

HuBMAP Portal Team (as of September 2019)

What is NOT in the Portal Team Scope?

The scope does NOT include many necessary pieces to Data Release but depends upon success of:

Data Release Teams metadata and pipeline and Q/A choices

Data Production, Q/A

UUIDs, Provenance, Registration UI, Data Ingest

The scope includes “hooks” to integrate, in the UI, with other components:

Exploration UI, Launch of Visualizations of Data, Perhaps Launch of Analyses

What is the HuBMAP Portal Team Scope?

User Interface (UI) MAY include:

Amount, breadth, provenance and growth of data, Anatomic Diagrams of Available Tissues, Search (e.g. by gene), Faceted Search, Data Tables, Visualize Pipeline Results, Visualize Expression by Tissues, Preview/Viz Imaging Data

Application Programming Interfaces (APIs):

API Gateway, Provenance APIs, Query APIs, User Info API, Data Identity APIs, Download APIs, Metadata APIs, Data Retrieval APIs

Infrastructure (I):

Data, Provenance, and Metadata Stores, Workflow and Container

Who are HuBMAP Portal Team Members?



cborromeo



Nils Gehlenborg ngehlenborg



Jonathan C. Silverstein computationdoc



Philip Blood pdblood



Desheng Li etlds



pkant



Joel Welling jswelling



Bill Shirey shirey Maintainer



Chuck McCallum mccalluc



Sushma Anand Akoju sushma-ananda



Matt Ruffalo mruffalo Maintainer



Zhou (Joe) Yuan yuanzhou

How is the HuBMAP Portal Team Doing its Work?

Agile Software Development with Scrum:

Multi-institutional Scrum Daily

Sprint Review/Planning: Every Two Weeks

GitHub *data-portal* Project (private project for now)

Multiple Repositories

HuBMAP Consortium

Repositories 22 Packages People 46 Teams 6 Projects 2

data-portal Updated 3 hours ago

Filter cards + Add cards Fullscreen Menu

26 Backlog

- Server-side rendering? Back to plain Django?** (bug, fundamental, question)
- Top of page hidden by header** (UI, bug)
- Visualize data processing pipeline runs** (Blocked)
- Support simple visualization of (expression) data** (enhancement)
- Create production instance of Airflow on HuBMAP infrastructure** (Automated as To do)

5 Ready

- Error is thrown if the description field is too long (10 paragraphs of lorem ipsum)** (uuid-ui#29)
- Show graph of how much data (cells/experiments/images/etc.) we have, by tissue** (enhancement, front page)
- Implement experiment detail interface** (mrufralo)
- Show male/female human anatomical diagrams on front page** (front page)
- Implement searching by gene, to show in which tissues a gene is expressed** (mrufralo)

15 In Progress

- show details for a given entity on a page** (API, UI, enhancement)
- API Gateway development version** (API)
- Initial Query API** (API)
- list all entities for a given type on a page** (API, UI, enhancement)
- list all entity types on a page**

1 Review/QA

- Production deployment through Docker containers** (mrufralo)

Where can I see Data Release Deliverables? Asana

<https://app.asana.com/0/1132791370113704/board>

This board is NOT low-level management tool for individual workstreams

not Portal Team's GitHub just shown, not the DRTs' individual Asana board

This board is high-level tool for HuBMAP Members (everyone)

Status of other teams' deliverables your team may be dependent upon

Product Owners (one from each workstream) expected to maintain/edit

Peter Kant (IEC) project manages the Data Release Deliverables

asana ☰

- Home
- My Tasks
- Inbox
- Portfolios
- HIVE TC-Harvard
 - HA C PK ma dp
 - Create a Project
- Working Groups
 - HA ZB dg cw
 - Data Science Working Group
- Data Release Teams (...)
- Data Release Deliverables
- Mass Spectrometry Data R...
- Microscopy Data Release
- Sequencing Data Release
- TEST Data Release Deli...

Invite your team and start collaborating!

Data Release Deliverables

Board | Timeline | Calendar | Progress | Forms | More...

Search | + New | Upgrade | JS

Last task completed yesterday | All tasks | Filter | Sort

Backlog	To Do	In Progress	Done
<p>Pipeline C: Valid QA/QC Report</p> <p>Port...</p>	<p>Data storage</p> <p>PB</p> <p>Port...</p>	<p>Data Ingest Portal</p> <p>sh</p> <p>Port...</p>	<p>June 2020 Data release team structure confirmed / adjusted</p> <p>Sep 13</p>
<p>Pipeline C: Portal Integration</p> <p>Port...</p>	<p>Metadata storage</p> <p>PB</p> <p>Port...</p>	<p>Provenance Tracking</p> <p>sh</p> <p>Port...</p>	<p>Refined Mass Spectrometry modality flow shared with team</p> <p>JS Aug 22</p> <p>Mas...</p>
<p>Pipeline C: Unit tests & containerization</p> <p>MR</p> <p>Port...</p>	<p>Data download capability</p> <p>PB</p> <p>Port...</p>	<p>API Gateway</p> <p>sh</p> <p>Port...</p>	<p>UUID System for biospecimens deployed</p> <p>sh</p>
<p>Pipeline B: Valid QA/QC Report</p> <p>MR</p> <p>Port...</p>	<p>Automate IMS QA/QC report generation</p> <p>JS</p> <p>Mas...</p>	<p>Initial Query API</p> <p>sh</p> <p>Port...</p>	<p>Diagram</p> <p>Mas...</p>
<p>Pipeline B: Unit Test and Containerization</p> <p>MR</p> <p>Port...</p>	<p>Assay: DartFISH: v1 metadata definition & extraction</p> <p>Yesterday</p>	<p>Finalize IMS Instrumental Metadata output</p> <p>JS Dec 31</p> <p>Mas...</p>	<p>Proteomics QC report</p> <p>Mas...</p>