

# RaMP-DB 3.0 Relational Database of Metabolomics Pathways

Workshop: An Introduction to NCATS Public Resources and Analytics for Rare Diseases, Targets, Drug Substances, and Analytes



## **Disclosure and Learning Objectives**

• I have no relevant relationships with commercial interests to disclose.

• Provide details on the data provenance for our resources, so that our users understand how the data could be useful for their work.

• Provide hands on experience interacting with our resources in various ways (e.g. web applications, APIs, or other packages).

• Interact and discuss how the data can be used, augmented, and further incorporated with other efforts. We anticipate new collaborative opportunities through the workshop.

### RaMP-DB: Comprehensive Metabolite/Gene Annotations

Up-to-date and comprehensive biological, chemical, ontology annotations from multiple sources

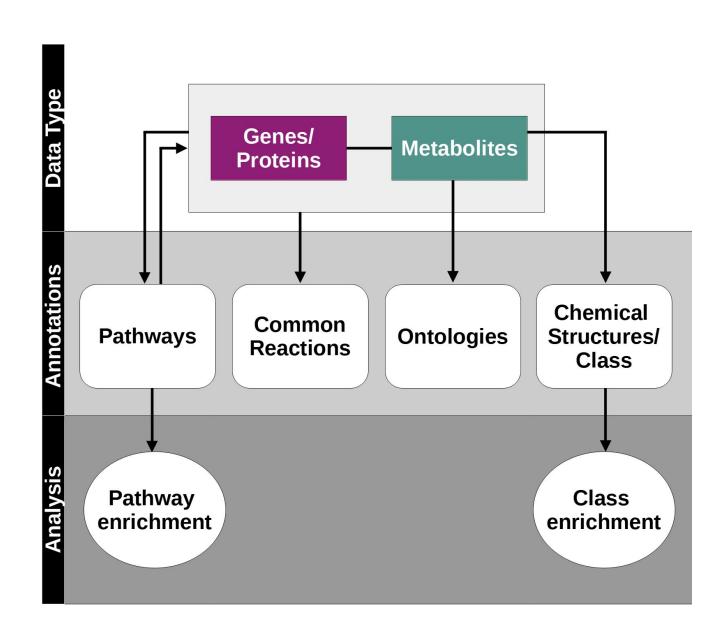
**Open-source**, stand-alone usable database

Multiple access points through the R package, SQLite database, APIs, interactive web app to perform queries and pathway analyses

**Transparent and reproducible process**, including semi-automated curation, code, and web app

Basis for improving pathway analysis methods

rampdb.nih.gov/



## RaMP-DB 3.0 - Major Improvements

- Expanded content
  - Source database updates
  - Increased focus on lipid annotations
  - Incorporation of curated reactions from Rhea
- Improved semi-automated integration scripts
- SQLite version (replacing MySQL)
   Greatly facilitates R package installation and integration with other tools (COMETS Analytics, MetaboAnalyst, Bioconductor Computational Metabolomics Ecosystem)
- New queries of biochemical reactions

# RaMP-DB 3.0 – Increased Annotations and Pathway Mappings

>230K metabolites, ~16K genes/enzymes, >53K pathways

| Total   | HMDB                                   | KEGG*   | Reactome   | WikiPathways   | LIPIDMAPS  | Rhea  |
|---------|--|---|--|--|--|---|
| 234,543 | 196,426                                | 5,578   | 2,239  | 3,569  | 39,368   | 12,155  |
| 16,443  | 7,073                                  |   | 11,229   | 13,841   |  | 3,498   |
| 53,952  | 49,613                                 | 363   | 2,627  | 1,349  |  |   |
| 412,990 | 367,370                                | 1,673   | 30,410   | 13,537   |  |   |
| 406,132 | 208,211                                | 8,459   | 127,077  | 62,385   |  |   |
|         | 234,543<br>16,443<br>53,952<br>412,990 | 234,543 196,426<br>16,443 7,073<br>53,952 49,613<br>412,990 367,370 | 234,543       196,426       5,578         16,443       7,073         53,952       49,613       363         412,990       367,370       1,673 | 234,543       196,426       5,578       2,239         16,443       7,073       11,229         53,952       49,613       363       2,627         412,990       367,370       1,673       30,410 | 234,543       196,426       5,578       2,239       3,569         16,443       7,073       11,229       13,841         53,952       49,613       363       2,627       1,349         412,990       367,370       1,673       30,410       13,537 | 234,543       196,426       5,578       2,239       3,569       39,368         16,443       7,073       11,229       13,841         53,952       49,613       363       2,627       1,349         412,990       367,370       1,673       30,410       13,537 |

<sup>\*</sup> From HMDB

## RaMP-DB 3.0 – Chemical Descriptions and Reactions (New)

### > 63K reactions, > 250K chemical property records

|                              | Rhea             |
|------------------------------|------------------|
| Reactions                    | 63,396 (15,849)  |
| Metabolite-Reaction Mappings | 299,272 (74,818) |
| Protein-Reaction Mappings    | 16,436 (4,109)   |

|                            | Total Distinct Compounds     | HMDB    | ChEBI  | LIPIDMAPS |
|----------------------------|------------------------------|---------|--------|-----------|
| Chemical Properties        | 256,537 (distinct InChlKeys) | 217,776 | 18,471 | 45,227    |
| Chemical Class Information | 155,293                      | 126,101 |        | 39,368    |
| Metabolite Ontology        | 134,653                      | 134,653 |        |           |

# New SQLite greatly improves accessibility and facilitates usage

### **Original MySQL Installation Process**

Install the RaMP R Package

Download the MySQL RaMP Data File

Download the MySQL Software and Install

Start the MySQL Server

Load the RaMP Data into MySQL Server

Load the RaMP Package

Establish RaMP to MySQL
Connection

Run RaMP Queries

### **New Simplified Process**

Install the RaMP R Package

Load the RaMP Package

Run RaMP Queries

RaMP 3.0 automatically downloads the SQLite Database file and stores it in a local file cache.

A RaMP function allows users to see and select RaMP DB versions on their computer OR in our RaMP-DB GitHub Large File Storage System.

## **Newly incorporate Biochemical Reactions**

**Input**: list of metabolite and gene/protein ids

Output: Rhea reactions, including

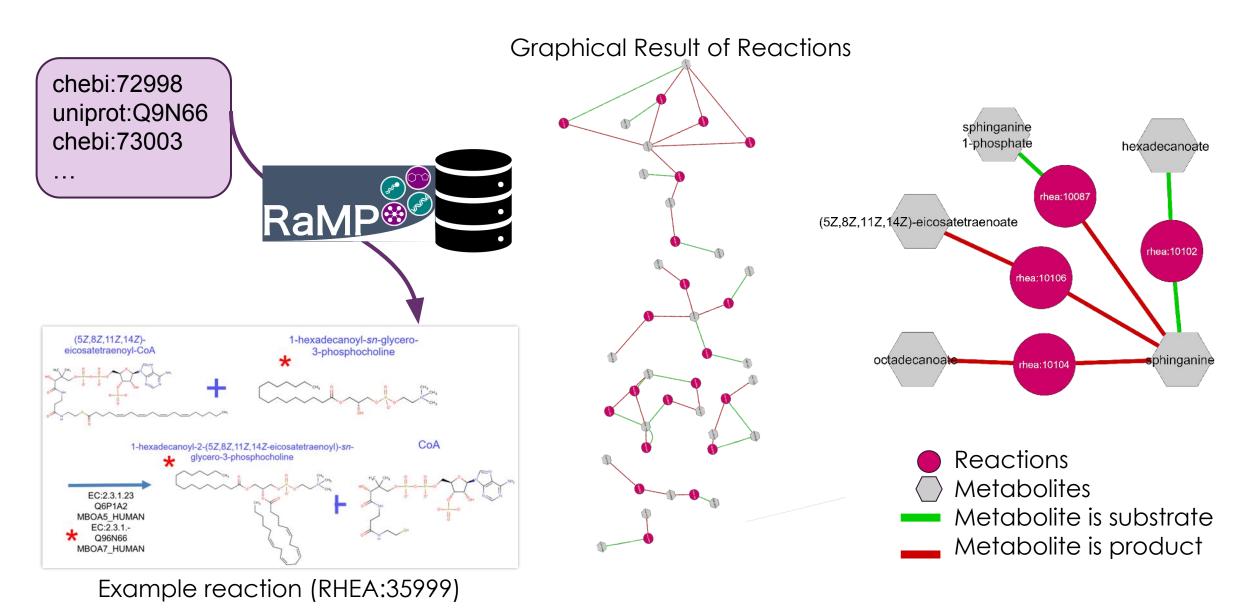
- Associated mappings to input metabolites
- Associated mappings to input genes/proteins

```
getReactionsForAnalytes(
   db = RaMP(),
   analytes,
   analyteType = "metabolites",
   namesOrIds = "ids",
   onlyHumanMets = F,
   humanProtein = F,
   includeTransportRxns = F,
   rxnDirs = c("UN")
)
```

### **Options**

- Exclusion of cofactors and transport reactions.
- Return of reactions involving human proteins or human metabolites only (as defined by ChEBI).
- Further filtering can be performed to report on reactions having more than one participant represented in the input query.

## Visualizing Returned RaMP-DB Reactions



## RaMP-DB – Many Uses and Accessibility

#### **User-friendly web app:**

https:rampdb.nih.gov

#### R package and Vignette/Tutorial:

https://github.com/ncats/RaMP-DB

https://ncats.github.io/RaMP-DB/RaMP Vignette.html

#### **Documented API Access:**

https://ramp-api-alpha.ncats.io/ docs /

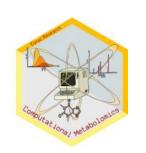
#### Code for building back-end and front-end:

https://github.com/ncats/RaMP-Client/ https://github.com/ncats/RaMP-Backend

#### Uses:

Reuse of parsing scripts, incorporate in data analysis workflows, develop new methods using RaMP-DB, etc.

Bioconductor Computational Metabol0mics ecosystem



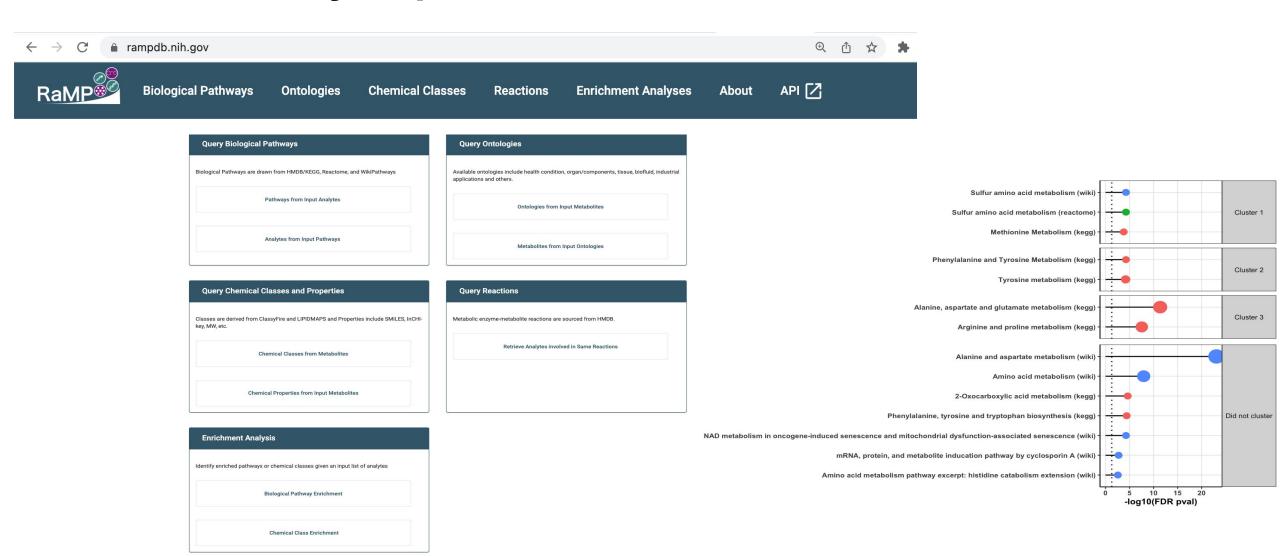








## We are always open to feedback!



### It's a team effort!









John Braisted

Tim Sheils

Andrew Patt Keith Kelleher

#### Also special thanks to

- Tara Eicher, Kyle Spencer, Haley Chatelaine for development/testing.
- Johannes Rainer for help in converting RaMP-DB to **SQLite**
- Sam Michael, Ke Wang, Betty Li, Amit Virakamath, and NCATS ITRB for computing infrastructure
- Egon Willighagen and WikiPathways team; Jeff ia and MetaboAnalyst team; David Wishart and HMDB team for feedback



https://ncats.nih.gov/preclinical/core/informatics (We're hiring!)

## Let's check it out and get started!



All workshop materials can be found here:



https://shorturl.at/kvzIZ

Some installation prerequisites for RaMP-DB:



https://shorturl.at/bgqrQ

Provide your comments/thoughts here:



https://shorturl.at/sS138