

# 10<sup>th</sup> Annual Course on Isotope Tracers in Metabolic Research: Principles and Practice of Kinetic Analysis – Oct 30th – Nov 3rd

Sponsored by NIDDK and organized on behalf of the NIH-sponsored  
Mouse Metabolic Phenotyping Centers (MMPC)

**October 30 – November 3rd, 2017**

Homewood Suites by Hilton® Nashville Vanderbilt \* Course Location  
2400 West End Avenue - Nashville, Tennessee 37203 (615-340-8000)

The tenth annual course provides basic introductory and comprehensive information on performing metabolic studies using tracers labeled with radioactive or stable isotopes, in humans and in animals. The course is designed for beginners as well as those with experience who wish to expand their capabilities to more sophisticated problems. The faculty is well-versed in a variety of applications and methodologies. Techniques will be presented for investigating whole body metabolism, for metabolite balance across organs, intracellular flux rates and pathway regulation. The basic aspects of modeling will be considered, as well as specific applications to the study of carbohydrate, fat, protein metabolism and energy balance. Theoretical and practical matters related to sample analysis by mass spectrometry and NMR will be discussed, including detailed numerical examples of calculations involved in determining isotopic enrichment and basic kinetic parameters. Advanced lectures will discuss in more detail the use of positional and mass isotopomer analysis for intracellular flux rates and various aspects of protein and amino acid metabolism. Course material will be available for download from <http://www.mmpc.org/shared/tracers.aspx> In addition to organized sessions, individual attendees will have ample opportunities for personal interaction with faculty members in the form of one-on-one mentoring sessions to discuss their research projects in more depth.

## Faculty (\*Co Directors)

\*Henri Brunengraber; Case Western Reserve U.  
Gary Cline, Yale U.  
Melanie Cree Green, U. of Colorado  
Joanne Kelleher, Mass. Inst. Tech.  
Maren Laughlin, National Inst. Health  
\*Owen McGuinness, Vanderbilt U.

Matthew Merritt, U. of Florida  
Elizabeth Parks, U. of Missouri  
Stephen Previs, Merck  
Michelle Puchowicz, Case Western Reserve U.  
Jamey Young, Vanderbilt U.  
\*Robert R. Wolfe, U. Arkansas

## Program Outline

### Monday

Basic characteristics of radioactive, stable isotope tracers.  
General principles of mass spectrometry.  
Isotopic enrichment using GC-MS.  
Methods of mass spectrometry analysis.  
Measurement of specific activity.

### Tuesday

Tracer kinetics (single pool models).  
Oxidation and synthesis rates.  
Glucose metabolism (clamp studies).  
Lipid metabolism (basic kinetics).

### Wednesday

Pathway fluxes using NMR isotopomer analysis.  
Methods in protein metabolism.

### Thursday

Energy expenditure with doubly labeled water.  
Synthesis rates with deuterated water: proteins, fatty acids  
sterols, glucose, nucleic acids.  
Mass isotopomer distribution analysis: polymer synthesis,  
multiple flux pathways, TCA cycle, anaplerosis.

### Friday

Pathway discovery via association of isotopomer analysis and  
metabolomics.  
Inherently difficult problems.

For additional information on this course please contact: [Amanda Zetans \(isotope.tracer@vanderbilt.edu\)](mailto:isotope.tracer@vanderbilt.edu) (615-343-1065)