## Introduction to Ingenuity Pathway Analysis

## **Hosted by the Cushing/Whitney Medical Library**

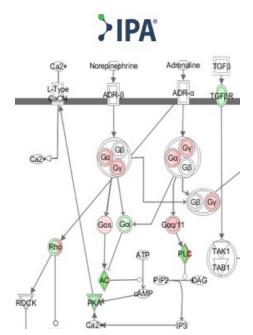
**Date/time**: Wednesday, July 15, 2015 from 9 AM - 12.00 PM **Location**: C-103 – 333 Cedar St, New Haven, CT 06520

**Presenter**: Dr. Kate Wendelsdorf, Applied Advanced Genomics, QIAGEN Informatics

Registration required due to limited seating: <a href="http://schedule.yale.edu/event.php?id=985558">http://schedule.yale.edu/event.php?id=985558</a>

What is IPA and what questions can it address? Overview of key features in IPA Ingenuity Knowledge Base

Search & Pathway Building - Gene/ Chemical, Functions, Drug Targets Advanced Search: Limiting results to a molecule type, family or disease-association.



Building pathways: Creating a pathway, pathway navigating, Using Build and Overlay tools Bioprofiler

Dataset Analysis: Interpretation of Gene, Transcript, Protein and Metabolite Data

Data Upload and Analysis: Uploading and formatting a dataset, setting analysis parameters and running an analysis

Pathway Analysis and Canonical Pathways

Downstream Effects Analysis and identifying downstream functions and processes that are likely affected

**Upstream regulators Analysis** 

Causal Network Analysis and identifying likely root regulators

Regulator Effects Analysis to link upstream regulators with downstream functions and

processes that are affected

Comparison analysis and comparing multiple observations

Questions: Rolando Milian – <u>rolando.milian@yale.edu</u>

