



The University of Manchester



Data Archiving and Networked Services



Welcome

researchobject

2018

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Center of Excellence for Computational Biomolecular Research

Twitter hashtags: [#eScience2018](#) [#ResearchObject](#)

Centre of Excellence for Computational
Biomolecular Research; H2020 grant [675728](#).

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The University of Manchester

researchobject.org



Research Object Community Update

Carole Goble, Stian Soiland-Reyes, Sean Bechhofer

The University of Manchester, UK

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RO2018, 29 October 2018, Amsterdam, 2018.

Satellite workshop of IEEE 14th International Conference on e-Science 2018



2010 – Research Objects not PDFs



Download Workflow Open in OnlineHPC

This workflow takes as input a file containing species occurrence points to create a model with the openModeller Web Service using the Bioclim algorithm. Environmental layers and mask are selected during the workflow. Points are filtered so that only environmentally unique points are used to create the model.

You can use this workflow to either filter a set of species occurrence points excluding redundant points with the same environmental conditions (you can get the result in the output "selected_csv_points") or to know the environmental ranges for each variable considering all input points. In the later case, you can inspect the "model_xml" to find this information in the Maximum and Minimum attributes of the Bioclim element. Values are separated by a space in the same order of the layers. Please note that there are no model tests or projections in this workflow.

For more information about the input file format, please look at the documentation for the corresponding parameter. If you use the default occurrence points you should know that Gammarus tigrinus is an aquatic species, so you need to choose marine environmental layers during the modelling procedure.

Workflow requirements: When running on Taverna workbench, this workflow requires Internet connection and the Taverna interaction plugin installed.

Preview

Run

Run this Workflow in the Taverna Workbench...

Option 1:
Copy and paste this link into File > Open workflow location...
<http://www.myexperiment.org/workflows/3725/download?version=4>
(More info [?])

Run this Workflow on the cloud with OnlineHPC...

Workflow Type: Taverna 2

Uploader: Renato De Giovanni

License: All versions of this Workflow are licensed under: [Creative Commons License]

Version 4 (latest) (of 4)

Credits (2): Renato De Giovanni, BioVet.

Attributions (0)

Tags (13): bio-oracle, bioclim, bioenv, eurora, filter, harmonize, world soil database, hess, hots4, hots4d, inofopen, openmodeller, points, range, workflow

Shared with Groups (2): BioVet.

Workflow

Results

Investigation

Workflow

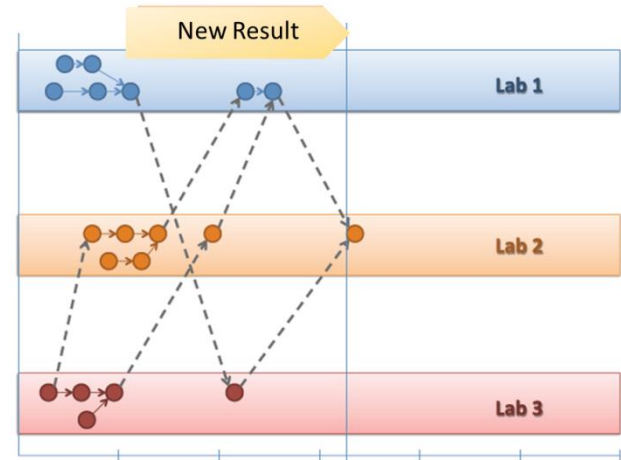
Data

Paper

Results

Investigations

Research has many components, of many types



Exchange of *all* the components of an investigation

Computational instruments break or need to be maintained

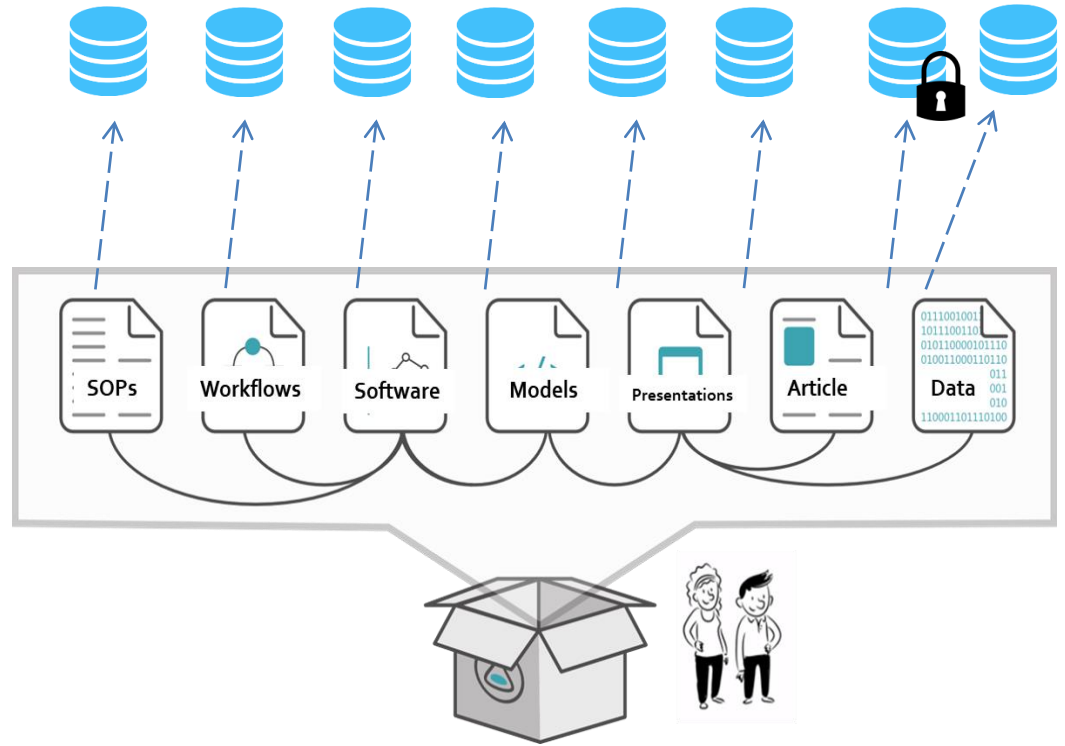
Science, and its products, evolve



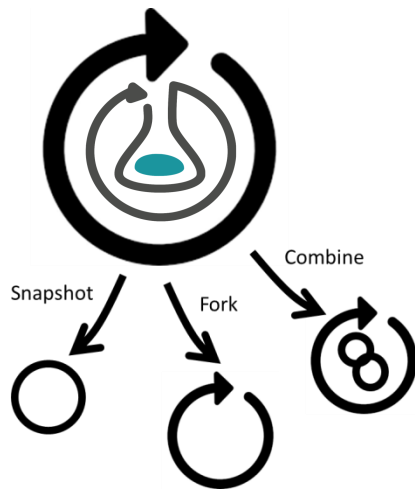


Overcome fragmentation

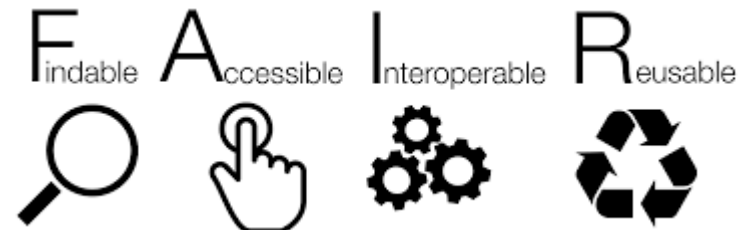
Bundle and relate components



Release evolving research
Accelerate exchange



Support preservation, reproducibility, reuse





Research Object Framework

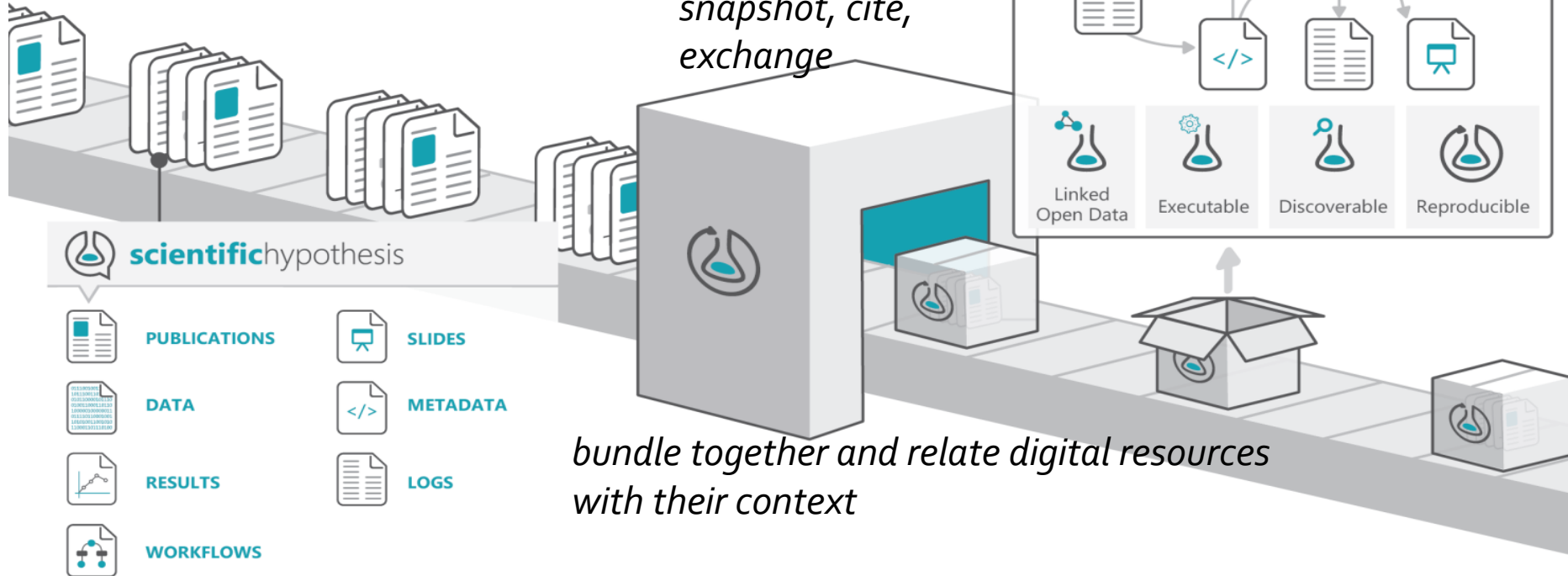
Standards-based generic metadata framework

Bechhofer et al (2013) <https://doi.org/10.1016/j.future.2011.08.004>

Bechhofer et al (2010) <https://eprints.soton.ac.uk/268555/>

 Enabling **reproducible**, transparent research.

carry machine processable metadata common and specific to different object types



Data used and results produced
Methods used to produce /analyse that data

Provenance and settings, **People** involved,
Annotations understanding & interpretation

Howard Ratner,
Chair STM Future Labs Committee, CEO EVP Nature Publishing Group
Director of Development for CHORUS (Clearinghouse for the Open Research of US) STM
Innovations Seminar 2012

Research Objects?

The diagram illustrates the concept of Research Objects (RO) and their relationships. A central grey box labeled 'RO' is connected to various components: Metadata, Workflow 13, Workflow 16, Results, Paper, Slides, and Logs. Red dashed arrows indicate aggregation, showing how individual components are grouped into a Research Object. Blue dashed arrows indicate representation, showing how the Research Object is represented by various formats like PDFs, images, and charts. Green solid arrows indicate domain relations, showing the flow of information between components. A legend at the bottom left defines the arrow types: blue dashed for Representation, red dashed for Aggregation, and green solid for Domain Relations.

Credit: S. Bechhofer et al., "Research Objects: Towards Exchange and Reuse of Digital Knowledge," 2010

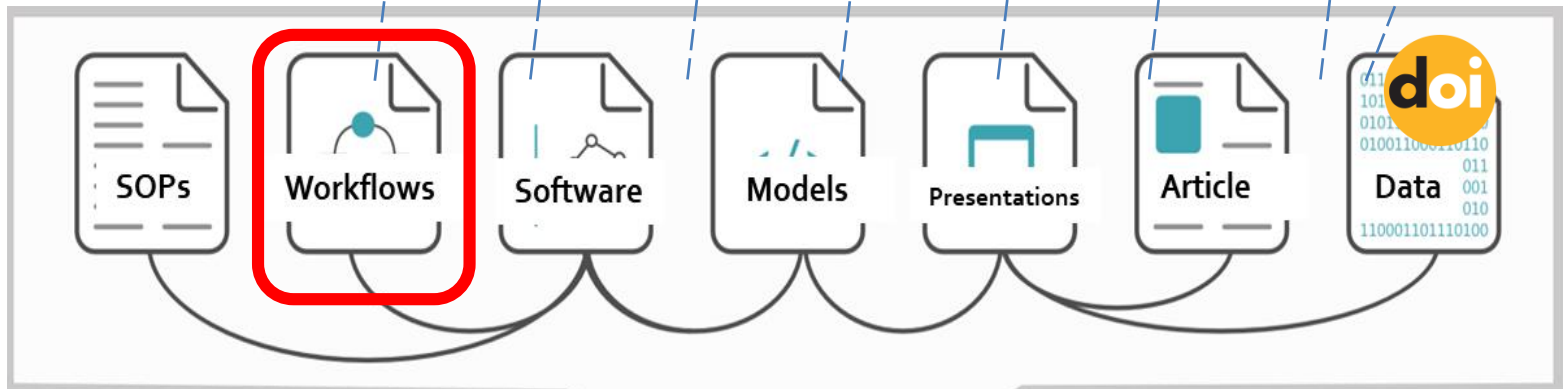
STM Innovations Seminar U.S. –
Reinventing Innovation
May 1, 2012
Washington DC, USA

Howard Ratner is shown in a video frame, speaking at a podium during the STM Innovations Seminar U.S. in Washington DC, USA.

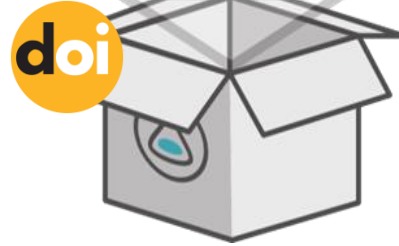


"Unbounded" Objects

Bags of things and external references to things

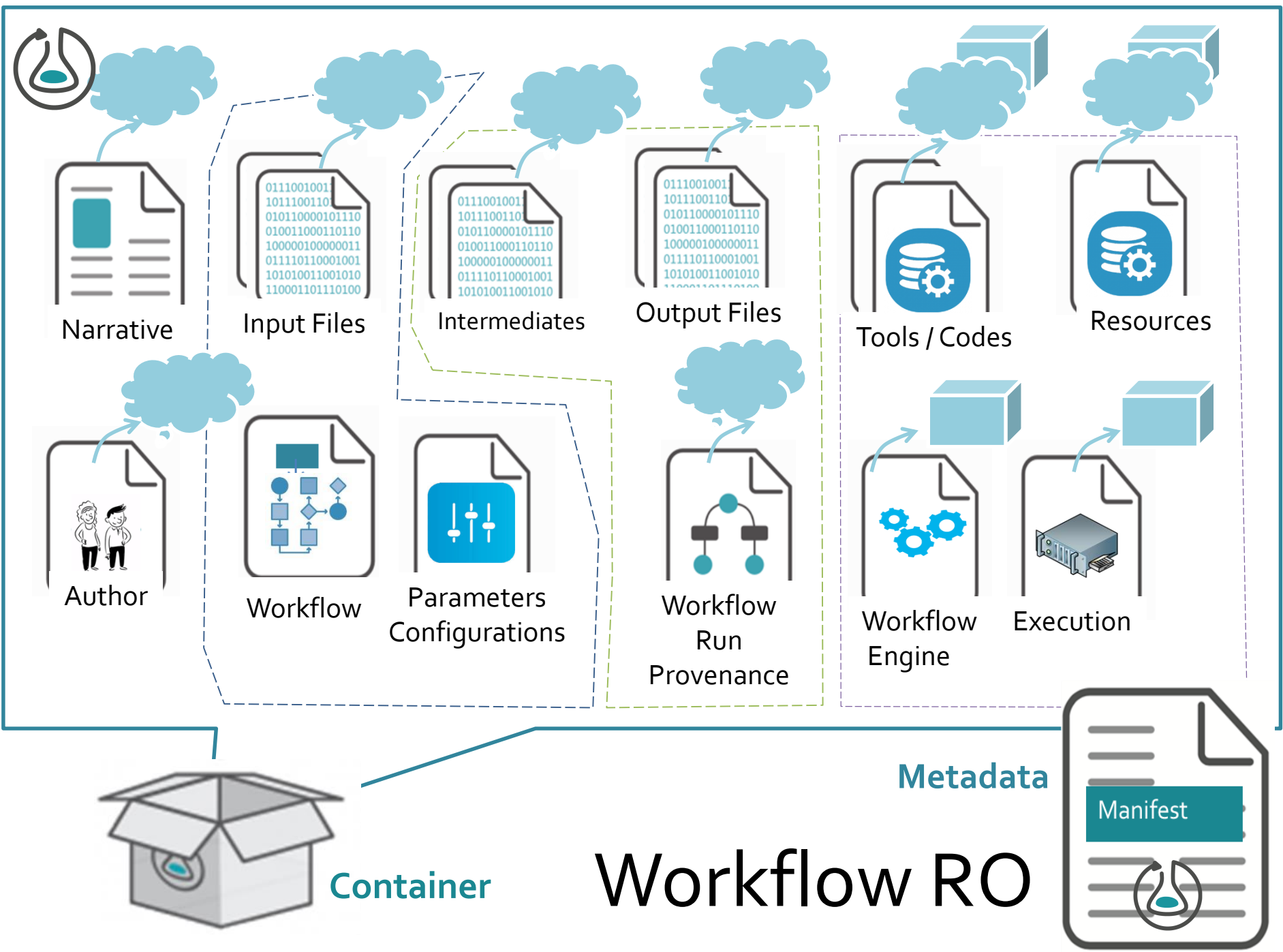


A Metadata Object
that represents
properties in common
across all research
artefacts types,
*common PIDs and
metadata*



Container

**A Digital Package Object
Type** composed of many
interrelated elements that
bundles together and
relates digital resources of
a scientific investigation
with context.





Describe and run workflows, and the command line tools they orchestrate, supporting containers to be portable, transparent and interoperable .



Describe the provenance of the workflow

EDAM

Describe the workflow inputs, outputs, tools and data with controlled vocabularies / ontologies



Workflow systems run the CWL workflow



Software components are containerised to be portable

Workflow RO



Metadata



Gather the CWL workflow descriptions + rich context, provenance using multi-tiered descriptions
Snapshot workflow.
Relate it to other objects.



Container

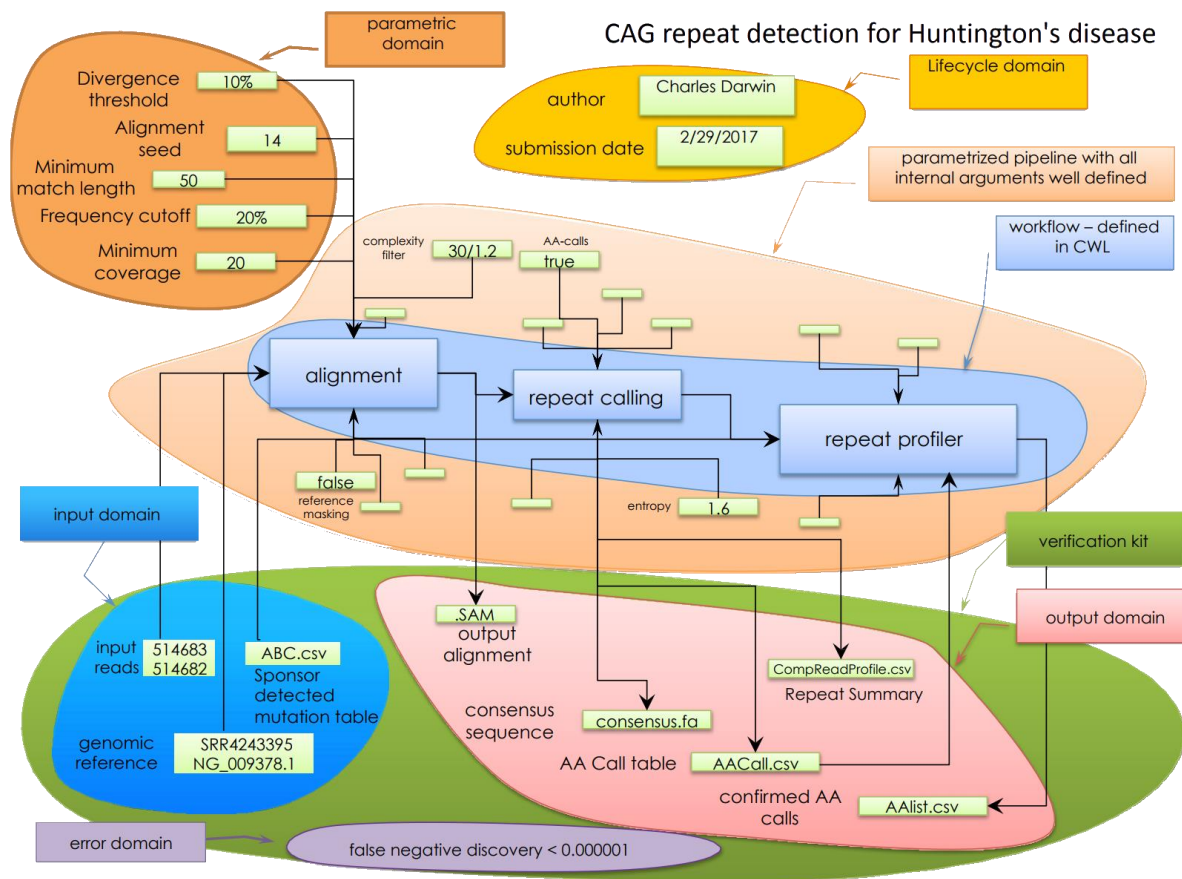


Archive formats to contain the object

<https://www.commonwl.org/>

Standardize exchange of HTS workflows for regulatory submissions between FDA, pharma, bioinformatics platform providers and researchers

Inspect and replicate the computational analytical workflow to review and approve the bioinformatics





Desiderata

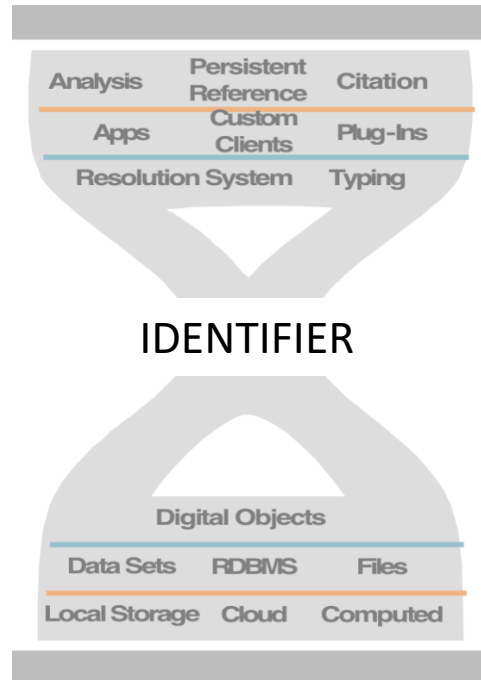
Use Open Standards.

Incremental.

**Graceful
degradation.**

Machine-
processable.

Technology
Independent.



The least possible.

Commodity
tooling.

Multi-platform.

The simplest feasible. Low tech.

Low user overhead and thin client



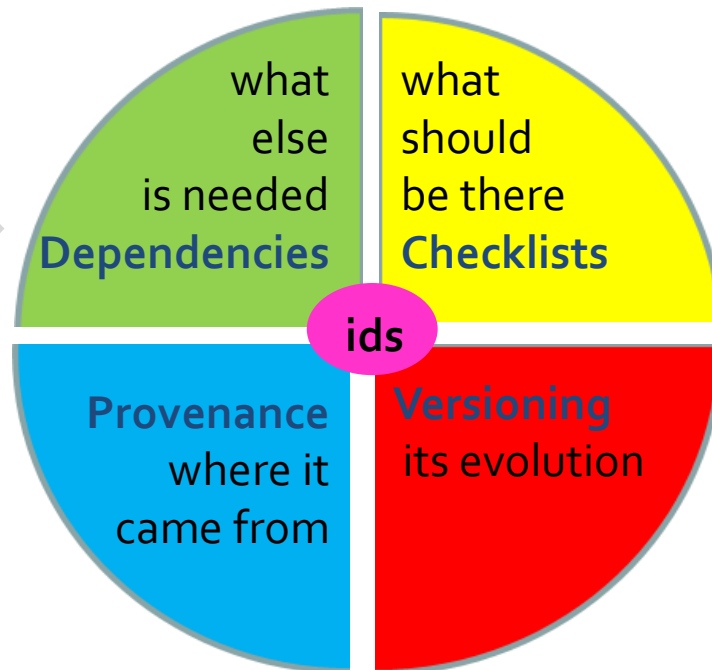
Tailored metadata profiles to describe a RO



Manifest Profile Descriptions



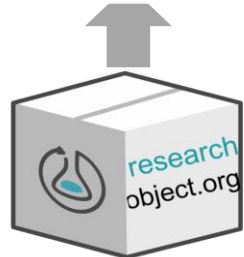
Container





Tailored metadata profiles to describe a RO *general purpose to drive scalable infrastructure*

Manifest Profile
Descriptions



Container

Overview Content Relations **Quality** Notifications History

Earth Science data-centric RO key requirements

✗ Target **Analyzing gene deregulation in Huntington's disease with respect to epigenetic information** *does not satisfy* checklist for *ready-to-release*.

- ✓ Research Object has title
- ✓ Research Object has description
- ✓ Research Object has creator
- ✗ Research Object does not have editor
- ✗ Research Object does not have indicated purpose
- ✓ Research Object has subject
- ✗ Research Object does not have copyright holder
- ✓ Data artifact is present (e.g., dataset, document, file, image)
- ✗ One or more data artifact has no format indicated, including http://sandbox.rohub.org/rodl/ROs/HD_chromatin_analysis/broad_hmm_3_Poised_Promoter.txt
- ✗ One or more data artifact has no filesize indicated, including http://sandbox.rohub.org/rodl/ROs/HD_chromatin_analysis/broad_hmm_3_Poised_Promoter.txt
- ✗ Research Object does not have access level specified
- ✗ Research Object does not have DOI

See quality history with RO Monitoring Tool

Validate



towards generic approaches ...

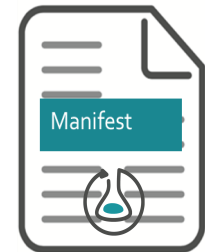


Tailored metadata profiles to describe a RO

general purpose to drive scalable infrastructure



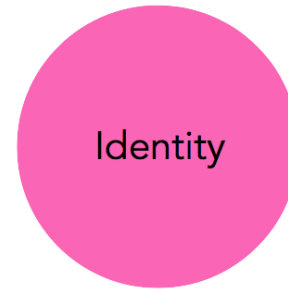
Manifest
Profile
Description



Manifest
Construction



Container
Profile

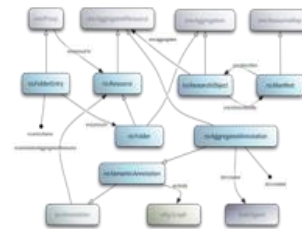


Identity

Identifiers: URI, RRI, DOI, ORCID



Identifiers.org **minid**



Research Object
Manifest Model



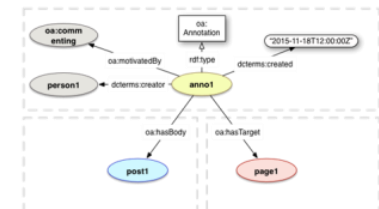
Aggregation



Open Archives Initiative
Object Exchange and Reuse
Aggregation to enumerate
and link together



Annotation



W3C Web Annotation Vocabulary
Annotation about RO, elements
and their relationships



Manifest Construction



<https://w3id.org/ro/2016-01-28>



Research Object ontology

Release 2016-01-28

This version:

<https://w3id.org/ro/2016-01-28/>

Latest version:

<https://w3id.org/ro/>

Previous version:

<https://w3id.org/ro/2013-11-30/>

Revision:

1.0.0-SNAPSHOT

Editors:

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Imported Ontologies:

[OAI-ORE](#)
[Metadata](#)

DOI: 10.5281/zenodo.12744



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Includes wfdesc and wfprov
Basis of CWL and CWLProv
Time for a review....?
e.g. RO type to help tools..



Container Profiles



Research Object Bundle 1.0

Specification for a structured ZIP-file, based on the ePub and Adobe UCF specifications

<https://researchobject.github.io/specifications/bundle/>



Specifies a file system structure for transferring and archiving a collection of files, **including their checksums to verify and validate content** and brief metadata.

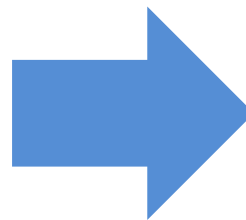
<https://github.com/ResearchObject/bagit-ro>



mechanism for serialization and transport consistency,



capture identity, annotations and provenance of the resources

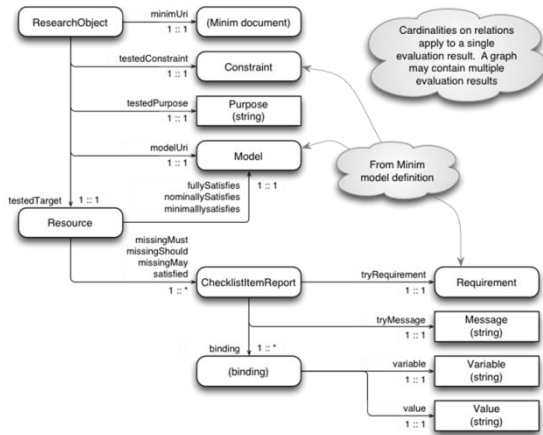


Big Data collections of arbitrary referenced content

<https://github.com/fair-research/bdbag>



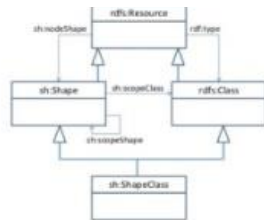
Manifest Profile Description general construction & validation tooling



Minim model for defining checklists



Shapes Constraint Language



Linked Data and RDF Shapes
Validate graph-based data against a set of conditions

- RO pre-processing to merge to single graph
- RDF Shape that indicates to follow links
- Bespoke validators / unpackers to iterate over the RO



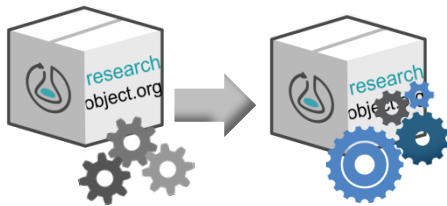
Research Object drivers



Exchange & Commons



Preservation and fixed point publishing



Reproducibility and execution



Active "release" research



LU
MC
Leids Universitair
Medisch Centrum



(GIGA)ⁿ_{DB}



everest

Workflows Models Data



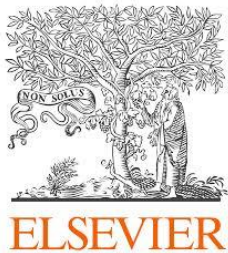
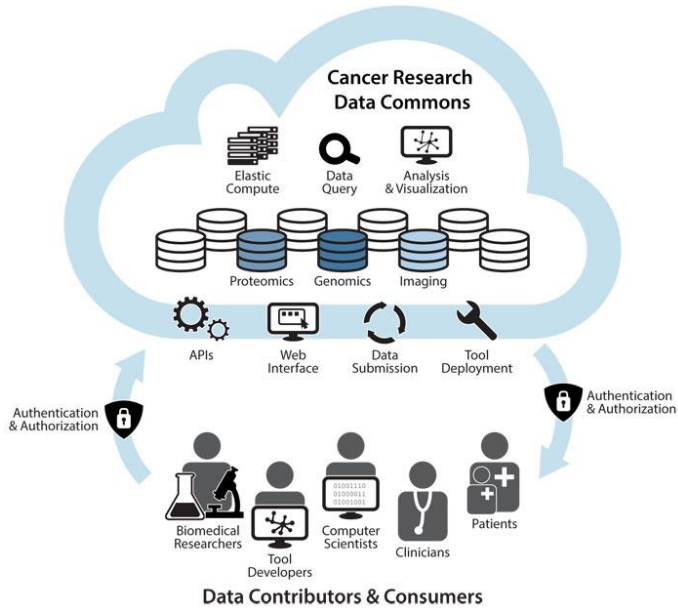
	Workflows	Models	Data	Research Object (Double Arrows)	Research Object (Document)	Research Object (Gears)	Research Object (Open Box)
LU MC	●	●	●	●	●	●	●
Workflow 4Ever	●	●	●	●	●	●	●
COMMON WORKFLOW LANGUAGE	●	●	●	●	●	●	●
CANCER GENOMICS CLOUD SEVEN BRIDGES	●	●	●	●	●	●	●
(GIGA) ⁿ _{DB}	●	●	●	●	●	●	●
elixir	●	●	●	●	●	●	●
BioCompute Objects	●	●	●	●	●	●	●
FAIRDOM	●	●	●	●	●	●	●
BIG DATA BAG	●	●	●	●	●	●	●
STELAR	●	●	●	●	●	●	●
Open PHACTS	●	●	●	●	●	●	●
everest	●	●	●	●	●	●	●



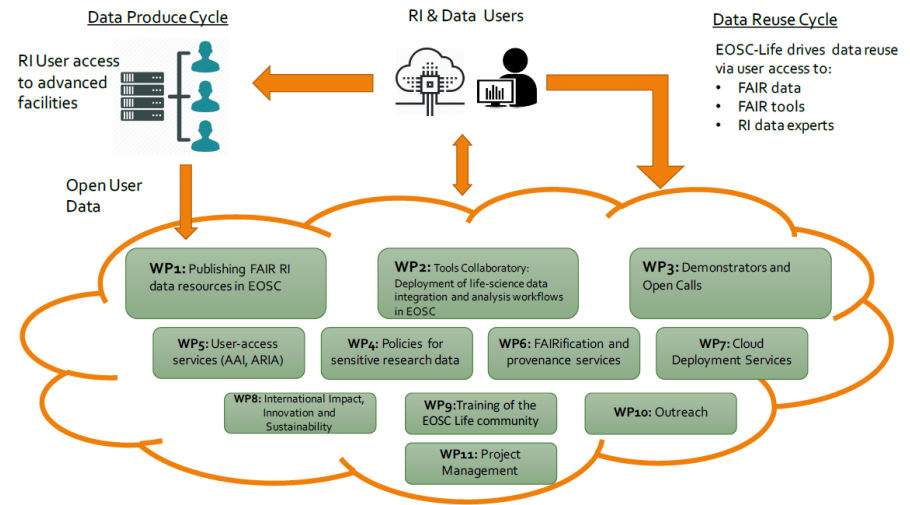
NIH Data Commons



CANCER GENOMICS CLOUD SEVEN BRIDGES



European Open Science Cloud



CWL Workflow Collaboratory





Research Object Gaps

research
object.org

RO Profiles

- Templates & Standard types

General tooling

- Construction, Validation, Viewing

Handling RO composition

- Nesting, Complex & mixed types
- Less flexibility -> easier tools

RO Life cycles stewardship

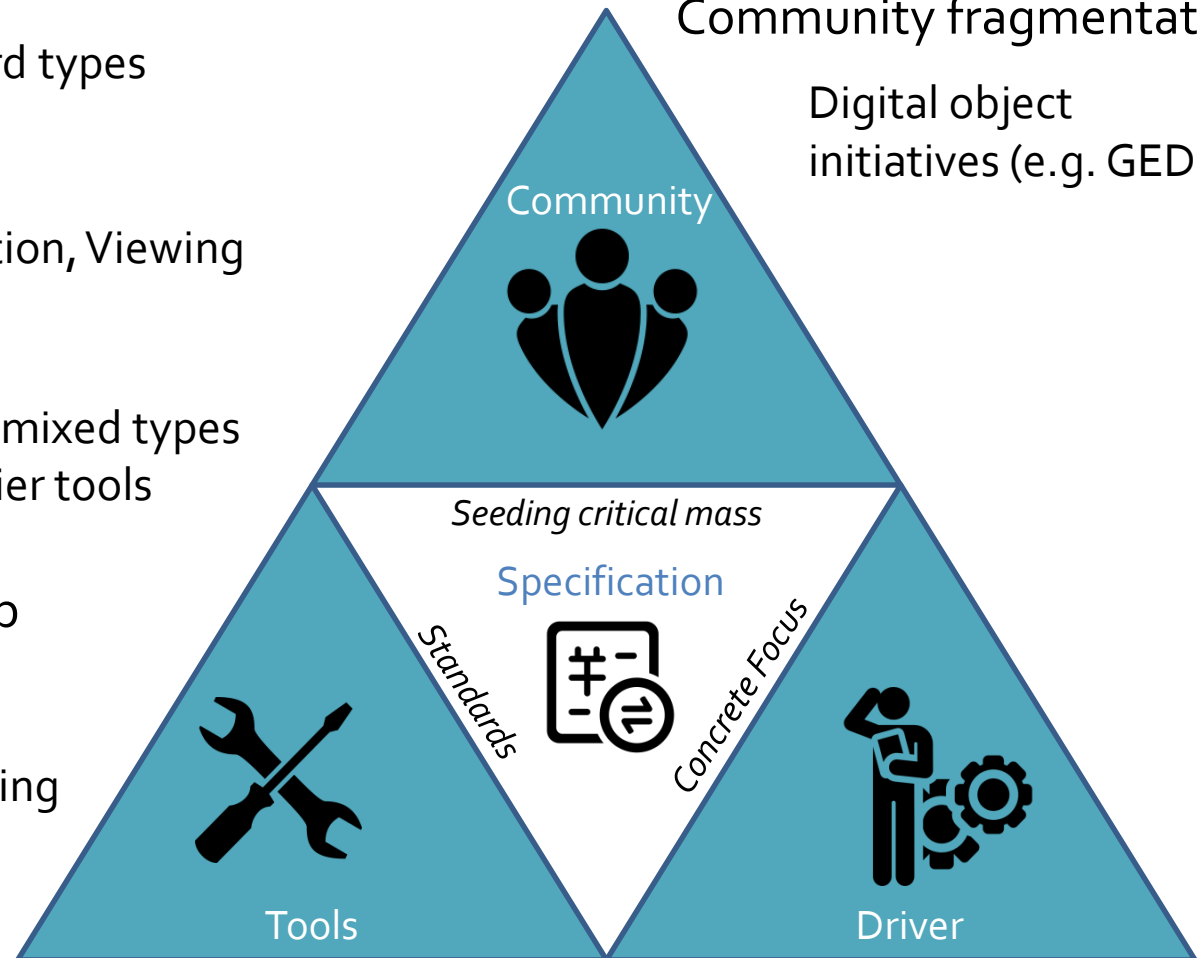
- Fixed snapshot
- Living objects
- Rot, mutations, cloning
- References

RO Circulation

- Credit, tracking

Community fragmentation

Digital object
initiatives (e.g. GEDE)





researchobject.org

Build a Community

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