



MERCK - MIT

Computational & Systems Biology Symposium

Original image courtesy of Thomas Rush, III, Ph.D., Drug Design & Optimization Merck Research Laboratories - Boston

Merck Research Laboratories Boston 2010 SEMINAR SERIES

Featured Speakers:

Merck

Hongyue Dai

"Identifying HCC siRNA targets by multiple components of molecular analysis"

Patrick Loerch

"A Network-based Approach to Integrating Diverse -omics Datasets and Identifying Disease Pathways"

MIT

Kristen Naegle

"A combinatorial unsupervised learning framework for prediction of post-translational modification functions in signaling networks"

Bracken King

"Analysis of biological systems with information theory"

Additional speakers from MIT's Department of Biological Engineering and the Department of Biology will also be participating.

Monday, April 5, 2010

9:00am: Breakfast

10:00am-5:00pm: Talks and Posters

Advance registration required at www.merckboston.com

Merck Research Laboratories Boston, Auditorium
33 Avenue Louis Pasteur, Boston (On-site parking is not available)
For directions, please visit www.merckboston.com