The NIH Common Fund Metabolomics Consortium

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Consortium Goals

1. *Establish an enduring public repository* for metabolomic data
2. *Overcome technical hurdles* in analyzing and interpreting metabolomics data, including the ability to determine metabolite identities
3. *Develop consensus* for, and promote adoption of, best practices and guidelines to promote accuracy, reproducibility, and re-analysis of metabolomics data in collaboration with the national and international communities.

Scan for additional information on consortium cores

Consortium Cores

National Metabolomics Data Repository, Metabolomics Workbench http://metabolomicsworkbench.org
Shankar Subramanian, University of California San Diego, National Metabolomics Data Repository NextGen Metabolomics Workbench
Five Compound Identification Cores
Arthur S. Edison, University of Georgia, Genetics and Quantum Chemistry as Tools for Unknown Metabolite Identification
Alexey Nesvizhskii, Charles R. Evans, University of Michigan, Michigan Compound Identification Development Cores (MCIDC)
Oliver Fiehn, University of California Davis, West Coast Metabolomics Center for Compound Identification
Dean Paul Jones, Shuzhao Li, Gary W. Miller, Edward T. Morgan, Emory University, Mega-scale Identification Tools for Xenobiotic Metabolism
Thomas O. Metz, Pacific Northwest National Laboratory, Pacific Northwest Laboratory Compound Identification Core
Seven Data and Tools Cores
John Weinstein, Rehan Akbani, Bradley M. Bloom, University of Texas MD Anderson Cancer Center, Computational Tools for Analysis and Visualization of Quality Control Issues in Metabolomic Data
Jamey Young, Doug Allen, Vanderbilt University, Tools for Leveraging High-resolution MS Detection of Stable Isotope Enrichments to Upgrade the Information Content of Metabolomics Datasets
XiuXia Du, University of North Carolina Charlotte, Cross-platform and Graphics Software Tool for Adaptive LC/MS and GS Metabolomics Data Processing
Shuzhao Li, Emory University, Gary Siuzdak, Scripps Research Institute, Mummichog 3, Aligning Mass Spectrometry Data to Biological Networks
Alla Karnovsky, University of Michigan, George Michailidis, University of Florida, Methods and Tools for Integrative Functional Enrichment Analysis of Metabolomics Data
Katerina Kechris, Debashis Ghosh, University of Colorado Denver, Addressing Sparsity in Metabolomics Data Analysis
Garry Patti, Washington University at St. Louis, A Comprehensive Platform for High-Throughput Profiling of the Human Reference Metabolome
Consortium Coordinating Center http://metabolomics.info
Richard A. Yost, Michael Conlon, University of Florida, Metabolomics Consortium Coordinating Center

Participating in the Consortium

❑ Consortium workgroups are open to all. Workgroups identify hurdles, develop best practices, contribute to repository development, and build consensus. Current workgroups include:
- Software standards for Metabolomics Tool Development, Jamey Young, Vanderbilt University, chair.
- Quantum Mechanical Computing Best Practices, Art Edison, University of Georgia, chair
- Unknown lipids data exchange, Charles Evans, University of Michigan, chair

❑ The Metabolomics Workbench is open to all. Find, use, and contribute FAIR metabolomics data. See https://www.metabolomicsworkbench.org/data/

❑ For more info, or to participate in a workgroup, visit http://metabolomics.info, contact us at *info@metabolomics.info*, or follow us on Twitter @metabinfo

The NIH Common Fund (http://commonfund.nih.gov) addresses emerging scientific opportunities and pressing challenges in biomedical research that no single NIH Institute or Center can address on its own, but are of high priority for the NIH as a whole.