

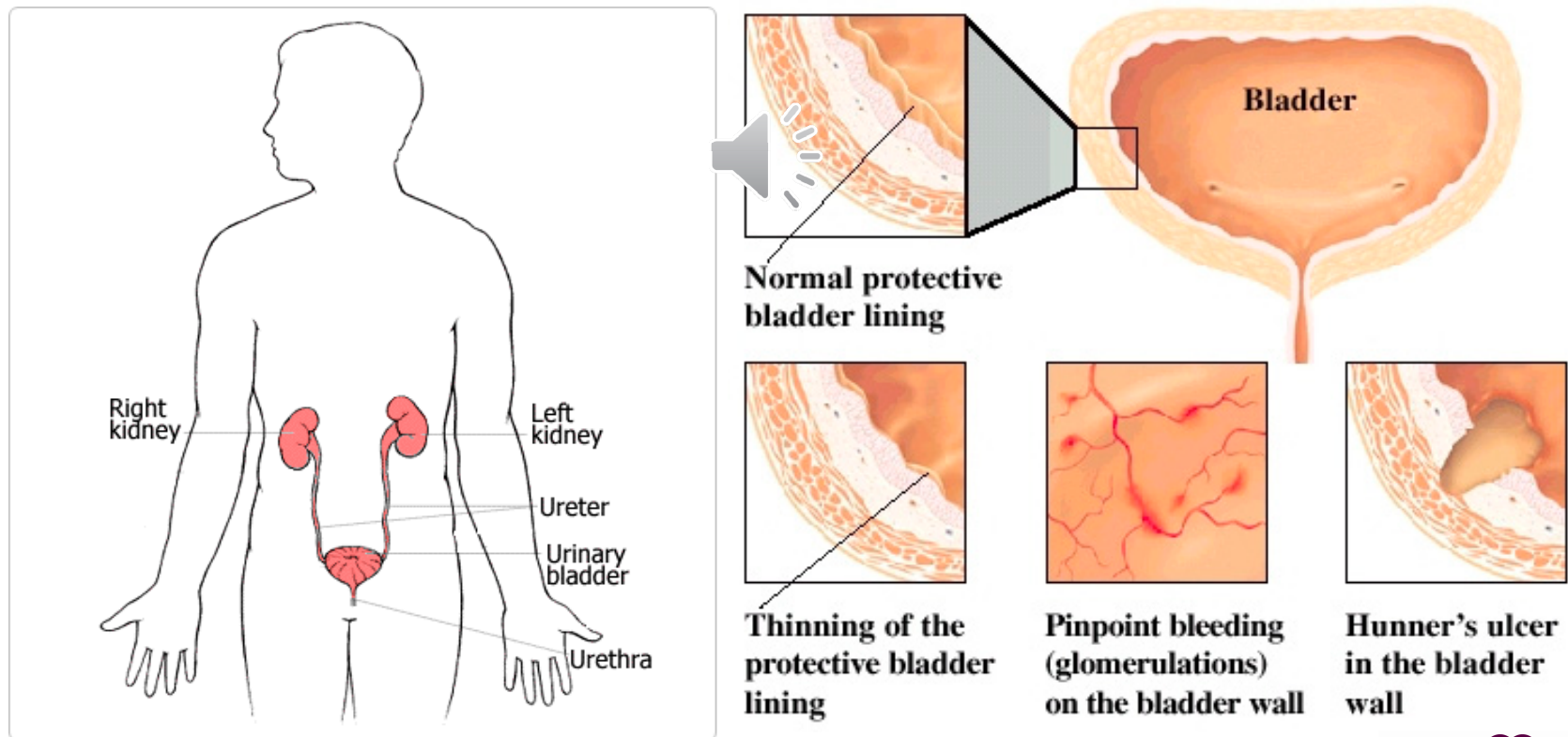
Interstitial Cystitis-Associated Urinary Metabolites Identified by Mass-Spectrometry Based Metabolomics Analysis

Jay Kim, PhD
Cedars-Sinai Medical Center



**Confidential information included*

Urinary Metabolite Profiling Combined with Computational Analysis Suggest Interstitial Cystitis-Associated Candidate Biomarkers



Interstitial Cystitis

- **A chronic syndrome of unknown etiology**
- **Very common bladder disease among old generation (more than one out of 77 people in USA)**
- **Affects quality of life, productivity and work performance—Public health burden**
- **Elmiron, the first FDA-approved oral drug for IC, shows unfavorable side effects**
- **Need for new medication for IC**
- **Need for objective and clinically relevant indicators**



IC-Associated Mechanistic Signaling Network 1:

The Frizzled 8-Associated Antiproliferative Factor Enhances p53 Stability Through USP2a and MDM2

FEBS Letters 581 (2007) 3795–3799

p53 mediates interstitial cystitis antiproliferative factor (APF)-induced growth inhibition of human urothelial cells

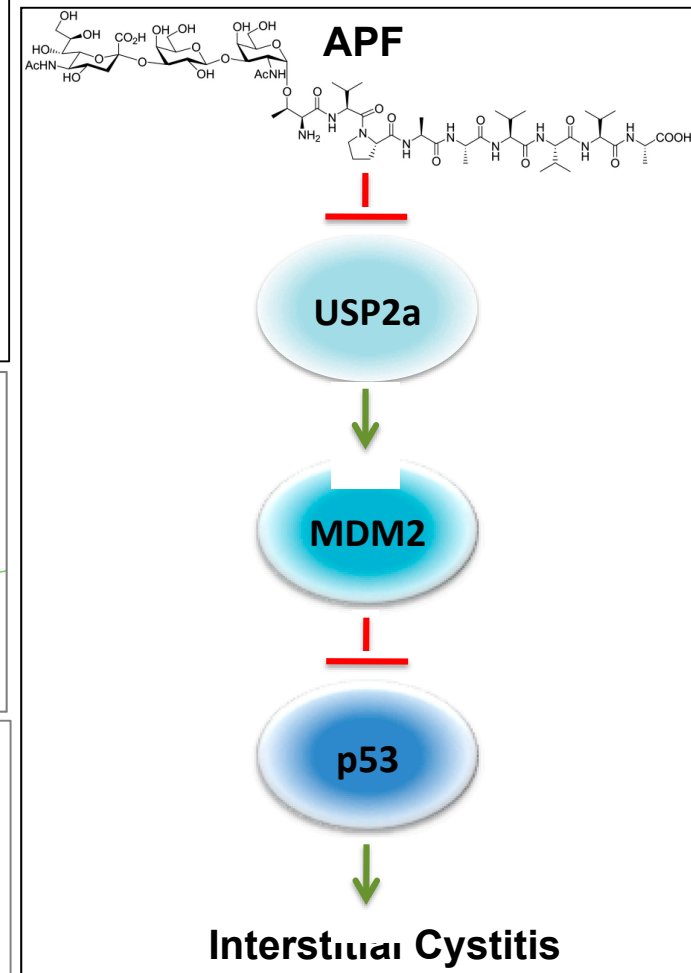
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Received 6 February 2007; accepted 21 June 2007



Heparin-binding epidermal growth factor-like growth factor functionally antagonizes interstitial cystitis antiproliferative factor via mitogen-activated protein kinase pathway activation

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Accepted for publication 10 July 2008

OPEN ACCESS Freely available online

PLOS ONE

A Synthetic Form of Frizzled 8-Associated Antiproliferative Factor Enhances p53 Stability through USP2a and MDM2

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IC-Associated Mechanistic Signaling Network 2:

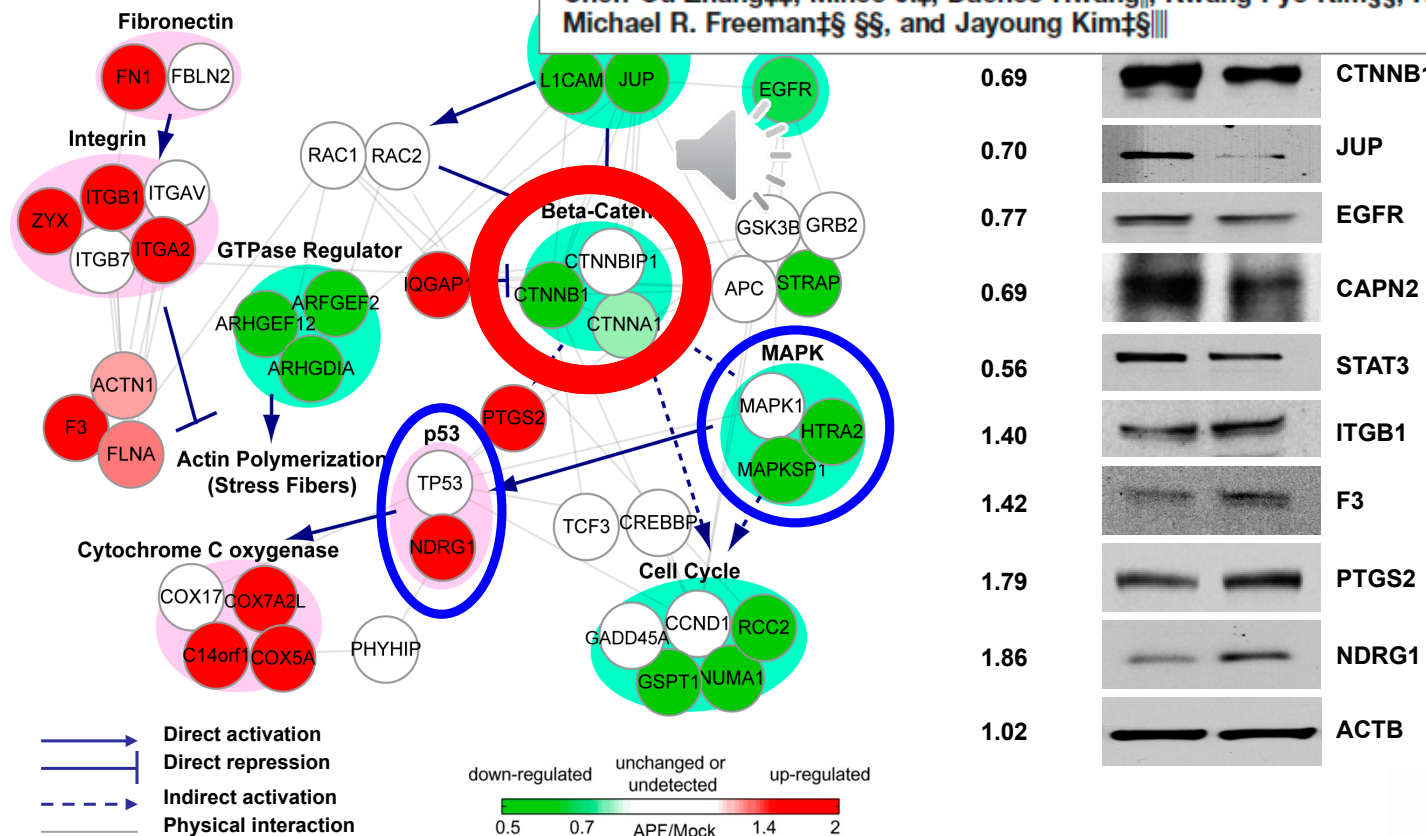
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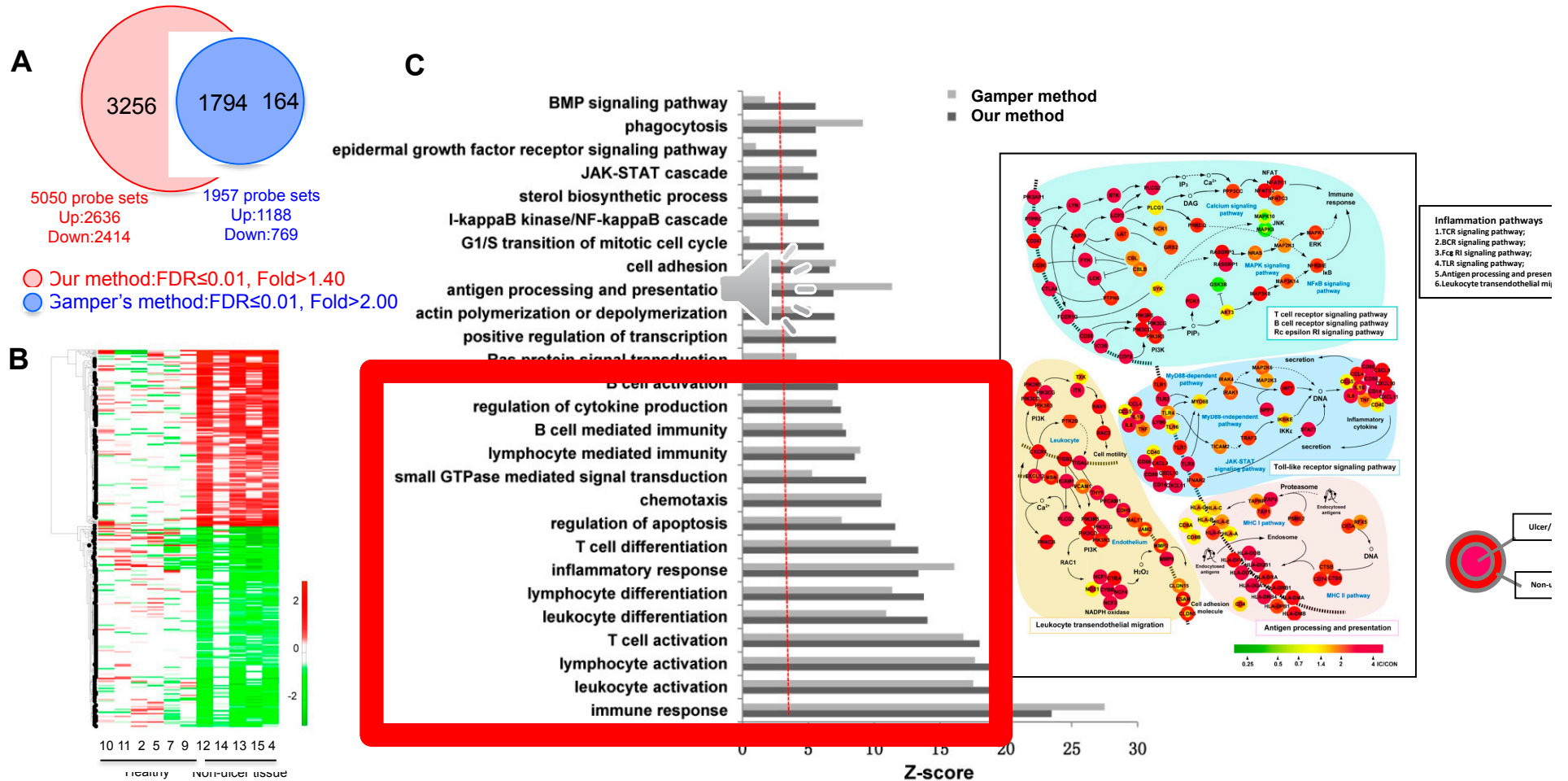
Quantitative Proteomics Identifies a β -Catenin Network as an Element of the Signaling Response to Frizzled-8 Protein-Related Antiproliferative Factor*

Wei Yang†§¶, Yeun Goo Chung‡, Yongsoo Kim||, Taek-Kyun Kim||, Susan K. Keay**, Chen-Ou Zhang‡‡, Mihee Ji‡, Daehee Hwang||, Kwang Pyo Kim§§, Hanno Steen¶¶¶, Michael R. Freeman‡§ §§, and Jayoung Kim‡§||



IC-Associated Mechanistic Signaling Network 3:

Integration Analysis of Quantitative Proteomics and Transcriptomics Data Identifies Potential Targets of Frizzled-8 Protein-related Antiproliferative Factor *In Vivo*



'OMICS' Approaches to Understand Interstitial Cystitis

More 'OMICS' Profiles using the **Cutting-Edge Technology** are needed

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Integration analysis of quantitative proteomics and transcriptomics data identifies potential targets of frizzled-8 protein-related antiproliferative factor *in vivo*

Wei Yang^{1,2,3}, Yongsoo Kim⁴, Taek-Kyun Kim⁴, Susan K. Keay⁵, Kwang Pyo Kim⁶, Hanno Steen^{3,7}, Michael R. Freeman^{1,2,6,8}, Daehee Hwang⁴ and Jayoung Kim^{1,2,8}



Review Article

Int Neurourol J 2012;16:159-168

<http://dx.doi.org/10.5213/inj.2012.16.4.159>

pISSN 2093-4777 · eISSN 2093-6931

International Neurology Journal

INJ

'Omics' Approaches to Understanding Interstitial Cystitis/Painful Bladder Syndrome/Bladder Pain Syndrome

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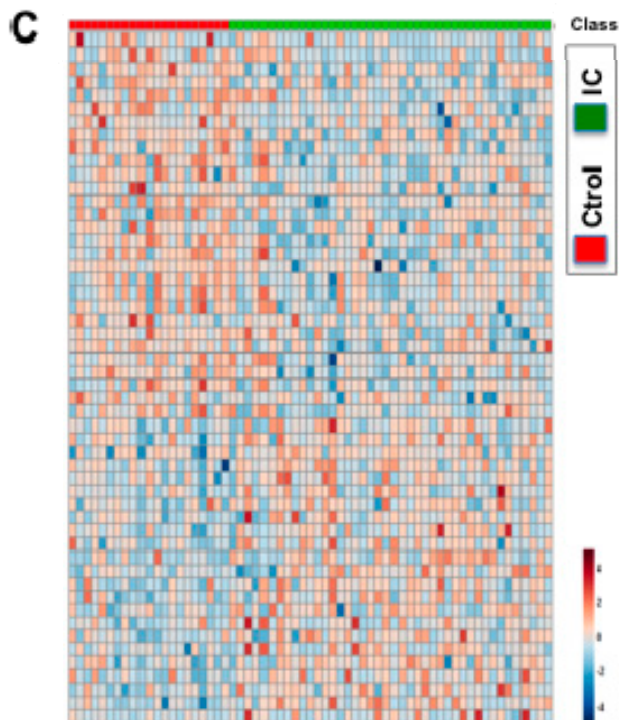
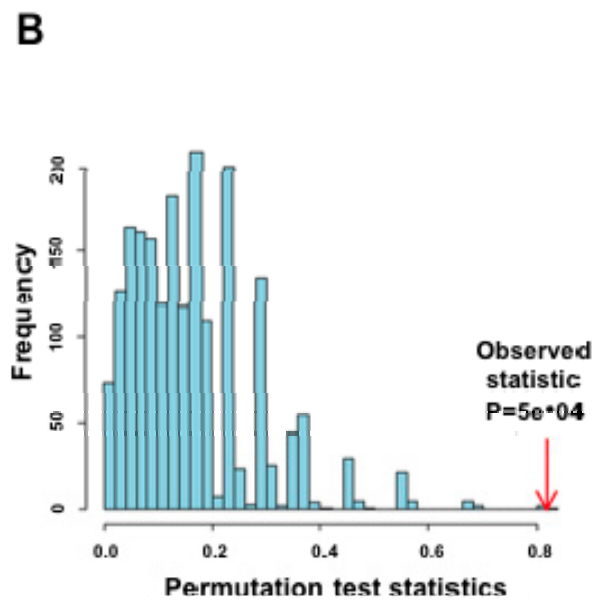
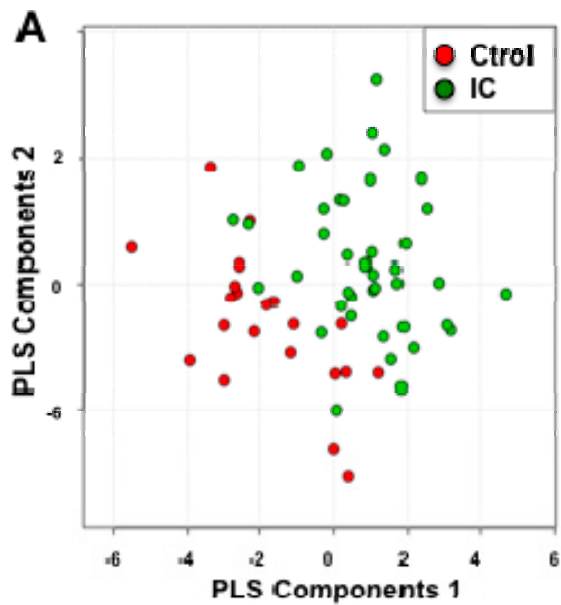
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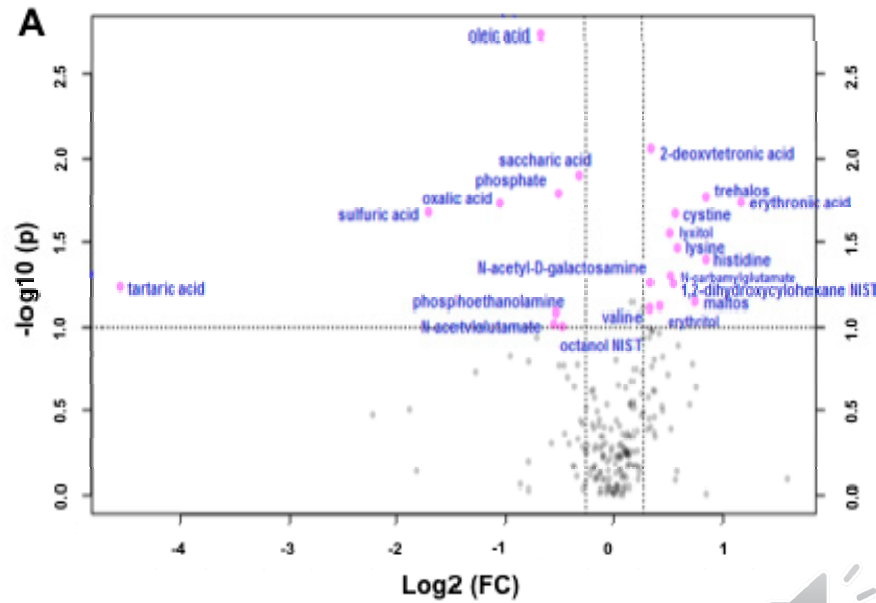
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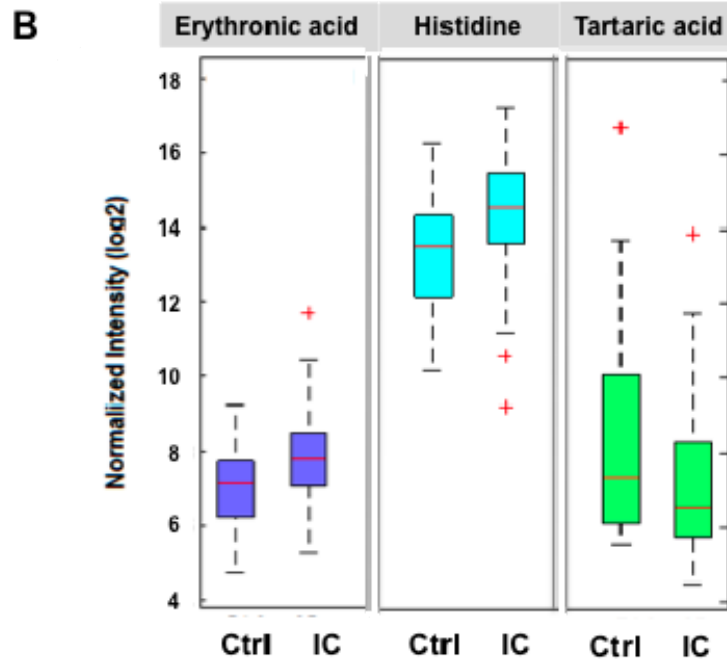
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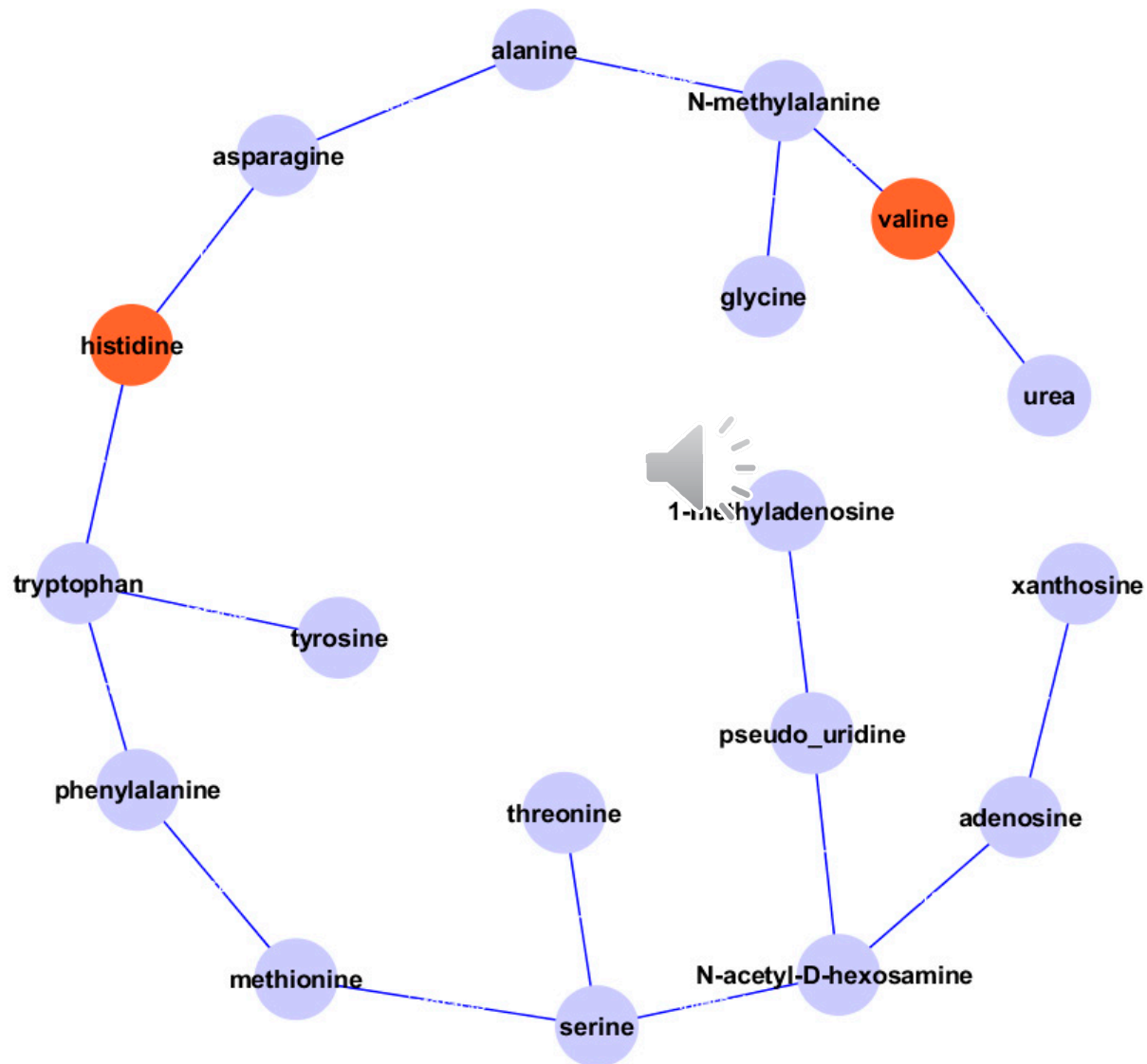
Differentiation of IC patients and healthy control groups using multivariate analysis



A volcano plot showing differentially expressed metabolites in IC patients.

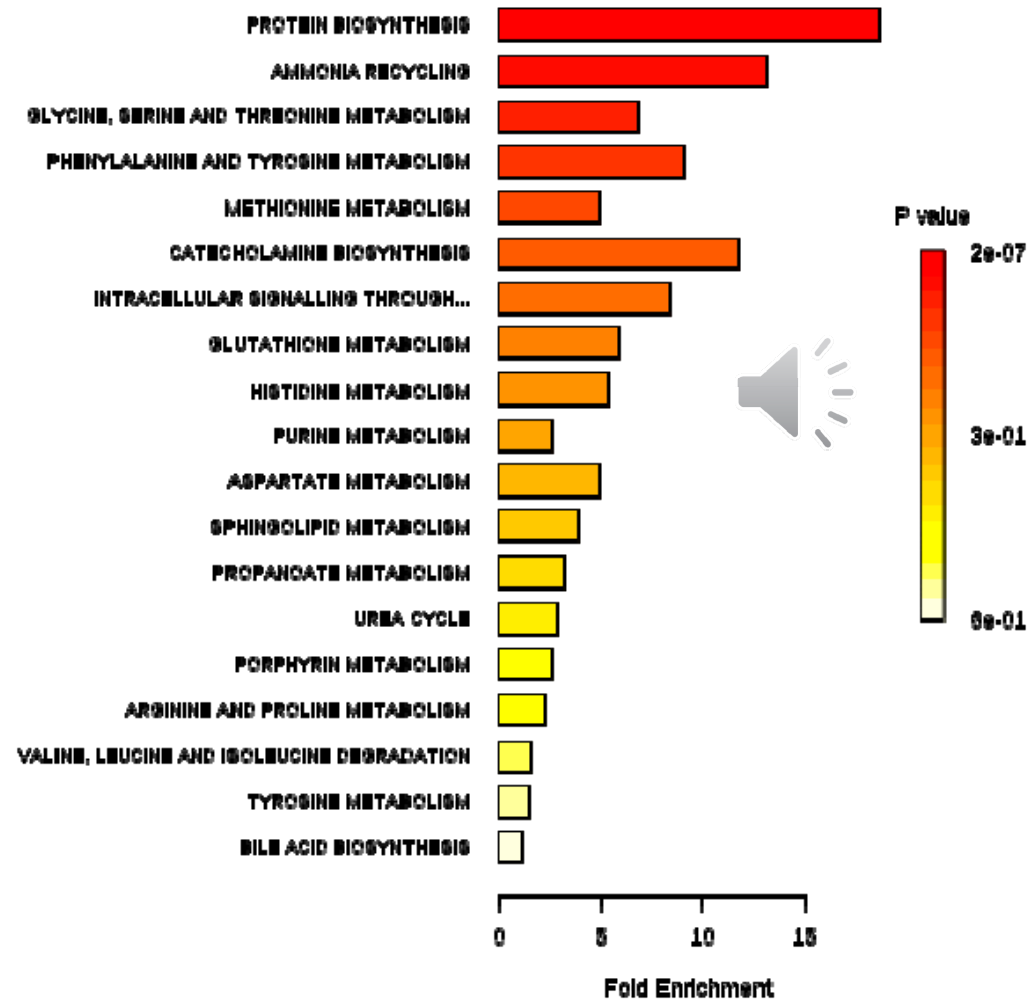


Network modeling derived from IC-associated metabolites




Differential network in IC is identified with multilevel local graphical model

Metabolite Sets Enrichment Overview



Acknowledgements

- NIDDK/NIH 1R01DK100974
- NIDDK/NIH 1U01 DK103260
- Department of Defense (W81XWH-15-1-0415)
- Centers for Disease Controls and Prevention (1U01DP006079)
- Steven Spielberg Discovery Fund Research Career Development Award
- U24 DK097154
- UCLA CTSI UL1TR000124
- Interstitial Cystitis Association (ICA,  grant)
- Fishbein Family IC Research Foundation
- New York Academy of Medicine
- Children's Hospital Boston Faculty Development
- J.K. is an IMAGINE NO IC Scholar, American Urological Association Foundation Research Scholar and an Eleanor and Miles Shore Scholar of Harvard Medical School.