- ➤ The behavioral changes that accompany drug addiction are believed to result from both short and long term adaptive changes in the neurochemistry of brain reward centers. To date, molecular studies have elucidated some of the transcriptional changes that occur in the addicted brain. However, little if anything is known about the effects of drugs of abuse on the neuronal proteome.
- ➤ The major goal of the Yale/NIDA Neuroproteomics Center is to bring together strong programs in proteomics, in drug abuse research, and in various aspects of neurobiology, to characterize using proteomic methods basic aspects of brain function and to use these methods and knowledge to identify adaptive changes in protein expression and regulation that occur in response to exposure to drugs of abuse.

- ➤ The Center will improve existing technologies as well as develop new proteomics technologies that can be applied to neurobiological questions, particularly relevant to the actions of drugs of abuse.
- > The Center will provide training in proteomics technologies.
- ➤ The Center will encourage collaboration between Center Investigators who might otherwise not interact, and to try to bring investigators with no prior experience in drug abuse research into the field of drug abuse.

NIDA Center Investigators

Yale - CNNR, Physiology, Child Study Yale - Psychiatry

Rockefeller, Stanford, Chicago, UT Southwestern

Pietro DeCamilli

Stephen Strittmatter

Thomas Biederer

Sreeganga Chandra

Susumo Tomita

Paul Lombroso

Ralph DeLeone

Angus Nairn

Marina Picciotto

Sam Sathyanesan

Arthur Simen

Rajita Sinha

Jane Taylor

Zoron Zimolo

Paul Greengard

Robert Malenka

William Green

James Bibb

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Susumo Tomita

Arthur Simen

Elizabeth Eipper

Paul Lombroso

Rajita Sinha

Maria Morabito

Len Kaczmarek

Dan Wu Jane Taylor

Richard Lifton_

Zoron Zimolo



Pre- and Post-Synaptic Signaling Transcriptional Regulation

Neuronal Development

Neuronal Plasticity

Regulation of the Cytoskeleton

Endocytosis and Exocytosis



1. Administration CD: Kenneth Williams

Angus Nairn

2. Protein Profiling, ID, and Lipid analysis

CD: Kathy Stone

3. Post-translational (PTM) profiling CD: Erol Gulcicek 4. Targeted Proteomics MRM, Antibody Microarray

CD: Chris Colangelo

5. Protein Database, Biostatistics, Bioinformatics, HPComputing

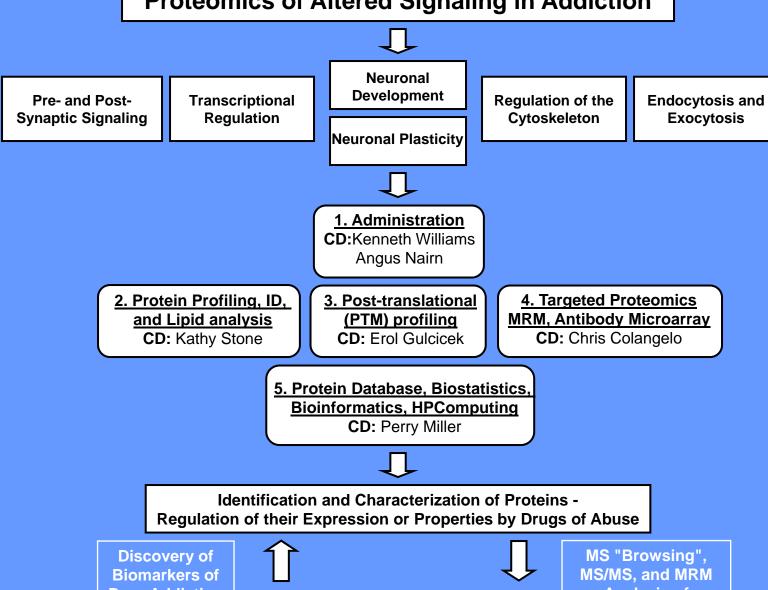
CD: Perry Miller



Identification and Characterization of Proteins - Regulation of their Expression or Properties by Drugs of Abuse



Clinical Studies of Drug Addiction



Drug Addiction

Analysis of Targeted Proteins

Clinical Studies of Drug Addiction



Neuronal Development

Thomas Biederer (2,5) Sreeganga Chandra (2,5) Dan Wu (2,5) Richard Lifton (2,3,4,5)

Regulation of the Cytoskeleton

Angus Nairn (2,3,4,5) Elizabeth Eipper (2,3,4,5) Pietro DeCamilli (2,3,4,5) Stephen Strittmatter (2,5)

<u>Pre- and Post-Synaptic</u> Signaling

Paul Greengard (2,3,4)
Angus Nairn (2,3,4,5)
Paul Lombroso (2,3,5)
Ralph DeLeone (2,3,5)
William Green (3,5)
Len Kaczmarek (3,5)
Pietro DeCamilli (2,3,4,5)
James Bibb (2,3,5)
Maria Morabito (2,5)

Neuronal Plasticity

Robert Malenka (3,5)
Susumo Tomita (3,5)
Elizabeth Eipper (2,3,4,5)
James Bibb (2,3,5)
Jane Taylor (2,3,4,5)
Marina Picciotto (2,3,4,5)
Angus Nairn (2,3,4,5)

Transcriptional Regulation

Arthur Simen (2,3,5) Angus Nairn (2,3,4,5) Sam Sathyanesan (2,3,5)

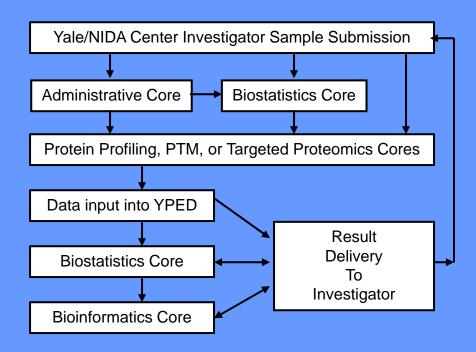
Endocytosis and Exocytosis

Pietro DeCamilli (2,3,4,5) Susuma Tomita (3,5)

Clinical Studies of

Drug Addiction
Rajita Sinha (2,4,5)
Zoron Zimolo (2,4,5)

General procedure for initiation of projects and analysis of samples



Collaborations between Center Investigators

