Metabolomics Workbench and the National Metabolomics Data Repository University of California San Diego and San Diego Supercomputer Center

Overview and Infrastructure

NIH Common Fund's National Metabolomics Data Repository (supported by NIH grant, U2C-DK119886)

Overview of the Metabolomics Workbench

The <u>National Institutes of Health (NIH) Common Fund Metabolomics Program</u> was developed with the goal of increasing national capacity in metabolomics by supporting the development of next generation technologies, promoting data/metadata sharing and collaboration and providing training and mentoring opportunities. In support of this effort, the Metabolomics Workbench website was created at the University of California, San Diego in 2013. The Metabolomics Workbench houses the National Metabolomics Data Repository (NMDR) which serves as a national and international center for metabolomics data and metadata and provides analysis tools and access to metabolite standards, protocols and other resources to the global community.



Metabolomics Workbench: <u>https://www.metabolomicsworkbench.org</u>

Contains the National Metabolomics Data Repository (NMDR)

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elcome to the UCSD Metabolo	mics Workbench, a resource sponsore	ed by the Common Fund of the Na	tional Institutes of Health.
National Metabolomics D	ata Repository	_	Quick Links - Key Resources 🗸 🗸
Upload and Manage Studies	Browse and Search Studies	Analyze Studies	Sollow @MetabolomicsWB
As of 02/14/22 a total of 2002 st	udies have been processed by the Nation	al Metabolomics Data Repository	Tweets by @MetabolomicsWB
(NMDR). There are 1/2/ public their embargo dates.	ly available studies and the remainder (2	(75) will be made available subject to	Metabolomics Workbench @MetabolomicsWB
Recently released studies on NMDR ST002058 - Muscle/Lung/Tumor metabolomics; <i>Mus musculus</i> ; <u>University of Colorado Anschutz Medical</u> <u>Campus</u>			The National Metabolomics Data Repository (NMDR) at @MetabolomicsWB has just processed its 2,000th study! MS/NMR data/metadata
ST002059 - 4T1 and SkM cells;	Homo sapiens; University of Colorado An	schutz Medical Campus	on studies covering over 130 species. Browse/search/download at
ST002067 - Time-Resolved Me (LC-MS); Mus musculus; Georgi	tabolomics of a Mouse Model of Ovaria a Institute of Technology	n High-Grade Serous Carcinoma	NIH Common Fund Stage 2 Metabolomics Consortium Centers
			Metabolomics Consortium Coordinating Center (M3C) 문 Bichard Yost U of Florida
Metabolite Structure Data	base		Metabolomics Workbench/NMDR 로 Shankar Subramaniam, UC San Diego (fbis website)
Updates to the Metaboli The updated Metabolite struc Metabolomics Workbench a chemical class. Over 164,000	te Structure Database (Februar ture database of primary and seconda Contains new substructure and text structures have been added including	ry 2, 2022) ary metabolites at the -based searches including by g over 10,000 sterols.	Compound Identification Cores (CIDCs) Arthur Edison, U. of Georgia Alexey Nesvizhskii, U. of Michigan Oliver Fiehn, UC Davis Dean Paul Jones, Emory University Thomas Metz, Pacific Northwest Nat. Lab.
			Data and Tools Cores (DTCs) John Weinstein, MD Anderson Cancer C. Jamey Young, Vanderbilt University Xiuxia Du, U. of North Carolina Charlotte Shuzhao Li, Emory University Alla Kamovsky, U. of Michigan Katerina Kechris, U. of Colorado, Denver Gary Patti, Washington U. at St. Louis
Holding and a set of se			Please cite:Metabolomics Workbench & You will get more info

Metabolomics Workbench website: what does it contain?

- National Metabolomics Data Repository (NMDR)
 - MS and NMR metabolomics studies
 - Metadata, targeted/untargeted measurements, raw data
- Metabolomics Workbench Metabolite database
- RefMet standardized metabolite nomenclature resource
- Online suite of statistical analysis tools
 For NMDR studies and ad-hoc user-supplied datasets
- MetStat summary reporting tool
- Human gene/protein database of metabolism-related genes
- Protocols for metabolomics experiments
- REST service
- MS search tools
- Other metabolomics software (MW group and collaborators)

Overview of NMDR cloud computing infrastructure

(located at the San Diego Supercomputer Center)



Public website

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	Summary of all studies						
	Study ID 술루	Study Title ✿基	Species ★↓	Institute ★寻	Analysis 술루	Submitted 會류	Download
	ST000001	Fatb Induction Experiment (FatBIE)	Arabidopsis thaliana	University of California, Davis	MS	2013-01-15	Raw data (476K)
	ST000002	Intestinal Samples II pre/post transplantation	Homo sapiens	University of California, Davis	MS	2013-01-23	Raw data (664K)
	ST000003	Metabolomic analysis of mouse embryonic fibroblasts, embryonic stem cells, and induced pluripotent stem cells	Mus musculus	University of California, Davis	MS	2013-01-16	Raw data (5.3G)
	ST000004	Lipidomics studies on NIDDK / NIST human plasma samples	Homo sapiens	LIPID MAPS	MS	2013-02-20	Raw data (48K)
	ST000005	Timecourse on RAW 264.7 cells treated with Kdo2-Lipid A and compactin	Mus musculus	LIPID MAPS	MS	2013-02-20	Raw data (56K)
	ST000006	White Wine Study	Vitis vinifera	University of California, Davis	MS	2013-02-21	Raw data (532K)
	ST000007	Rice Infection Study	Oryza sativa	University of California, Davis	MS	2013-02-22	Raw data (1.7M)
	ST000008	Metabolomics Analysis of Population Genetics (PopGen)	Homo sapiens	RTI International	NNR	2013-02-17	Raw data (39M)

NMDR online portals

Data submission/review portal

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Study ID	ary of all studies	Species 술루	Institute 全导	Analysis +	Submitted	Download
ST000001	Fatb Induction Experiment (FatBIE)	Arabidopsis thaliana	University of California, Davis	MS	2013-01-15	Raw data (476
ST000002	Intestinal Samples II pre/post transplantation	Homo sapiens	University of California, Davis	MS	2013-01-23	Raw data (664
ST000003	Metabolomic analysis of mouse embryonic fibroblasts, embryonic stem cells, and induced pluripotent stem cells	Mus musculus	University of California, Davis	MS	2013-01-16	Raw data (5.3)
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Development websites

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Open access

(non-embargoed studies)

Access controlled

Submit/View data/metadata submitted by your institution (or other institutions for which you have permission)

Metabolomics study workflow



Data analysis

Overview of NMDR Inputs and Outputs





NMDR Online submission overview



The **mwTab** format: A "common currency" for metadata/data sharing and storage

SECTIONS:	#METABOLOMICS WORKBENCH Nagireddy_Put VERSION 1	luri_20211115_090410 DATATRACK_ID:2929 STUDY_ID:ST002005 ANALYSIS_ID:AN003268 PROJECT_ID:PR001271
Matadata	CREATED_ON November 29, #PROJECT	2021, 7:26 pm
<u>ivietadata</u>	PR:PROJECT_TITLE PR:PROJECT_TITLE	Alterations of lipids in tumor tissues from African American and European American patient with bladder cancer
Project	PR:PROJECT_SUMMARY PR:PROJECT_SUMMARY PR:PROJECT_SUMMARY PR:PROJECT_SUMMARY	Cancer affects all individuals in the United States, unfortunately due to socioeconomics, and environmental disadvantages, certain group of populations especially African American (AA) community bear a high burden of cancer than the other community. Based or different again a condensional study expected that
Study	PR: PROJECT_SUMMARY PR: PROJECT_SUMMARY PR: PROJECT_SUMMARY	higher incidence and mortality rate of bladder cancer in AA community. To understand and reveal the biological mechanism in terms of lipidomics, lipidomics profile were performed in 98 bladder cancer (African American and
Experimental variables (factors)	PR:PROJECT_SUMMARY PR:INSTITUTE PR:LAST_NAME DD:FLDST_NAME	European America) tissues including benign. Baylor College of Medicine Putluri Nagi peddy
Subject	PR:ADDRESS PR:EMAIL PR:PHONE	One Baylor Plaza, Houston, Texas 77030 putluri@bcm.edu (713) 798-3139
Collection	#STUDY ST:STUDY_TITLE ST:STUDY_TITLE ST.STUDY_CLAMMARY	Alterations of lipids in tumor tissues from African American and European American patient with bladder cancer
Treatment	ST:STUDY_SUMMARY ST:STUDY_SUMMARY ST:STUDY_SUMMARY ST:STUDY_SUMMARY	socioeconomics, and environmental disadvantages, certain group of populations especially African American (AA) community bear a high burden of cancer than the other communities. Based on different social epidemiological study reported that
Sample preparation	ST:STUDY_SUMMARY ST:STUDY_SUMMARY ST:STUDY_SUMMARY	higher incidence and mortality rate of bladder cancer in AA community. To understand and reveal the biological mechanism in terms of lipidomics, lipidomics profile were performed in 98 bladder cancer (African American and
Chromatography	ST:STUDY_SUMMARY ST:INSTITUTE ST:DEPARTMENT ST:LAST NAME	European America) tissues including benign. Baylor College of Medicine Molecular and Cellular Biology Putluri
Analysis	ST:FIRST_NAME ST:ADDRESS ST:EMAIL	Nagireddy One Baylor Plaza, Houston, Texas 77030 putluri@bcm.edu
MS	ST: PHONE #SUBJECT SU: SUBJECT_TYPE SU: SUBJECT_SDECIES	(713) 798-3139 Human
NMR	SU: JOBOBCI_SECTES SU: TAXONOMY_ID #FACTORS #SUBJECT SAMPLE FACTORS:	NOME SAFENS 9606 SUBJECT (optional) [tab]SAMPLE[tab]FACTORS (NAME:VALUE pairs separated by]) [tab]Faw file names and
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Data	SUBJECT_SAMPLE_FACTORS SUBJECT_SAMPLE_FACTORS	- 03-22-2021-Pos-35599 BENIGN-40 Group:Benign - 03-22-2021-Pos-35774 BENIGN-42 Group:Benign

Data

Named metabolite measurements (table) Named metabolites and annotations File names for untargeted datasets

Additional

Comments preceded by a #

MwTab specification: https://www.metabolomicsworkbench.org/data/mwTab_specification.pdf