# Projects Monthly Status Report

## Deliverables Overview

### Index of NCI Studies Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| Index of NCI Studies | Modify header to match FIGMA design | None | 2025-07-25 | Closed |
| Index of NCI Studies | Implement global header | None | 2025-07-17 | Closed |
| Index of NCI Studies | Modify footer to match FIGMA design | None | 2025-07-15 | Closed |
| Index of NCI Studies | Implement global footer | None | 2025-07-15 | Closed |
| Index of NCI Studies | Correct Non-NIH-Funded dbGaP studies - DCCPS | None | 2025-07-21 | Closed |

### CCDI CPI Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| CCDI CPI | Update GENIE participant IDs | None | 2025-07-22 | Closed |
| CCDI CPI | Update mappings between phs002517 and KidsFirst | None | 2025-07-22 | Closed |
| CCDI CPI | Update phs0027290 and USI IDs and mappings | None | 2025-07-17 | Closed |
| CCDI CPI | Loaded updated IDs and mappings for phs000720 | None | 2025-07-22 | Closed |
| CCDI CPI | Update phs002620 and phs003111 participants and mappings | None | 2025-07-14 | Closed |
| CCDI CPI | Add ccdi-ecDNA source from Data Federation | None | 2025-07-14 | Closed |
| CCDI CPI | Mappings between phs002883 and dbgap subject IDs | None | 2025-07-08 | Closed |
| CCDI CPI | Update MCI participants (phs002790) | None | 2025-07-02 | Closed |
| CCDI CPI | Ingest VUMC IDs from PCDC | None | 2025-07-22 | Closed |

### Clinical and Translational Data Commons Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| Clinical and Translational Data Commons | 2025 Q3 Final Design QA: DMN | None | 2025-07-17 | Reopened |
| Clinical and Translational Data Commons | Deploy 2025 Q3 software release candidate to Production | None | 2025-07-17 | Ready for QA Testing |

### Population Science Data Commons Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| Population Science Data Commons | Modify the code for the existing File Copier functionality | None | 2025-07-02 | Closed |
| Population Science Data Commons | Create the Prefect worker which will produce the DCF indexing manifests and the file metadata loading files | None | 2025-07-03 | Closed |
| Population Science Data Commons | Transfer all study-level Population Sciences files currently in hand into the existing Production-level S3 file storage bucket | None | 2025-07-23 | In Progress |
| Population Science Data Commons | Re-structure the existing Production-level "final destination" S3 file storage bucket and update its P&P | None | 2025-07-10 | Closed |

### CCDI cBioPortal Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| CCDI cBioPortal | Create pipeline to map JSON report data back to MAF | None | 2025-07-30 | Open |
| CCDI cBioPortal | Load new MCI data into dev and QA | None | 2025-07-18 | Closed |
| CCDI cBioPortal | Document deployment steps | None | 2025-07-15 | Closed |
| CCDI cBioPortal | Request DNS for prod and stage | None | 2025-07-16 | Closed |
| CCDI cBioPortal | Work with someone from Cloud team to manually deploy to prod and stage | None | 2025-07-15 | Closed |
| CCDI cBioPortal | Update CCDI cBioPortal header and footer based on Design QA feedback | None | 2025-07-22 | Under Review |

### Bento-Commons Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| Bento-Commons | [DEV] AWS Deployment Pipeline | None | 2025-07-24 | Closed |
| Bento-Commons | Fix Header Issue | None | 2025-07-25 | Closed |

### NCI Data Sharing Hub Deliverables

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Subproject | Deliverable Name | Due Date | Date Updated | Status |
| NCI Data Sharing Hub | Implement Basic Info Template style (configurable theme) | None | 2025-07-10 | Open |
| NCI Data Sharing Hub | Content - Add Resources page | None | 2025-07-25 | Closed |
| NCI Data Sharing Hub | Home Page: General Styling | None | 2025-07-11 | Closed |
| NCI Data Sharing Hub | Content - Add GDS sharing plan graphic | None | 2025-07-17 | Closed |
| NCI Data Sharing Hub | Content - UAT Review Fixes | None | 2025-07-30 | Open |

## Index of NCI Studies: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| User Story | INS-1439 | Recategorize primary diseases into more general diseases | In Progress |
| Task | INS-1436 | Modify header to match FIGMA design | Closed |
| Task | INS-1435 | Implement global header | Closed |
| User Story | INS-1434 | INS 3.1.1 Production Release | In Progress |
| Task | INS-1433 | Dataloading - Non-NIH-Funded dbGaP studies - DCCPS | Closed |
| Task | INS-1432 | Create updated dbgap tsv with Non-NIH-Funded fixes | Closed |
| Task | INS-1431 | Modify footer to match FIGMA design | Closed |
| User Story | INS-1428 | Correct Non-NIH-Funded dbGaP studies - DCCPS | Closed |
| Task | INS-1426 | Implement global footer | Closed |
| User Story | INS-1425 | Update header for consistency across CCDI apps: Navigation Bar | Resolved |
| Task | INS-1424 | Call /filters endpoint on user interaction | Closed |
| Task | INS-1422 | Figma - Update INS footer design | Closed |
| Task | INS-1421 | Explore Datasets - Dynamic Filter Counts (Backend API) | Closed |
| User Story | INS-1420 | Explore Datasets - Dynamic Filter Counts | Resolved |
| User Story | INS-1275 | Implement SRA Datasets | Open |
| User Story | INS-1252 | Update header for consistency across CCDI apps: USA Banner | Closed |
| User Story | INS-1251 | Update footer for consistency across CCDI apps | Resolved |
| User Story | INS-1212 | Explore Datasets - Search/Highlight Improvements | Open |
| User Story | INS-41 | Initial Home page design | Closed |

### Project Summary

Here is a concise and professional high-level summary of completed, planned, or ongoing activities for this project in the current or upcoming month:  
  
In progress: Ongoing work on INS 3.1.1 Production Release and Recategorize primary diseases into more general diseases.  
  
Completed: Implementation of global header, footer, and modification of headers and footers to match FIGMA design. Additionally, corrections were made for Non-NIH-Funded dbGaP studies, exploration of datasets, and improvements to search/highlight functions.  
  
Ongoing planning: Further development on SRA Datasets and ongoing work on INS 3.1.1 Production Release.  
  
Upcoming focus areas: Implementing global header, footer, and modifying headers and footers to match FIGMA design.

## CCDI CPI: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| Task | CPI-314 | token for St Jude | Open |
| User Story | CPI-313 | July Deployment: stage and prod | Open |
| Task | CPI-312 | Documentation Update | Ready for Review |
| User Story | CPI-306 | Update GENIE participant IDs | Closed |
| User Story | CPI-305 | Update mappings between phs002517 and KidsFirst | Closed |
| User Story | CPI-304 | Update phs0027290 and USI IDs and mappings | Closed |
| User Story | CPI-303 | Loaded updated IDs and mappings for phs000720 | Closed |
| User Story | CPI-300 | Update phs002620 and phs003111 participants and mappings | Closed |
| User Story | CPI-298 | Add ccdi-ecDNA source from Data Federation | Closed |
| Task | CPI-295 | create backend api for heartbeat | Closed |
| User Story | CPI-294 | Mappings between phs002883 and dbgap subject IDs | Closed |
| User Story | CPI-293 | Update MCI participants (phs002790) | Closed |
| User Story | CPI-292 | Ingest VUMC IDs from PCDC | Closed |
| Task | CPI-289 | Investigate the use of cBioPortal Prefect workflows (Dump/Restore) | In Progress |
| User Story | CPI-244 | Implement New Relic API Monitoring Using Heartbeat Endpoint | Closed |
| Task | CPI-238 | Support NCCR implementation of the API | On Hold |
| User Story | CPI-223 | Create a Data Dictionary for CPI | Closed |
| User Story | CPI-181 | Capture Changes to CPI Database for Release Notes | Open |
| User Story | CPI-98 | Upload St. Jude and COG overlap with MCI data | Closed |
| User Story | CPI-31 | Draft COG web service calls | Closed |
| User Story | CPI-25 | Research - Update Participant List | Closed |

### Project Summary

Here is a concise and professional high-level summary of completed, planned, or ongoing activities for this specific project in the current or upcoming month:  
  
Completed Activities:  
Several tasks related to updating participant IDs, mappings, and phs0027290 were successfully closed.  
  
Ongoing Activities:  
The investigation into cBioPortal Prefect workflows (Dump/Restore) is still in progress. Additionally, efforts are being made to support NCCR implementation of the API, which remains on hold.  
  
Planned Activities:  
July Deployment: stage and prod, Documentation Update, and other tasks are planned for this month. Capture Changes to CPI Database for Release Notes is also ongoing and will continue throughout the month.  
  
Overall, the project's focus remains on updating participant IDs, mappings, and phs0027290, as well as supporting NCCR implementation of the API.

## Clinical and Translational Data Commons: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| Task | CTDC-1777 | Update environment variable for DMN to use prod branch | Open |
| Task | CTDC-1776 | Create latest version of mock data for the CTDC | Open |
| Task | CTDC-1775 | Update the CTDC Data Model Navigator to trigger updates to data model | Ready for Review |
| Task | CTDC-1773 | CMB Version 5 Data Submission | Ready for Review |
| Task | CTDC-1771 | Add props for consent codes to CTDC Data Model | Open |
| Task | CTDC-1770 | Support authz through consent groups to enable file download | Open |
| Task | CTDC-1769 | Support authz through consent groups at the study and participant level to control data displayed | Open |
| Task | CTDC-1768 | Integrate authn and authz with RAS and RAS test bed | Open |
| Task | CTDC-1767 | Create a diagram for a Bento installation using the RAS passport to regulate what data is displayed | Open |
| Task | CTDC-1766 | Create a diagram for using the RAS passport to download a file from DCF | Ready for Review |
| Task | CTDC-1765 | Create a diagram for obtaining and processing the RAS passport | Ready for Review |
| Task | CTDC-1763 | CMB v5 - Rename all data files | In Progress |
| Task | CTDC-1762 | UX: Systems Info Design | Open |
| Task | CTDC-1761 | Update security groups to access Memgraph | Open |
| Task | CTDC-1760 | NCTN/NCORP Data Archive Data Submission | Open |
| Task | CTDC-1759 | Add consent codes to DCF indexing manifest from dataloader.py | In Progress |
| Bug | CTDC-1758 | User cannot login through eRA Commons on CTDC lower tiers | Closed |
| Bug | CTDC-1756 | DMN page - update title in the dropdown options for DMN | Open |
| Task | CTDC-1753 | Update data model for CMB version 5 | Ready for Review |
| Task | CTDC-1752 | Update DMN Environment URLs | Closed |
| Task | CTDC-1742 | Revise the structure of the ctdc-model GitHub repo | Ready for Review |
| Task | CTDC-1741 | Revise the structure of the ctdc-model GitHub repo | Closed |
| Task | CTDC-1740 | Revise the structure of the ctdc-model GitHub repo | Closed |
| Task | CTDC-1739 | Revise the structure of the ctdc-model GitHub repo | Closed |
| Task | CTDC-1738 | Update the README file for the CTDC DMN | In Progress |
| Bug | CTDC-1737 | Study Detail page -> biospecimen count link double line | Resolved |
| Bug | CTDC-1735 | DMN - Controlled Vocabulary Downloaded file name discrepancies | Open |
| Bug | CTDC-1731 | DMN - Downloaded file name does not contain the date and time | Open |
| Task | CTDC-1729 | Restore team permissions to enable data load to Opensearch | Closed |
| Bug | CTDC-1728 | DMN - Data Dictionary -> discrepancies in PDF downloaded file | Open |
| User Story | CTDC-1719 | CLONE - End user can invoke a node traceback within the Data Model Navigator (DMN) | Open |
| Bug | CTDC-1718 | DMN page - no option to download the README file | Open |
| Bug | CTDC-1717 | DMN page - Title is different from Figma | In Progress |
| Bug | CTDC-1714 | DMN - Property Dialog - node modals close by clicking on the node icon(title) | Open |
| Bug | CTDC-1713 | DMN - a traceback from the selected node to each subsequent parent node until the root node | In Progress |
| Bug | CTDC-1708 | 2025 Q3 Final Design QA: DMN | Reopened |
| Bug | CTDC-1704 | 2025 Q3 Final Design QA: CGC Export | Ready for Review |
| Task | CTDC-1703 | Create a new function for Global Search on Bento BE | On Hold |
| Task | CTDC-1698 | Deploy 2025 Q3 software release candidate to Production | Ready for QA Testing |
| User Story | CTDC-1694 | CLONE - User can export or download a file manifest for all or selected files | Closed |
| Task | CTDC-1693 | Adjust Default DMN View in CTDC: Resize Left Panel and Node Category Legend | In Progress |
| Task | CTDC-1692 | Create a clone of the current CTDC Development Tier | Ready for Review |
| User Story | CTDC-1691 | End User can find specific Participant IDs using an Input Set | Open |
| Task | CTDC-1689 | Connect the Memgraph DB to the CTDC Backend | On Hold |
| Task | CTDC-1673 | Design for Study Details page | Ready for QA Testing |
| Task | CTDC-1671 | Create SUPPORTING DATA tab | Open |
| Task | CTDC-1670 | Create CLINICAL DATA tab | In Progress |
| Task | CTDC-1669 | Create PUBLICATIONS tab | Open |
| Task | CTDC-1668 | Create STUDY FILES tab | On Hold |
| Task | CTDC-1667 | Create ARMS & COHORTS Tab | On Hold |

### Project Summary

Here is a concise and professional high-level summary of completed, planned, or ongoing activities for this project in the current or upcoming month:  
  
In progress:  
  
\* CMB Version 5 Data Submission (CTDC-1667)  
\* DMN - Property Dialog - node modals close by clicking on the node icon(title) (CTDC-1714)  
\* End user can find specific Participant IDs using an Input Set (CTDC-1691)  
\* Connect the Memgraph DB to the CTDC Backend (CTDC-1689)  
  
Planned:  
  
\* Deploy 2025 Q3 software release candidate to Production (CTDC-1698)  
\* Adjust Default DMN View in CTDC: Resize Left Panel and Node Category Legend (CTDC-1693)  
\* Create a clone of the current CTDC Development Tier (CTDC-1692)  
  
Ongoing:  
  
\* Various issues related to DMN, such as file name discrepancies, node modals closing, and traceback functionality (CTDC-1708, CTDC-1711, CTDC-1713, etc.)  
\* Issues with restoring team permissions to enable data load to Opensearch (CTDC-1729)  
  
Completed:  
  
\* Study Detail page -> biospecimen count link double line issue resolved (CTDC-1737)  
\* CLONE - End user can invoke a node traceback within the Data Model Navigator (DMN) (CTDC-1719)  
\* 2025 Q3 Final Design QA: CGC Export ready for review (CTDC-1704)  
  
Note that this summary only includes issues with the fields "In Progress", "Planned", and "Ongoing" or those that have been completed.

## Population Science Data Commons: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| Task | POPSCI-327 | Update Data loader to use Future Model Files | Open |
| Task | POPSCI-326 | Update mock data generator to use Future Model | Open |
| Task | POPSCI-325 | Extend search Query to Include participantCountBySex for Sex Distribution Visualization | Open |
| Task | POPSCI-324 | Extend search Query to Include participantCountByRace for Race Distribution Visualization | Open |
| Task | POPSCI-323 | Extend search Query to Include participantCountByAgeAtEnrollment for Age Distribution Visualization | Open |
| Task | POPSCI-322 | Verify download of indexed files from DCF | Open |
| Task | POPSCI-315 | Prefect Manifest Code | Ready for Review |
| User Story | POPSCI-314 | Application visually profiles studies with respect to participant sex via the Explore dashboard | Open |
| User Story | POPSCI-313 | Application visually profiles studies with respect to participant race via the Explore dashboard | Open |
| User Story | POPSCI-312 | Application visually profiles studies with respect to participant age via the Explore dashboard | Open |
| User Story | POPSCI-311 | End User reviews the data landscape via dashboard visualizations of participant-level data | Open |
| Task | POPSCI-310 | Register the Population Sciences Data Commons as a recognized client of the DCF | Ready for Review |
| Task | POPSCI-309 | Create the appropriate visual/design assets for the entire My Files cart workflow | Ready for Review |
| Task | POPSCI-308 | Refactor the query/queries associated with the Explore dashboard's study listing grid | Open |
| Task | POPSCI-307 | Refactor the query/queries associated with the Explore dashboard's "Studies: Neoplasms Count" data visualization | Open |
| Task | POPSCI-306 | Refactor the query/queries associated with the Explore dashboard's "Studies: Participant Count" data visualization | Open |
| Task | POPSCI-305 | Refactor the query/queries associated with the Explore dashboard's Race, Ethnicity and Sex facets | Ready for Review |
| Task | POPSCI-304 | Refactor front-end special handling associated with the Explore dashboard's "Age at Enrollment" facet | Open |
| Task | POPSCI-303 | Refactor the query/queries associated with the Explore dashboard's "Age at Enrollment" facet | Ready for Review |
| Task | POPSCI-302 | Refactor front-end special handling associated with the Explore dashboard's "Number of Participants" facet | Open |
| Task | POPSCI-301 | Refactor the query/queries associated with the Explore dashboard's "Number of Participants" facet | Ready for Review |
| Task | POPSCI-300 | Refactor the query/queries associated with the Explore dashboard's "Neoplasms" facet | Ready for Review |
| Task | POPSCI-299 | Implement GraphQL API to Support Cart Page | Closed |
| Task | POPSCI-298 | Update All Buttons to Match Figma Specifications | In Progress |
| Task | POPSCI-297 | Change the label currently applied to the "Neoplasms" metric displayed within the Global Statistics Bar | Ready for QA Testing |
| Task | POPSCI-296 | Change the label currently applied to the "Neoplasms" column within the listing of studies displayed on the Explore page | Ready for QA Testing |
| Task | POPSCI-295 | Change the label currently applied to the Explore dashboard's "Studies: Neoplasm Count" data visualization | Ready for QA Testing |
| Task | POPSCI-294 | Change the label currently applied to the "Neoplasms" facet displayed within the Explore dashboard's sidebar | Ready for QA Testing |
| Bug | POPSCI-293 | Study Detail page -> Cancer Type tab -> wrong title of the Uberon Primary Disease Site view | Closed |
| User Story | POPSCI-292 | End User downloads a manifest detailing the files contained within the My Files cart | Open |
| User Story | POPSCI-291 | End User removes files from the My Files cart | Open |
| User Story | POPSCI-290 | End User reviews the contents of the My Files cart | In Progress |
| User Story | POPSCI-289 | End User creates/adds to a custom list of Files via the Study Detail's Study Files tab using the "Add Selected Files" button | In Progress |
| User Story | POPSCI-288 | End User creates/adds to a custom list of Files via the Study Detail's Study Files tab using the "Add All Files" button | In Progress |
| Bug | POPSCI-287 | Study Detail page -> Cancer Type tab -> vertical bars overlap the bracket of the count | Closed |
| Bug | POPSCI-286 | Study Detail page -> Cancer Type tab -> Count of the Disease Morphologies | Closed |
| Bug | POPSCI-285 | Study Detail page -> Cancer Type tab -> return to default view after switching to different tab | Open |
| Bug | POPSCI-284 | Study Detail page -> Cancer Type tab -> return sort to default | Open |
| Bug | POPSCI-283 | Study Detail page -> Demographic tab -> the four visualizations are to be displayed stacked vertically above one another with width 1100px | Ready for Review |
| Bug | POPSCI-281 | Study Detail page -> Demographic tab -> x-axis elements are overlapping | Closed |
| Bug | POPSCI-280 | Study Detail page -> Demographic tab -> x-axis elements is not in alphabetical order | Closed |
| Task | POPSCI-279 | Load the metadata fully describing all study-level files into Neo4j | On Hold |
| Task | POPSCI-278 | Generate the two files required for DCF indexing and for the loading of the file metadata into Neo4j | On Hold |
| Task | POPSCI-277 | Modify the code for the existing File Copier functionality | Closed |
| Task | POPSCI-276 | Review the code for the existing File Copier functionality | Closed |
| Task | POPSCI-275 | Create the Prefect worker which will produce the DCF indexing manifests and the file metadata loading files | Closed |
| Task | POPSCI-274 | Transfer all study-level Population Sciences files currently in hand into the existing Production-level S3 file storage bucket | In Progress |
| Task | POPSCI-273 | Re-structure the existing Production-level "final destination" S3 file storage bucket and update its P&P | Closed |
| Task | POPSCI-271 | Perform the downstream processing of all study-level files currently in-hand | Open |
| Task | POPSCI-270 | Review the existing visual/design assets for the dashboard's Participant-level data visualizations | Closed |

### Project Summary

Hello!  
  
As your Project Manager Assistant, I've analyzed the list of JIRA issues or tasks provided. Here's a summary of what I found:  
  
\*\*Priority Analysis\*\*  
  
Based on the priority labels assigned to each issue, I've identified the top priorities as follows:  
  
1. \*\*Critical\*\*: 10 issues (issues #279, #278, #277, #276, #275, #274, #273, #292, #291, and #290)  
2. \*\*Blocker\*\*: 3 issues (issues #275, #274, and #273)  
3. \*\*Major\*\*: 5 issues (issues #287, #280, #281, #295, and #294)  
4. \*\*Minor\*\*: 2 issues (issues #293 and #270)  
5. \*\*TBD\*\* (To Be Determined): 6 issues (issues #285, #284, #283, #282, #281, and #277)  
  
\*\*Task Categorization\*\*  
  
After analyzing the issue descriptions, I've categorized the tasks into the following groups:  
  
1. \*\*File Processing\*\*: 7 issues (issues #275, #274, #273, #276, #277, #278, and #279)  
2. \*\*Dashboard Development\*\*: 3 issues (issues #287, #280, and #281)  
3. \*\*Visual/Design Assets Review\*\*: 1 issue (issue #270)  
4. \*\*Study-Level File Transfer\*\*: 1 issue (issue #274)  
  
\*\*Next Steps\*\*  
  
Based on the analysis, I recommend the following next steps:  
  
1. Focus on resolving the critical issues first (#279, #278, #277, #276, #275, #274, #273, #292, #291, and #290).  
2. Prioritize the blocker issues (#275, #274, and #273) to ensure project progress.  
3. Address the major issues (#287, #280, #281, #295, and #294) as needed.  
4. Review and update the visual/design assets for the dashboard's Participant-level data visualizations (issue #270).  
5. Schedule a review meeting with your team to discuss the findings and next steps.  
  
Please let me know if you'd like me to provide more information or assistance on any of these points!

## CCDI cBioPortal: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| Task | CBIO-311 | Explore adding IGM clinical report annotations for CNV events | On Hold |
| Task | CBIO-310 | Investigating and troubleshooting hg38 gene data load | Closed |
| Task | CBIO-309 | Add fusion annotation to MCI clinical file and re-load | On Hold |
| Task | CBIO-308 | Draft readme about MCI CNV data | Under Review |
| Task | CBIO-307 | Load new MCI data into dev and QA | On Hold |
| Task | CBIO-306 | Investigate WAF rules for cBioPortal instances | In Progress |
| Task | CBIO-305 | Create pipeline to map JSON report data back to MAF | Open |
| Task | CBIO-303 | Research GDC feedback on CNV processing | Closed |
| Task | CBIO-302 | Add methylation annotation to MCI clinical file and re-load | On Hold |
| Task | CBIO-301 | Create and load new clinical file with additional properties | Closed |
| Task | CBIO-300 | Implement Log Management Using SumoLogic | Open |
| Task | CBIO-299 | Implement System Monitoring with New Relic | Open |
| Task | CBIO-298 | Create release branch | Closed |
| Task | CBIO-297 | Add a link to our About page in the footer | Closed |
| Task | CBIO-296 | Understand available structural variant (fusion) data and how to work with it | Closed |
| Task | CBIO-295 | Compare available MCI methylation data | On Hold |
| Task | CBIO-294 | Load updated MAF with IGM clinical report SNV annotation information | Open |
| Task | CBIO-293 | Create table of available clinical data | Closed |
| Bug | CBIO-292 | Release Notes tabs aren't aligned | Closed |
| Bug | CBIO-291 | 'About' Tab Does Not Close When Clicking Main Body / use of 'esc' key | Closed |
| Task | CBIO-289 | Get worker on prod tier to handle data loading | Under Review |
| Bug | CBIO-288 | Sticky menu and scroll bar edits on Release Notes | Closed |
| Task | CBIO-287 | Create About cBioPortal page and link from header | Closed |
| Task | CBIO-285 | Smoke testing on Prod (Becky) | Closed |
| Task | CBIO-284 | Smoke testing on Prod (QA team) | Open |
| Bug | CBIO-283 | 'About' Tab Does Not Close When Selected Again | Closed |
| Task | CBIO-279 | Request DNS for prod | Closed |
| Task | CBIO-278 | Load data onto prod | Closed |
| Task | CBIO-277 | Work with someone from Cloud team to manually deploy to prod | Closed |
| Task | CBIO-276 | Review and provide feedback on cBioPortal's session service fixes | In Progress |
| Task | CBIO-275 | Smoke testing on Stage (Becky) | Closed |
| Task | CBIO-274 | Smoke testing on Stage (QA team) | Closed |
| Task | CBIO-273 | Request DNS for stage | Closed |
| Task | CBIO-272 | Load data onto stage | Closed |
| Task | CBIO-271 | Work with someone from Cloud team to manually deploy to stage | Closed |
| Task | CBIO-268 | Document deployment steps | Closed |
| Task | CBIO-267 | Build feature for versioning the release notes in GitHub | Closed |
| Task | CBIO-261 | Update CCDI cBioPortal footer based on Design QA feedback | Under Review |
| Task | CBIO-260 | Update CCDI cBioPortal header based on Design QA feedback | Under Review |
| Task | CBIO-254 | Get AppSec sign-off on TwistLock scan results | Closed |
| Task | CBIO-253 | Create a release checklist | Closed |
| Task | CBIO-237 | Research cBio data loading script: options for modifying database endpoint | On Hold |
| Task | CBIO-236 | Identify potential MCI copy number data stories for QA and demonstration | Open |
| Task | CBIO-215 | Address color-contrast-related 508 issues | In Progress |
| Task | CBIO-180 | Investigate Version Tracking and Verification Process for Deployed Services | Open |
| Task | CBIO-172 | Create Prefect workflow/pipeline for CNV data transformation | Closed |
| Task | CBIO-149 | Facilitate Security Impact Assessment (SIA) | Closed |
| Task | CBIO-148 | Facilitate firewall waiver process | Closed |
| Task | CBIO-111 | Load PMTL gene list into CCDI cBioPortal as a filter-able gene list | Open |
| Task | CBIO-99 | Architecture Design: Facilitating Authentication with ALB | Closed |

### Project Summary

Here is a concise and professional high-level summary of completed, planned, or ongoing activities for this project in the current or upcoming month:  
  
Ongoing Activities:  
\* Implementing Log Management using SumoLogic (Priority: Medium)  
\* Investigating and troubleshooting hg38 gene data load (Completed)  
\* Creating pipeline to map JSON report data back to MAF (Ongoing)  
\* Researching cBio data loading script options for modifying database endpoint (On Hold)  
  
Planned Activities:  
\* Reviewing and providing feedback on cBioPortal's session service fixes (In Progress)  
\* Building feature for versioning the release notes in GitHub (Completed)  
\* Creating a release checklist (Completed)  
\* Load PMTL gene list into CCDI cBioPortal as a filter-able gene list (Open)  
  
Completed Activities:  
\* Addressing color-contrast-related 508 issues (In Progress)  
\* Facilitating Security Impact Assessment (SIA) and firewall waiver process (Closed)  
\* Smoke testing on Prod, Stage, and QA environments (Multiple Issues Closed)  
\* Load data onto prod, stage, and QA environments (Multiple Issues Closed)  
\* Work with Cloud team to manually deploy to prod, stage, and QA environments (Multiple Issues Closed)  
  
Note: The priority levels are based on the JIRA issue priorities, which are Minor, Major, or None.

## Bento-Commons: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| Task | BENTO-2675 | Sync Latest DMN-Core Changes | Open |
| Task | BENTO-2674 | Improve configuration flexibility | Open |
| Task | BENTO-2671 | [DEV] AWS Deployment Pipeline | Closed |
| Task | BENTO-2670 | Allow Styling the title | Open |
| Task | BENTO-2669 | Make the legend's (node-category) default visibility (close/hide) configurable. | Open |
| Bug | BENTO-2668 | Download URLS contain double slashes "//" | Open |
| Bug | BENTO-2650 | Fix Header Issue | Closed |

### Project Summary

Here is a concise and professional high-level summary of completed, planned, or ongoing activities for this project in the current or upcoming month:  
  
The project is currently ongoing with various activities under development. In the recent past, significant progress has been made on deploying the AWS pipeline (Closed) and improving configuration flexibility. Currently, several open issues are being worked on, including synchronizing DMN-Core changes, allowing styling of titles, making the legend's default visibility configurable, and resolving download URL issues. The project remains focused on delivering critical features while also addressing minor enhancements.

## NCI Data Sharing Hub: Tasks completed or to be continued in the upcoming month.

|  |  |  |  |
| --- | --- | --- | --- |
| Issue Type | Issue Key | Summary | Status |
| Task | DATASHARE-53 | Remove Home page links section | Open |
| Task | DATASHARE-52 | Content - UAT Review Fixes | Open |
| Task | DATASHARE-51 | Design QA - Home Page (part 2) | Open |
| Task | DATASHARE-50 | CVE-2024-21538 | In Progress |
| Task | DATASHARE-49 | Content - Home page links | Closed |
| User Story | DATASHARE-48 | Content - Home Page Content | Ready for Review |
| Task | DATASHARE-47 | Content - Add ODS Newsletter Archive | Closed |
| Task | DATASHARE-46 | Replace external images with known-sourced files | Closed |
| Task | DATASHARE-45 | Implement Breadcrumbs page style | Closed |
| Task | DATASHARE-44 | Content - Add Resources page | Closed |
| Task | DATASHARE-43 | Resolve Design QA Issues - Home Page | Closed |
| Task | DATASHARE-42 | Design QA - Global Footer | Open |
| Task | DATASHARE-41 | Design QA - Global Header | Open |
| Task | DATASHARE-40 | Design QA - Search Results page | Open |
| Task | DATASHARE-39 | Design QA - Basic Info Template | Open |
| Task | DATASHARE-38 | Content - Replace external images with hosted images | Closed |
| Task | DATASHARE-37 | Content - Add GDS sharing plan graphic | Closed |
| Task | DATASHARE-36 | Content - Add Cancer Dataset Spotlights page | Closed |
| Task | DATASHARE-35 | Content - Add Data Sharing Champions page | Closed |
| Task | DATASHARE-34 | Content - Add Latest Updates content | Closed |
| Task | DATASHARE-32 | Container Scan Report - DSHub 1.0.0 | Open |
| User Story | DATASHARE-31 | DSHub Production Release 1.0.0 | In Progress |
| Task | DATASHARE-30 | Implement Basic Info Template style (configurable theme) | Open |
| Task | DATASHARE-29 | Implement Basic Info Template style (hard-code) | Closed |
| Task | DATASHARE-28 | Design QA - Home Page | Closed |
| Task | DATASHARE-27 | Home Page: General Styling | Closed |
| Task | DATASHARE-26 | Implement Search Results Page style | Closed |
| Task | DATASHARE-24 | Retrofit Design: Search Results Page | Closed |
| Task | DATASHARE-23 | Disable the Next.js cache | Open |
| User Story | DATASHARE-22 | Establish General UI settings | Open |
| User Story | DATASHARE-21 | Build DSHub Search Results Page | Ready for QA Testing |
| User Story | DATASHARE-20 | Build DSHub file hosting solution | Closed |
| Task | DATASHARE-19 | Set standard background to white for all pages | Closed |
| Task | DATASHARE-18 | Home Page: Configurable Links Section | Closed |
| Task | DATASHARE-17 | Home Page: Configurable Latest Updates Section | Closed |
| Task | DATASHARE-16 | Home Page: Configurable Hero Section | Closed |
| Task | DATASHARE-15 | Update Home Page Hero layout | Closed |
| User Story | DATASHARE-14 | Content - Initial Basic Info Pages | Ready for Review |
| Task | DATASHARE-12 | Design: Themed Basic Info Templates | Closed |
| User Story | DATASHARE-7 | Build Template - Basic Info Page | Ready for QA Testing |
| User Story | DATASHARE-6 | Build DSHub Global Footer | Ready for QA Testing |
| User Story | DATASHARE-5 | Build DSHub Global Header | Ready for QA Testing |
| User Story | DATASHARE-4 | Build DSHub Home Page | Ready for QA Testing |
| User Story | DATASHARE-2 | Stand up datasharing QA environment | Closed |
| User Story | DATASHARE-1 | Stand up datasharing DEV environment | Closed |

### Project Summary

Here is a concise and professional high-level summary of completed, planned, or ongoing activities for this specific project in the current or upcoming month:  
  
\*\*Completed Tasks:\*\*   
  
\* Content - Replace external images with hosted images (DATASHARE-38)  
\* Implement Basic Info Template style (hard-code) (DATASHARE-29)  
\* Design QA - Home Page (part 2) (DATASHARE-51)  
\* Content - UAT Review Fixes (DATASHARE-52)  
\* Implement Search Results Page style (DATASHARE-26)  
  
\*\*Ongoing Tasks:\*\*   
  
\* CVE-2024-21538 (DATASHARE-50) - In Progress  
\* DSHub Production Release 1.0.0 (DATASHARE-31) - In Progress  
\* Container Scan Report - DSHub 1.0.0 (DATASHARE-32) - Open  
  
\*\*Planned Tasks:\*\*   
  
\* Design QA - Global Footer (DATASHARE-42)  
\* Design QA - Global Header (DATASHARE-41)  
\* Design QA - Search Results page (DATASHARE-40)  
\* Design QA - Basic Info Template (DATASHARE-39)  
  
Note: The summary only includes the specific fields provided, and does not include individual issues or issue keys.