
ArrayExpress	Functional genomics data. Stores data from high-throughput functional genomics experiments.	
BioModels	Computational models of biological processes.	
EGA	Personally identifiable genetic and phenotypic data resulting from biomedical research projects.	European Bioinformatics Institute and the Centre for Genomic Regulation
ENA	Nucleotide sequence information, covering raw sequencing data, contextual data, sequence assembly information and functional and taxonomic annotation.	International Nucleotide Sequence Database Collaboration
IntAct	IntAct provides a freely available, open source database system and analysis tools for molecular interaction data.	The International Molecular Exchange Consortium
MetaboLights	Metabolite structures and their reference spectra as well as their biological roles, locations and concentrations, and experimental data from metabolic experiments.	
PDBe	Biological macromolecular structures.	
PRIDE	Mass spectrometry-based proteomics data, protein expression information (2D values) and the supporting	

- The ELIXIR Deposition Databases meet the technical quality and governance criteria expected of ELIXIR [Core Data Resources](#)
- Agreed collectively by 21 Heads of Nodes
- International collaborative effort

<https://elixir-europe.org/platforms/data/elixir-deposition-databases>



“All data submitted to Open Data archives must be annotated in accordance with community-defined standards”



<https://elixir-europe.org/platforms/interoperability>



Identifiers.org and N2T.net help establish global standards for citation of biomedical data

SCIENTIFIC DATA

MENU ▾

Search E-alert Submit Login

Altmetric: 38 More detail >>

Editorial | **OPEN**

On the road to robust data citation

PDF Tools ▾

Associated Content

Scientific Data | Article | **OPEN**

[Uniform resolution of compact](#)

SCIENTIFIC DATA

MENU ▾

Search E-alert Submit Login

Altmetric: 38 More detail >>

Article | **OPEN**

Uniform resolution of compact identifiers for biomedical data

PDF Tools ▾

Associated Content

Scientific Data | Editorial | **OPEN**

[On the road to robust data citation](#)

Sarala M. Wimalaratne, Nick Juty, John Kunze, Greg Janée, Julie A. McMurry, Niall Beard, Rafael

ELIXIR FAIR-CMM Model

Level	Process	Datasets and Linksets
1. Initial	Processes are disorganized. Success is dependent on specialised, heroic and one-off efforts, considered unrepeatable, because processes are not sufficiently defined and documented to be replicated.	Datasets are disorganised and may well be unstructured. No Linksets (i.e. explicitly published mappings between datasets).
2. Repeatable	Basic processes are established, created and maintained. Successes could be repeated, because the processes are defined, and documented.	Basic levels of FAIR are implemented by the dataset. Linksets are implied as the links are intermingled with the data. Descriptions of the links are not available.
3. Defined	An organization has its own process through greater attention to documentation, standardization, and integration.	Datasets have limited metadata and access capabilities. Linksets are identified, i.e. there are descriptions of the datasets that are linked to.
4. Managed	Organization monitors and controls its own processes through data collection and analysis	Datasets have further metadata and access capabilities. Linksets are explicitly managed (but can be idiosyncratic).
5. Optimizing	Processes are constantly being improved through monitoring feedback from current processes and introducing innovative processes to better serve the organization's particular needs	Datasets are fully annotated with metadata and access capabilities, i.e. they fully satisfy all the FAIR principles. Linksets are managed as first class objects, i.e. regarded as datasets in their own right, and accessible to mapping services.

“ELIXIR Nodes implement a harmonised FAIR Data Management programme for the life sciences”

Support via Implementation studies

- Genomics Data Management with EGA / TraIT (NL, EBI, ES)
- Data Management Implementation study



elixir
CZECH
REPUBLIC

Navigation: Vision | Action steps | Lifecycle | Data | Roles | Managerial Questions

Let's dig the gold mine!

Bioinformatics produces a lot of data that is very valuable and that's our gold mine. In our working group, we realize this value of data. We set ourselves the following goal: help BioMed researchers mine their gold:

- Collect and provide the information about bioinformatics data produced
- Help the data producers to take care about their data (a.k.a. Data Stewardship)
- Help the data producers share their data with others.
- Connect and help interested parties to use the available data sources.

Sharing the data sources contents poses challenges regarding technical solutions and legal issues. The ultimate goal is making the data F.A.I.R., i.e. Findable, Accessible, Interoperable, and Reusable, while maintaining all the necessary constraints.

Check our action steps.

Contact
Phone: +420 220 183 267



ELIXIR Core Data Resources – fundamentally important to life-science research

ELIXIR Core Data Resource list

Core Data Resource	Data type
ArrayExpress	Functional Genomics Data from high-throughput functional genomics experiments.
CATH	A hierarchical domain classification of protein structures in the Protein Data Bank.
ChEBI	Dictionary of molecular entities focused on 'small' chemical compounds.
ChEMBL	Database of bioactive drug-like small molecules, it contains 2-D structures, calculated properties and abstracted bioactivities.
EGA	Personally identifiable genetic and phenotypic data resulting from biomedical research projects.
ENA	Nucleotide sequencing information, covering raw sequencing data, sequence assembly information and functional annotation.
Ensembl	Genome browser for vertebrate genomes that supports research in comparative genomics, evolution, sequence variation and transcriptional regulation.
Ensembl Genomes	Comparative analysis, data mining and visualisation for the genomes of non-vertebrate species.
Europe PMC	Europe PMC is a repository, providing books, patents and clinical publications.

<https://www.elixir-europe.org/platforms/data/core-data-resources>

- 16 Core Data Resources Nominated
- ELIXIR is committed to Open Access as a core principle for publicly funded research.
- Discussions on-going with Nodes, Resources and funders on high-quality, non-Open Access resources
- ELIXIR Core Data Resources should reflect this commitment and have terms of use or a license that enables the reuse and remixing of data.
- See "[Identifying ELIXIR Core Data Resources](#)"
- Agreed collectively by 21 Node directors



Open access life science data is part of the bioeconomy infrastructure



2010-2015:

30 771 patents used bioinformatics repositories to identify genes, enzymes, SNPs, ...



Vaccines



Pharmaceuticals



Beauty care



Industry enzymes ...

Open Data as an Innovation Driver – ELIXIR SME Programme

- 4 Innovation and SME Forums
- 2 events in 2018 (Cambridge & Frankfurt)
- Growing ELIXIR industry [intranet group](#) (23 people) and mailing list (550)
- External events: BioVision, London Festival of Genomics, ENVRI Plus, IMI stakeholder event



ELIXIR Industry strategy

April 2016

Background

Industry usage of many key bioinformatics resources within Europe is high. Users of ELIXIR services range from large multinationals to micro-SMEs and cover fields including pharma, biotech, food and agriculture and blue biotech. These industries are major employers globally, generating wealth and supporting transformation to a knowledge-based economy.

The biomedical sector is worth 460 billion annually¹. The pharmaceutical industry alone directly employs over 700,000 people in the EU, generating three to four times more downstream jobs and contributing to a trade surplus of 480 billion². As a sector, the bioinformatics industry is forecast to be worth 43.3 billion by 2020³. Europe is a rich breeding ground for biotech start-ups, with 1,799 healthcare biotech companies operating in Europe in 2012. The healthcare sector as a whole accounts for 8% of the total European workforce and for 10% of the EU's GDP⁴.

In the food sector, the EU was the top world exporter of food and drink products in 2012⁵, exporting 28.7 billion (20.5% of world total) and importing 48.5 billion (38.1%). Fish production in aquaculture is increasing on average 8.8% per annum globally⁶, driven by technological developments and better molecular understanding of species and parasites.

Industry itself sees added value in ELIXIR in terms of reduced costs and decreased duplication of effort, access to common

data and interface standards and much better public-private data integration. Many of these requirements are addressed through the general implementation of ELIXIR and are relevant for all users in industry and academia.

Given the number of research-intensive companies in Europe, and the increasing reliance within industrial R&D upon computational methods, stimulating innovation and supporting industry is a key priority for ELIXIR. Indeed, ELIXIR's Scientific Programme for the period 2015-2018 includes a Strategic Objective to "Support innovation in 'big data biology'".

The range of industry sectors that ELIXIR has the potential to support is also broad – from publishers to SMEs to HPC and cloud providers. ELIXIR's Industry Strategy aims therefore to be comprehensive in its scope, yet focused and tailored enough to address the needs of each industry sector.

Understanding Industry's needs

In order to understand the needs of industry and ensure that proposed activities are fit for purpose, ELIXIR's Industry Advisory Committee⁷ provides high-level strategic advice on the activities included in this strategy. The ELIXIR Industry Advisory Committee's recommendation reports^{8,9}, map closely against the activities within this strategy.

ELIXIR's Industry Strategy responds to input from several other reports and analyses of industry needs. The ELIXIR Preparatory Phase Industry report¹⁰, and a report entitled "Developing ELIXIR Interactions with Industry", commissioned in 2013 and carried out by ConnectedDiscovery, clearly articulates a number of key value drivers for

¹ Chakri et al. NEJM. 2015. doi:10.1056/NEJMp1502068

² European Federation of Pharmaceutical Industries and Associations: http://www.efpia.eu/facts_figures

³ Bioinformatics Market by Sector (Molecular Medicine, Agriculture, Forensic, Animal, Research & Gene Therapy), Segment (Sequencing Platforms, Knowledge Management & Data Analysis & Application (Genomics, Proteomics & Metabolomics) - Global Forecast to 2020": <http://www.marketresearch.com/Market-Report/bioinformatics-2016.html>

⁴ EC: Investing in Health 2015: http://ec.europa.eu/health/strategy/docs/wh_investing_in_health.pdf

⁵ FoodDrinkEurope: 2014: <http://www.food-drinkeurope.eu/industry-statistics/industry-statistics/2014-report-on-the-european-food-and-drink-sector.pdf>

⁶ Sustainable agriculture, forestry and fisheries in the bioeconomy: http://ec.europa.eu/economy_finance/docs/2014_11_19_1499_en.pdf

⁷ http://www.efpia.eu/facts_figures

⁸ http://www.efpia.eu/facts_figures

⁹ http://www.efpia.eu/facts_figures

¹⁰ http://www.efpia.eu/facts_figures

⁷ ELIXIR Scientific Programme 2015-2018: <http://www.elixir.eu/elixir-science-programme-2015-2018>

⁸ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

⁹ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

¹⁰ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

¹¹ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

¹² ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

¹³ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

¹⁴ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

¹⁵ ELIXIR Industry Strategy: <http://www.elixir.eu/elixir-industry-strategy>

Extending the outreach via local Industry Clusters



flanders.bio



Outcome from innovation events:



Node - collaboration



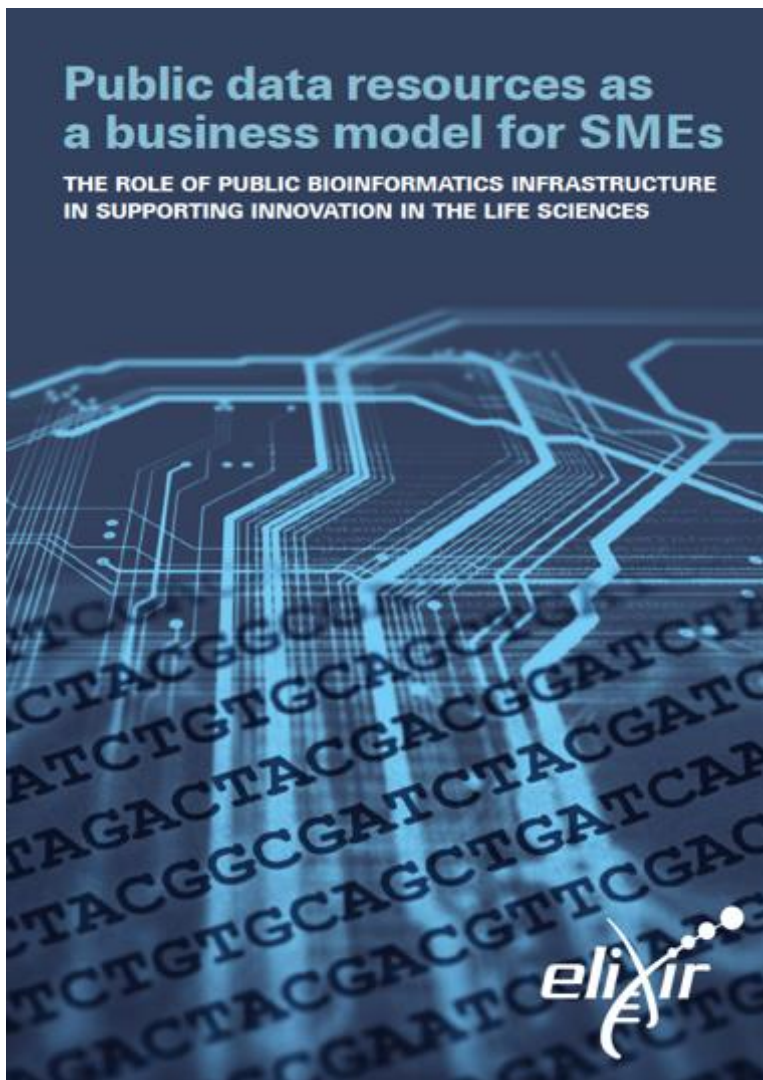
Node - collaboration



Service - exchange



Public data resources as a business model for SMEs



ELIXIR in numbers

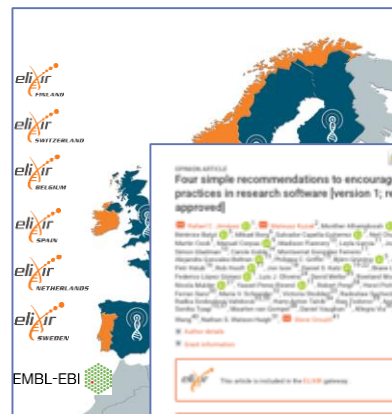
- 21 Members and 1 Observer
- ~ 180 institutes involved
- 600+ staff
- 16 Core Data Resources
- 23 Implementation Studies ongoing or soon to start
- 20 papers in ELIXIR F1000 channel
- 264 live events in TeSS
- 350 companies attended Innovation and SME programme

The image shows a screenshot of the F1000Research Channels page. At the top, there is a navigation bar with 'F1000Research / Channels'. Below this, there are three article thumbnails: 'Making Babies in the 21st Century' (03 Jul - 14 Aug 2017), 'Efficient Parallel Programming with GASPI @ HLRS' (3 - 4 Jul 2017), and 'Foundation skills for HPC in computational biomolecular research - BioExcel Summer School'. The main content area features a large group photo of many people standing in front of a modern building. Below the photo, there is a 'METHOD ARTICLE' section with a 'REVISED' badge and the title 'Identifying ELIXIR Core Data Resources; referees: 2 approved]'. The authors listed are Christine Durinx, Jo McEntyre, Ron Appel, Rolf Appel, and others. At the bottom, there is a section for 'InterPro' with the text 'Functional analysis of protein sequences predicting the presence of...'. The entire screenshot is overlaid on a large orange map of Europe with numerous blue pins indicating member locations.

ELIXIR Outcomes

ELIXIR Core Data Resource list

Core Data Resource	Data type
ArrayExpress	Functional Genomics Data from high-throughput functional genomics experiments.
CATH	A hierarchical domain classification of protein structures in the Protein Data Bank.
CHEBI	Dictionary of molecular entities focused on 'small' chemical compounds.
CHEMBL	Database of bioactive drug-like small molecules, it contains 2-D structures, calculated properties and abstracted bioactivities.
EGA	Personally identifiable genetic and phenotypic data resulting from biomedical research projects.
ENA	Nucleotide sequencing information, covering raw sequencing data, sequence assembly information and functional annotation.
Ensembl	Genome browser for vertebrate genomes that supports research in comparative genomics, evolution, sequence variation and transcriptional regulation.
Ensembl Genomes	Comparative analysis, data mining and visualisation for the genomes of non-vertebrate species.
Europe PMC	Europe PMC is a repository providing full-text access to biomedical journals, books, patents and clinical trial records.



Bioschemas.org



Secure long-term haping data landscape and science policy

Development of shared standards / conventions

Developed long-term, stable foundation services



Integration of node services

Community Standards



Tools



Soren Brunak



Alfonso Valencia

Data



Jo McEntyre



Christine Durinx

Human data



Serena Scollen



Tom Keane



Jordi Rambla

Compute



Tommi Nyrönen



Ludek Matyska



Steven Newhouse

Rare diseases



Ivo Gut



Marco Roos

Interoperability



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Many thanks
to Platform
and Use Case
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the hard
work in 2017!

ELIXIR's Coordination Groups



Training Coordinators Group

- Chair: Louisa Bellis



Technical Coordinators Group

- Chair: Rafael Jimenez



Thank you!



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