

WfInstances Browser

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Introduction:

Scientific workflow applications have been used by scientists to support some of the most significant discoveries of the past several decades and are executed daily to serve a wealth of scientific domains.

Workflow research and development activities rely on the use of **workflow instances** to describe, execute, profile, and evaluate production workflows, which can be applied in simulated and real world conditions.

The WfCommons project provides users a with a collection of open access production workflow instances from various scientific applications shared in a common instance format. **However, these instances contain information that is currently only accessible by manually reading through the instance json files, which is a labor- and time-consuming process.**

Methodology

- **Agile Project Management** in the form of a Kanban Board utilizing Issue-Driven Project Management
- **Defined Team Roles** split among group members at the beginning of the project to ensure an equal work split
- **Weekly Team Meetings** to keep team members and the sponsor well informed throughout the entirety of the development process

Technology Stack:

Front End:

Back End:

TS

Remix

FastAPI

mongoDB

Objective:

Create a web application that allows users to easily browse, sort, visualize, select and download workflow instances.

Download

Workflow

Visualize Workflow

id

Filter by id

Filter Mode: Fuzzy

Github Repo

Filter by Github Repo

Filter Mode: Includes

of Tasks

Min

Max

Filter Mode: Between

of Files

Min

Max

Filter Mode: Between

Metrics

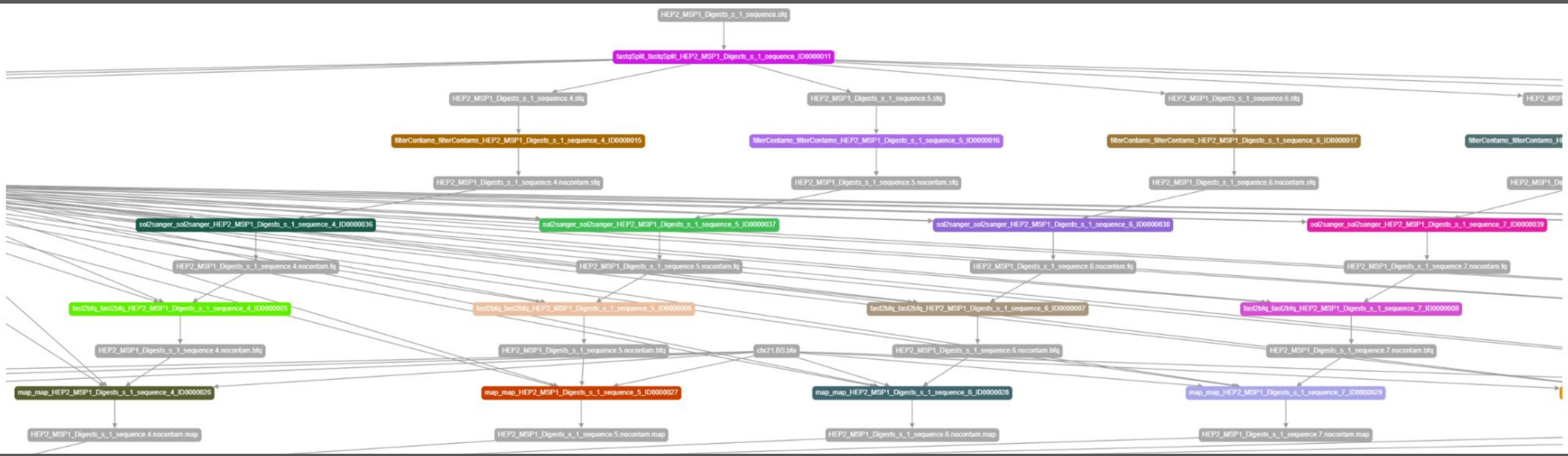
Total MB Read

Filter by Total

Filter Mode: Fuzzy

<input type="checkbox"/>	blast-chameleon-large-001.json	wfcommons/WfInstances	103	307	3222.10 MB
<input checked="" type="checkbox"/>	blast-chameleon-large-002.json	wfcommons/WfInstances	103	307	3202.14 MB
<input type="checkbox"/>	blast-chameleon-large-003.json	wfcommons/WfInstances	103	307	3220.94 MB
<input checked="" type="checkbox"/>	blast-chameleon-large-004.json	wfcommons/WfInstances	103	307	3199.19 MB
<input type="checkbox"/>	blast-chameleon-large-005.json	wfcommons/WfInstances	103	307	3201.58 MB
<input checked="" type="checkbox"/>	blast-chameleon-medium-001.json	wfcommons/WfInstances	303	907	961.92 MB
<input type="checkbox"/>	blast-chameleon-medium-002.json	wfcommons/WfInstances	303	907	962.49 MB

Browsing, sorting, and selection of workflow instances and their metrics



Interactive graphical visualization of workflow instances as directed acyclic graphs

Solution:

- Implemented Features
- Browsable list of workflow instances
 - Viewable summarized workflow metrics
 - Sorting for workflow instances by metrics
 - Selection and download of workflow instances
 - Visualization of workflow instances using Cytoscape

Challenges:

- The project utilized a combination of old and new frameworks that were not compatible with each other at different points in the project
 - Documentation for framework compatibility was either outdated or did not exist due to the framework being too recent
- The structure of workflow instances required us to research a variety of different graph algorithms in order to construct and traverse a graph of nodes in a timely and efficient manner
- The project requirements were modified to include a visualization of the workflow instances, which required us to add the visualization framework Cytoscape into our tech stack in the middle of the development process

Learnings:

- Built a full-stack web application from the ground up with a TypeScript frontend and a Python backend that communicate via a custom REST API
- Learned differences between TypeScript and Javascript
- Used a data visualization framework