



A new Approach for Publishing Workflows: Abstractions, Standards and Linked Data

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 - **Publishing abstract workflows and specific workflows**
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 - **Linked Data Publication**
- 4. Workflow querying and Linked Data consumption**
- 5. Conclusions**

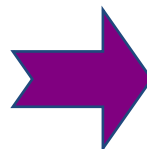
Typical Published Article

Text:
Narrative of method,
software packages used

Data:
Key datasets and figures/plots

NOT published,
loosely recorded:

Software:
scripted codes + manual steps +
notes/emails



Reproducible Article: Weaver, GenePattern GRRD, etc.

Text:
Narrative of method,
software packages used

Data:
Key datasets and figures/plots

Workflow:
Workflow/scripts describing
dataflow, codes, and parameters

Reproducible Article: Weaver, GenePattern GRRD, etc.

Text:

Narrative of method,
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Data:

Key datasets and figures/plots

Workflow:

Workflow/scripts describing
dataflow, codes, and parameters

Only executable workflow is published:

1. Must have the **same codes** to re-execute the workflow, but:
 - Codes become unavailable
 - Eg: eHits was proprietary and replaced by AutodockVina
 - Different labs prefer different codes
 - Eg: R vs Matlab
 - Eg: viz in Cytoscape vs yEd
2. Must have the **same workflow framework** to re-execute the workflow
 - Must have R for Weaver
3. Must **import files** to local file system and workflow framework
 - Must import bundle of workflow/data/code files to reproduce

- Publish an **abstract workflow** in addition to executable workflow
 - Description of workflow that is independent of the codes executed
 - Maps to the codes executed (the “executable workflow”)
- Publish both abstract and executable workflow **using the OPM standard**
 - OPM (Open Provenance Model) is independent of workflow framework and is widely implemented
 - Other groups can import to their own workflow framework
- Publish data and workflows as **Linked Data** on the Web
 - All workflows and related files are web-accessible
 - Simple mechanism to share across local file systems

1. Use URIs as names for things.

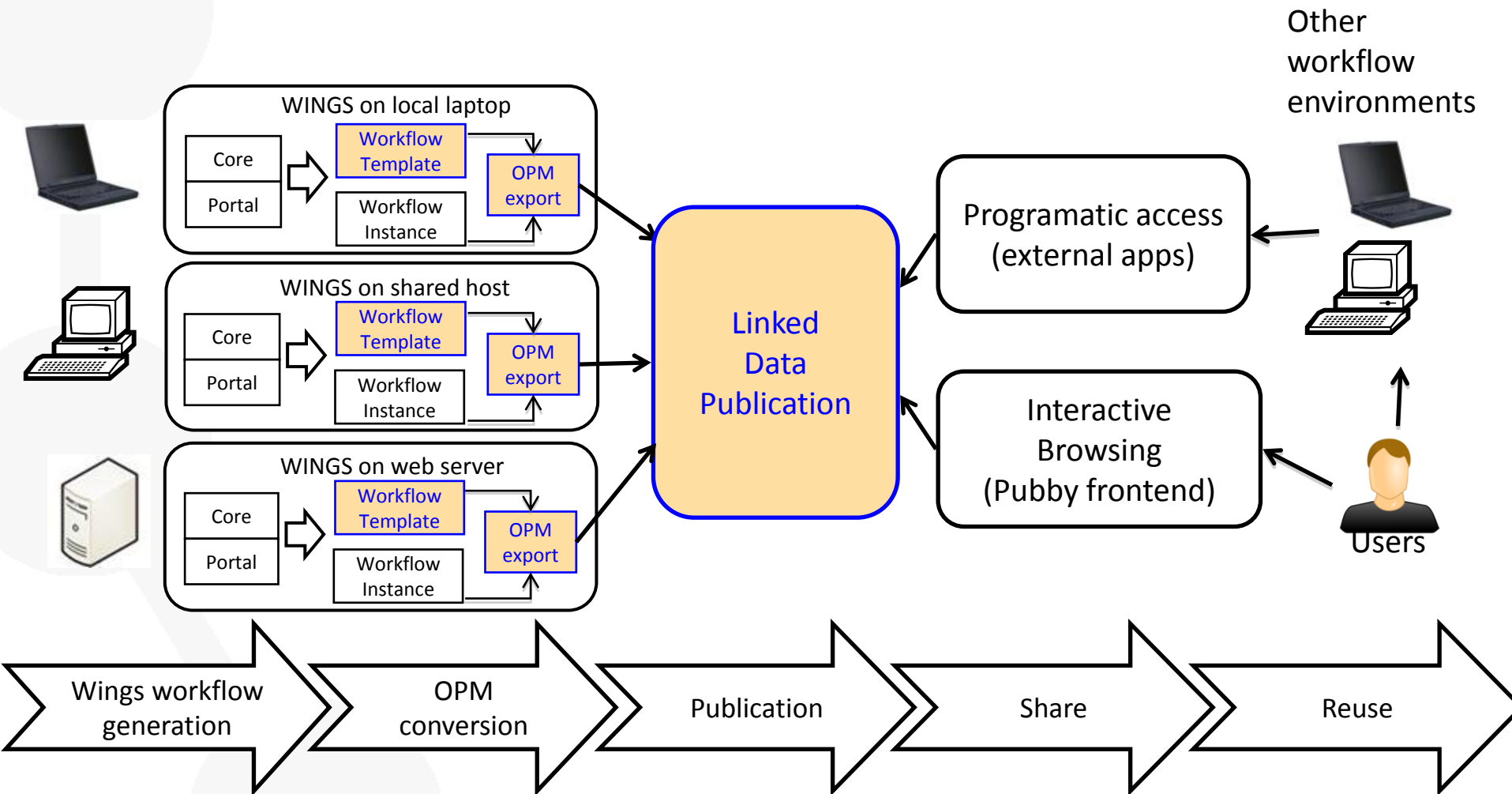
2. Use HTTP URIs so that people can look up those names.

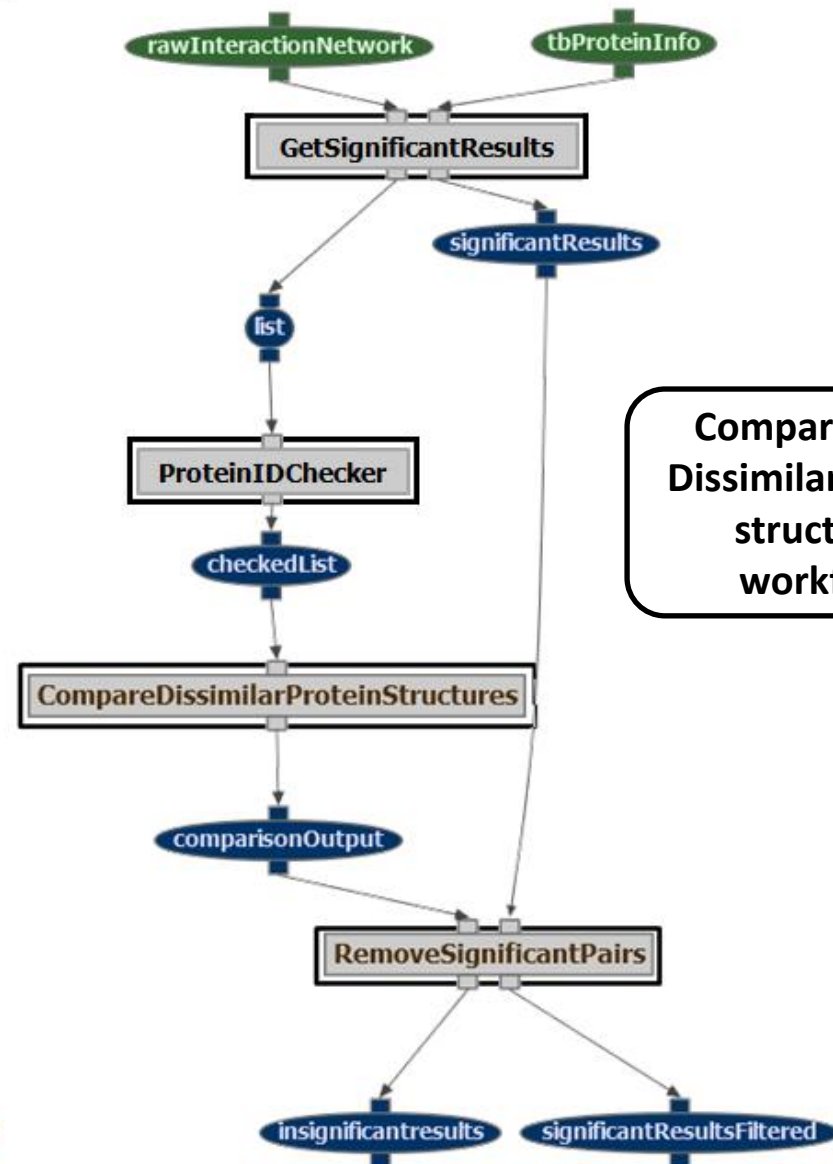
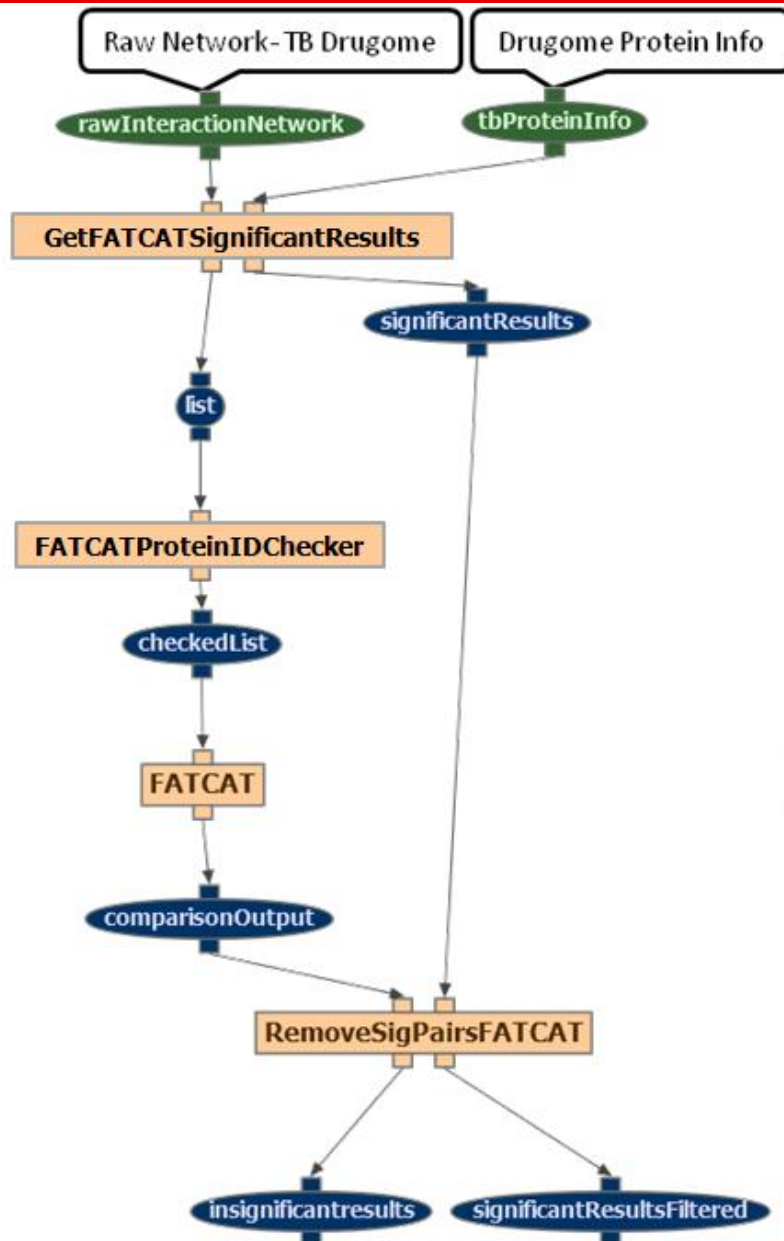
3. When someone looks up a URI, provide useful information.

4. Include links to other URIs.

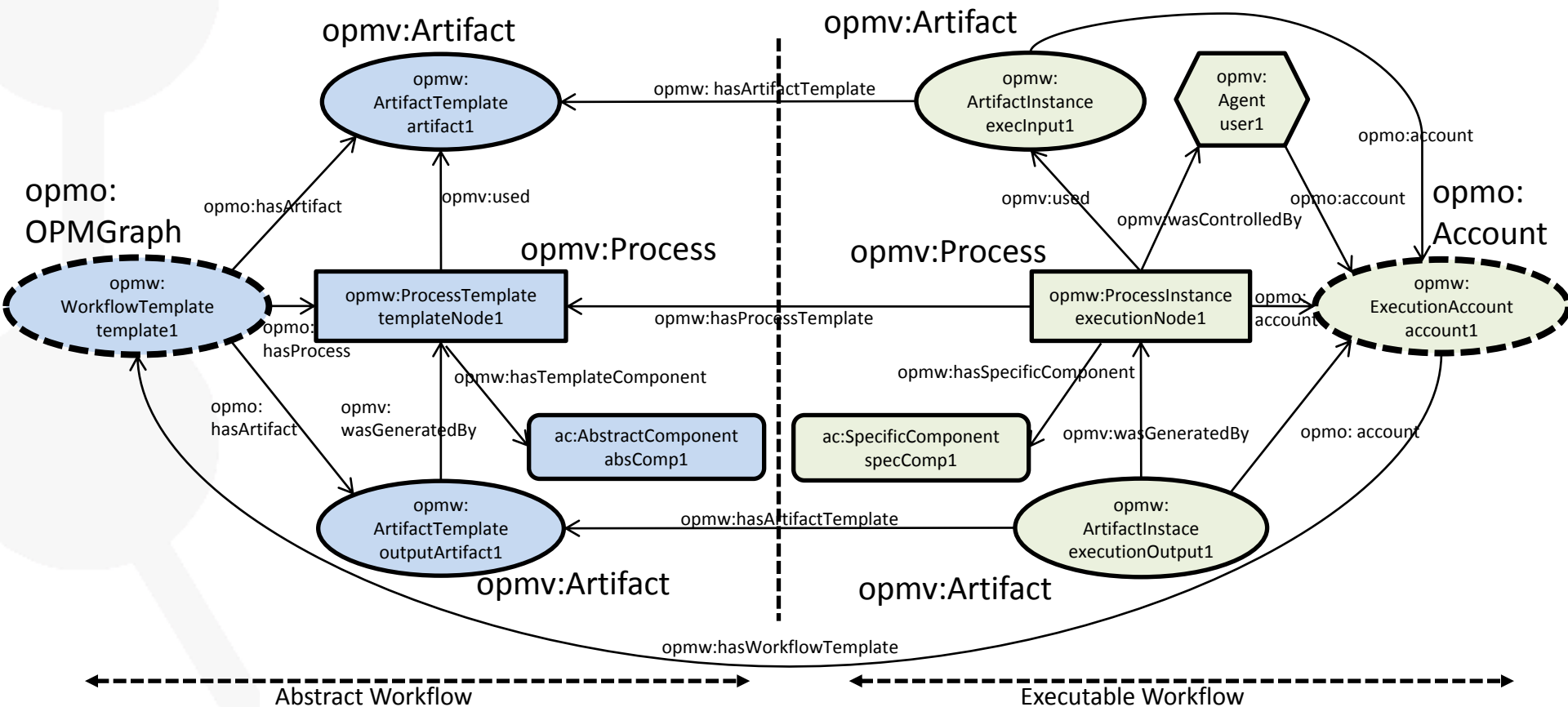


“Linking Open Data cloud diagram, by Richard Cyganiak and Anja Jentzsch. <http://lod-cloud.net/>”

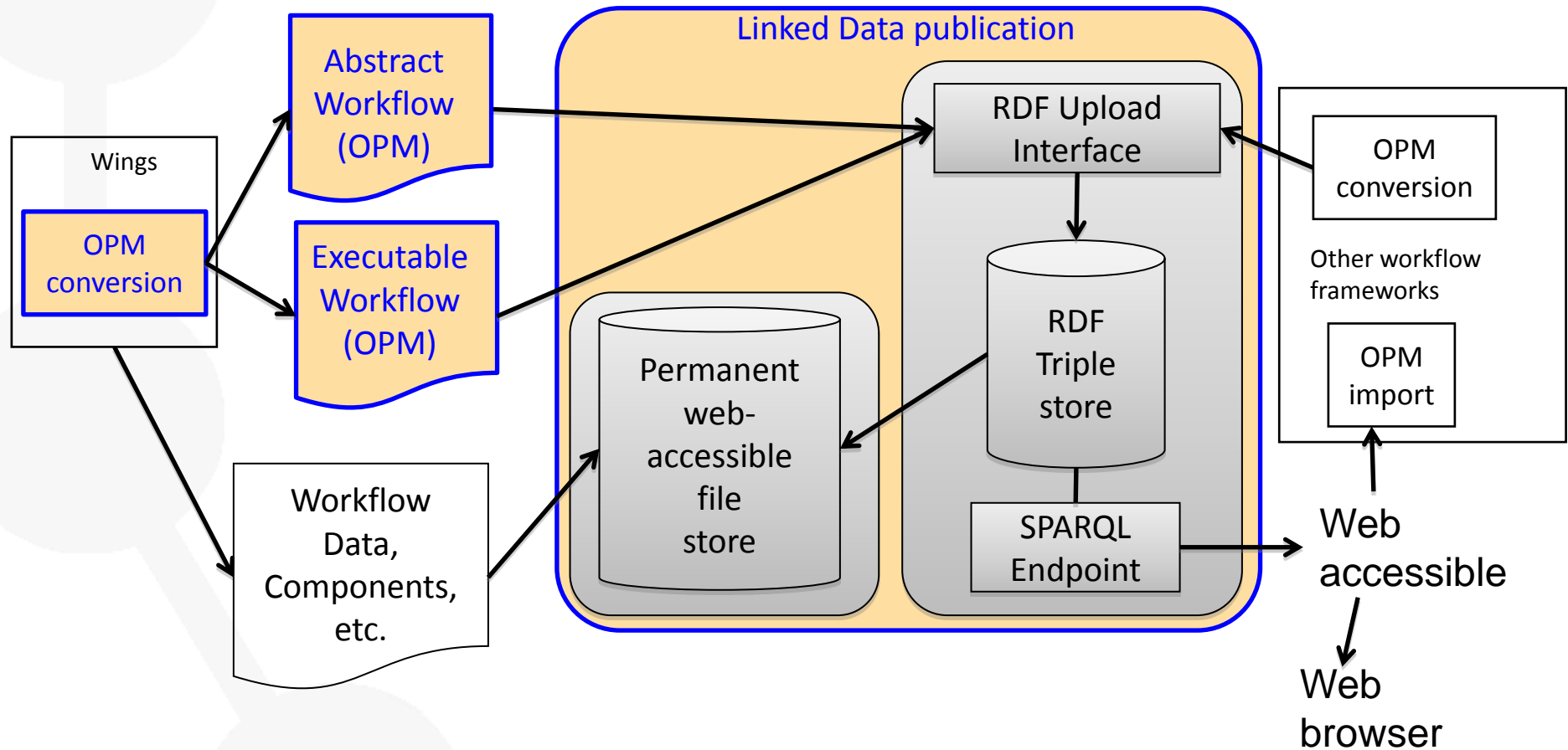




Comparison of
Dissimilar protein
structures
workflow



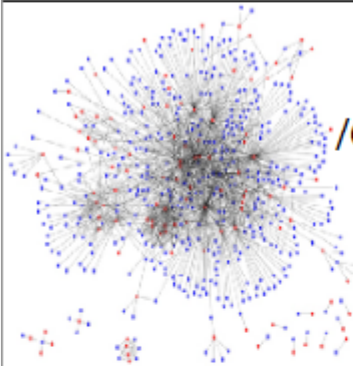
Publication of Workflows as Linked Data





Types of search

[Workflow search](#) | [Author and contributor search](#) | [Global search](#)



<http://wings.isi.edu/opmexport/resource/ProcessInstance/COMPAREDISSIMILARPROTEINSTRUCTURES1314662017568>

Global search (all resources)

Insert the name of a resource to check its related information

Resource URI
(Process instance)

compareDissimilarProteinStructures
<http://wings.isi.edu/opmexport/resource/ProcessInstance/FORMATSTEPTOCOMPAREDISSIMILARPROTEINSTRUCTURES1314662017568>
<http://wings.isi.edu/opmexport/resource/ProcessInstance/COMPAREDISSIMILARPROTEINSTRUCTURES1314662017568>

Autocomplete search bar

Properties

| | |
|-----------------------------|--|
| hasSpecificComponent | FATCAT/COMPONENTFATCAT |
| hasProcessTemplate | ProcessTemplate/ABSTRACCTSUBWFFATCAT_COMPAREDISSIMILARPROTEINSTRUCTURES1314662017568 |
| wasControlledBy | Agent/DANIEL |
| used | ArtifactInstance/31A41E8CDA244334B692F13A2CB7C4C1 |
| account | Account/ACCOUNT1314662017568 |
| type | ProcessInstance |

Specific component for this process instance

Searching/Browsing Workflows as Linked Data

| | | |
|----------------------|---|---|
| Component Name | COMPAREDISSIMILARPROTEINSTRUCTURES | Component Name |
| INPUTS | | Component Inputs |
| uses | DataVariable/INPUT137 | |
| OUTPUTS | | Component Outputs |
| generates | DataVariable/COMPARISONOUTPUT | |
| IMPLEMENTATIONS | | Code Implementations |
| code | COMPAREDISSIMILARPROTEINSTRUCTURESImpl0 | |
| code | COMPAREDISSIMILARPROTEINSTRUCTURESImpl1 | |
| code | COMPAREDISSIMILARPROTEINSTRUCTURESImpl2 | |
| TEMPLATE METADATA | | Template additional metadata |
| type | WorkflowTemplate | |
| label | fatcatTest | |
| contributor | Agent/DANIEL | |
| hasDocumentation | | |
| hasVersion | 2 | |
| lastUpdateTime | 2011-10-04T01:58:33-07:00 | |
| AVAILABLE EXECUTIONS | Account/ACCOUNT1319113809483 | Record of the different executions of this workflow |

1. Publication of an **abstract workflow** that represents the computational method in an **execution-independent manner**.
2. Publication of **the abstract workflow** and the **executed workflow** using the **OPM standard** that is independent of the execution environment used.
3. Publication of the workflows, components, codes and datasets **as Linked Data** on the web.

- Extensions to abstract workflow publication
 - Be able to provide abstractions **on several steps**.
 - Incomplete provenance.
- Create an OPMV/W3C PROV-O profile for common workflow representation.
 - **Increase interoperability** with other workflow representation systems.
- Workflow reuse in **different workflow systems**.
 - Import and execute workflows in other workflow frameworks.

- WINGS workflow system: <http://seagull.isi.edu/marbles/>
- The Open Provenance Model Specification: <http://openprovenance.org/>
- OPMO: <http://openprovenance.org/model/opmo>
- OPMV: <http://open-biomed.sourceforge.net/opmv/ns.html>
- TB Drugome Wiki (Evolution of this work):
http://seagull.isi.edu/wings-drugome/index.php/Main_Page
- W3C PROV-O current ontology (draft):
http://www.w3.org/2011/prov/wiki/PIL_OWL_Ontology
- Principles of Linked Data:
<http://www.w3.org/DesignIssues/LinkedData.html>

- UCSD people:

- Li Xie

- Lei Xie

- Sarah Kinnings

- Phil Bourne

- ISI people:

- Varun Ratnakaar

- OEG people:

- Oscar Corcho





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