

*User Guide*

## Feature Manager

Web UI for managing Quantitative Image Biomarkers

## Introduction

This is the user guide for feature-manager, a React web application to manage quantitative image biomarker (QIB) collections. This application is designed to consume a Flask-based REST API, whom code can be found in the following link:

<https://github.com/genttunn/python-rest-api>

The application has 3 main functions, which corresponds to 3 application pages:

1. Loading, viewing, and managing QIB collections: Grid page.
2. Visualizing QIB data into bivariate plot: Plot page.
3. Managing QIB metadata: Database page.

## Grid: Load and View and QIBs

The screenshot shows the 'Feature Manager' application with the 'Grid' tab selected. The interface is divided into a sidebar on the left and a main content area on the right. The sidebar contains a list of QIBs with details like Name, Description, Date, and outcome\_column. The main content area displays a 'QIB Table' with columns for PatientName, pic\_status, Modality, ROI, Series\_region, and original\_firstorder\_10Percentile. The table is sorted by PatientName. The top navigation bar includes tabs for Feature Manager, Grid, Plot, and Database. The sidebar has buttons for 'By Album' and 'By Date'. The main table has buttons for 'Upload QIB', 'Filter selected rows', 'Tag columns', and 'Current Export Mode: All'. The table rows are numbered 1 through 7, corresponding to the callouts in the list below.

Actions	PatientName	pic_status	Modality	ROI	Series_region	original_firstorder_10Percentile
<input type="checkbox"/>	Tom_1	0	PT	GTV_N	365	0.161817
<input type="checkbox"/>	PatientLC_10	1	PT	GTV_N	366	0.613304
<input type="checkbox"/>	PatientLC_11	0	PT	GTV_N	367	0.178194
<input type="checkbox"/>	PatientLC_12	1	PT	GTV_N	368	0.250006
<input type="checkbox"/>	PatientLC_13	1	PT	GTV_N	369	0.465724
<input type="checkbox"/>	PatientLC_14	1	PT	GTV_N	370	0.383604
<input type="checkbox"/>	PatientLC_15	1	PT	GTV_N	371	0.755934

1. Left scrollable column shows a list of QIBs in database. To sort them, click on By Album or By Date.
2. Click on Load to load the QIB to the table on the right (loaded QIB is bolded), Edit to change its metadata information, and Delete to remove it.
3. Click Upload QIB to upload a new CSV file. CSV files, depending on CSV type, will be checked for valid column names (see end of guide).
4. Hover mouse over a column's name too reveal a sorting arrow. Typing in Search and in each column Filter field will also allow for searching by name.
5. Click on tag columns to tag outcome columns and metadata columns, which will turn red and blue, respectively.
6. Exporting QIBs to CSV: click the checkboxes next to rows and **Filter selected rows** to select rows, click to choose columns, and to export.
7. Click on the moon symbol to switch between Light and Dark theme. The theme will persist in other tabs and next sessions.

## Plot: Visualize QIBs



1. Upper row shows general statistics of the database.
2. To make a plot, first choose the QIB.
3. Then select two features and click Make Plot.
4. Hover mouse over data point to see coordinate values.

## Database: Edit metadata entities



Click on each tab in the left column to manage Albums, Patient, Modality & Region of Interest, and Feature Families & Features.

## Appendix

Valid CSV by Type:

PatientID	Modality	ROI	original_fi
PatientLC_1_201601	CT	GTV_N	-866.528

*Valid new QIB file*

PatientName	plc_status	Modality	ROI	Series_region	original_first	original_fi
PatientLC_On	0	CT	GTV_L	1	82.1093	-711.881

*Valid custom QIB file*

patient_id	is_chuv	plc_status
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*Valid outcome list file*

Examples valid CSV files can be found in the backend application python-rest-api, in /csv.