May 16-18, 2019, Niagara Falls, NY

3-DAY WORKSHOP ON MONOCLONAL ANTIBODY

PHARMACOKINETICS & PHARMACODYNAMICS Concepts, Applications, & Case Studies

University at Buffalo
School of Pharmacy and
Pharmaceutical Sciences

COURSE OUTLINE

BMSR Biomedical Simulations Resource

This workshop has been designed to provide a detailed discussion of issues relevant to the pharmacokinetic / pharmacodynamic (PK/PD) modeling of antibody drugs, and to provide a series of "hands-on" case studies describing the development and application of mathematical models to simulate and characterize antibody PK/PD. Lectures will address primary determinants of antibody pharmacokinetics (PK) and pharmacodynamics (PD), the design and implementation of pre-clinical investigations of antibody PK/PD, and state-of-the-art mathematical models to characterize and predict antibody PK and PD. Case studies, featuring use of the ADAPT program, will include application of models for antibody ligand binding at



use of the ADAPT program, will include application of models for antibody-ligand binding and disposition, target-mediated disposition, and physiologically-based pharmacokinetic modeling. Special emphasis is placed on discussion of the role of FcRn on the absorption, distribution, and elimination of antibodies, on the mathematical modeling of target-mediated antibody disposition, and on physiologically-based modeling of antibody pharmacokinetics. The workshop content is provided as a combination of formal lectures and informal review sessions.

Subjects that will be presented include:

Determinants of antibody pharmacokinetics and pharmacodynamics: mechanisms of antibody elimination, the role of convection in the kinetics of antibody distribution, the role of FcRn in antibody absorption, distribution, and elimination

Interspecies Scaling of Antibody PK: considerations and examples involving the use of allometric methods to scale protein pharmacokinetics from pre-clinical models to humans

Drug-drug interactions: mechanistic considerations and examples for drug-drug interactions involving monoclonal antibodies, including consideration of antibodies as perpetrators and as victims of DDI

Target-Mediated Antibody Disposition: modeling, implications for interspecies scaling, implications for First-in-Human studies **Modeling of bimolecular antibody-ligand interaction**

Physiologically-based pharmacokinetic modeling: Incorporation of FcRn-mediated antibody transport in PBPK models, incorporation of target-mediated disposition, use of PBPK and preclinical data to predict antibody disposition in humans

COURSE DIRECTION

Joseph P. Balthasar, PhD. Dr. Balthasar is Professor of Pharmaceutical Sciences at the University at Buffalo, State University of New York and Director of the Center for Protein Therapeutics. His PK/PD modeling interests and capabilities include the development and preclinical evaluation of anti-cancer immunotherapies (including immunoconjugate immunotherapies), and the development and preclinical evaluation of novel immunotherapies for humoral autoimmune conditions.



David Z. D'Argenio, PhD is Professor of Biomedical Engineering at the University of Southern California and holder of the Chonette Chair of Biomedical Technology. He is a Fellow of the American Institute for Engineering in Medicine and Biology, American Association of Pharmaceutical Sciences, International Society of Pharmaceutics, and a past member of the FDA Advisory Committee for Pharmaceutical Science and Clinical Pharmacology. Since 1985 he has served as co-director of the Biomedical and Simulations Resource (BMSR) at USC, which develops, applies and disseminates advanced modeling methods for studying biological systems, where he has also led the development of the ADAPT software for PK/PD modeling and analysis.



Donald E. Mager, PharmD, PhD is Professor of Pharmaceutical Sciences at the University at Buffalo, State University of New York. He was also a Visiting Professor at the University Paris Descartes (Jan. 2007-2013). He currently serves on the Pharmaceutical Sciences and Clinical Pharmacology Advisory Committee to the FDA, and as an Associate or Consulting Editor at CPT:Pharmacometrics & Systems Pharmacology and Pharmacology, Research & Perspectives. His research involves identifying molecular and physiological factors that control the pharmacological properties of drugs, with a focus on anti-cancer and immunomodulatory agents.



Dhaval K. Shah, PhD is Assistant Professor of Pharmaceutical Sciences at the University at Buffalo, State University of New York. His research focuses on understanding the determinants for the ADME of protein therapeutics. He is involved in the development of a platform PBPK model for biologics that can characterize and predict the pharmacokinetics of diverse protein therapeutics in several preclinical species and human. Dr. Shah also directs the discovery, development and clinical translation of novel protein therapeutics like antibody-drug conjugates and bi-specific molecules in his laboratory. His research is supported by NIH & pharmaceutical industry.



AGENDA

Day 1		Day 2	
08:45-09:00	Introductions	09:00-10:00	Mathematical Modeling of Bimolecular Antibody-Antigen
09:00-11:00	Introduction to Antibody Pharmacokinetics (Balthasar)		Interaction and ADC PKPD (Shah)
	 Introduction to antibodies (isotypes, polyclonal vs. 		• Review of binding kinetics (Law of Mass Action, equilibrium
	monoclonal, humanization, etc.)		vs. non-equilibrium binding)
	 Mechanistic determinants of antibody absorption, 		Mathematical modeling of antibody binding: Examples from
	distribution, and elimination (contrasting with determinants		antibodies used for immunotoxicotherapy
	of small-molecule ADME)		Introduction to mechanistic modeling of ADC PKPD
	• Comments on the mathematical modeling of antibody PK		• introduction to mechanistic modernig of ADC FRFD
	• Recent research relating to the role of FcR and mAb PK	10:00-11:00	Dhysiologically, Rosed DV Modeling of Moh (Bolthoson)
11:00-11:15	Break	10.00-11.00	Physiologically-Based PK Modeling of Mab (Balthasar)
11:15-11:40	Analytical Assays for Antibodies: Implications for		Review of PBPK models Application of PBPK models applied to Make
11.13-11.40	PK/PD Analyses (Balthasar)		Application of PBPK models applied to Mab
	Discussion of major types of analytical assays for		Discussion of major features of PBPK models for Mab &
	monoclonal antibodies (ELISA, RIA, LC MS/MS, SPR,		discussion of associated physiology (convection, restriction
	"direct" labeling)		coefficients, sites of catabolism, "two-pore formalism",
	E,		incorporation of specific binding, incorporation of FcRn)
11.40 12.00	• Examples / case-studies	11:00-11:15	Break
11:40-12:00	Immunogenicity and Macromolecule PK/PD (Shah)	11:15-12:00	Review Module #2: Design & Analysis of a Preclinical
	• Factors associated with immunogenicity		Investigation of Antibody PK (Part 2, Balthasar)
	• Identification of host "anti-drug" antibodies		Development of mechanistic mathematical models
12 00 01 00	• PK modeling	12:00-01:00	Lunch
12:00-01:00	Lunch	1:00-02:00	Application of PK/PD Theory to Guide the Discovery and
01:00-01:30	Use of PK/PD Studies to Support Comparability		Development of New Immunotherapies (Balthasar)
01 20 02 15	Assessments of Therapeutic Proteins (Balthasar)	2:00-3:15	Background: Modeling with ADAPT (D'Argenio)
01:30-02:15	Interspecies Scaling of Antibody PK (Balthasar)	3:15-3:30	Break
	General review of interspecies scaling	3:30-4:15	Review Module #3: Prediction of the Influence of Shed
	Considerations for scaling antibody pharmacokinetics		Antigen on the Distribution of Mab in Solid Tumors
00 15 00 00	• Examples / case-studies		(Balthasar)
02:15-03:30	Mathematical Modeling of Target-Mediated Disposition		Model building exercise
	of Monoclonal Antibodies (Balthasar)		Simulations
	• Introduction to TMD of Mab with examples	4:15 - 5:00	Review Module #4 (Balthasar)
	• Review of mathematical models that have been applied to		Discussion questions and review
	characterize Mab TMD		
	 Comparison of model performance; discussion of 	Day 3	
	implications for predicting Mab PK/PD	9:00 - 9:45	Case Study: Target Mediated Drug Disposition (Mager)
03:30-03:45	Break 3:45 - 4:15 Biologics and Drug-Drug	9:45 - 10:15	Case Study: Antibody-Ligand Interaction and Disposition
	Interactions (Balthasar)		(Balthasar)
	Mechanisms	10:15 - 10:30	Break
	 Examples / case-studies 	10:30 - 11:15	Case Study: Physiologically-Based Pharmacokinetic
04:15-05:00	Review Module #1: Design & Analysis of a Preclinical		Modeling for mAb (Balthasar)
	Investigation of Antibody PK (Balthasar)	11:15 - 12:00	Case Study: PKPD for Antibody Drug Conjugates (Shah)
	 Study objectives 	12:00 - 1:00	Lunch
	 Consideration for study design 	1:00 - 1:45	Modeling Bispecific mAb PKPD (Shah)
	Assay considerations	1:45 – 2:30	Case Study: Denosumab PKPD (Mager)
	• Initial evaluation of data (Additional studies needed?)	2:30 – 2:45	Break
	• Initial characterization of ADME (NCA vs. modeling)	2:45 – 3:30	Estimation Principles & Challenges for Biologics (D'Argenio)
	• Evaluation of NCA results	3:30 – 4:15	Case Study: Non-depleting Anti-CD4 mAb (D'Argenio)

REGISTRATION INFORMATION

4:15

Course location: The course will be held at The Conference Center Niagara Falls, 101 Old Falls Street, Niagara Falls, NY 14303. USA. Phone: (716) 278-2100. Fax: (716) 278-0008. The Center is 28 min from Buffalo International Airport. Website: http://www.ccnfny.com

Reception and Group Dinner

05:00-07:30

Hotel location: *Sheraton at the Falls*, 300 Third St., Niagara Falls, NY 14303. USA. Phone: (716) 285-3361. The price is \$93/night single/double occupancy (triple/quadruple add \$10 per person). *Hotel Deadline: April 8th*, 2019.

Website: https://www.starwoodmeeting.com/Book/Pharmacokinetic2019

Fee: Individual fee: \$2800. This includes course documentation, continental breakfasts, mid-session refreshments, lunches and opening dinner. Graduate Students may enroll at a fee of \$1400 (registered MS and PhD).

Registration: Online registration begins October 1st, 2018.

The course is limited to the capacity of 30 participants. Confirmation email of registration will be returned upon successful registration and payment at the following website: http://pharmacy.buffalo.edu/ under Quick Links.

Discussion / Adjourn

Cancellations: Cancellations with a full refund may be made until March 12, 2019. No refund is possible on cancellations received after this date. Substitutions may be made at any time.

Payment: Mastercard, Visa, American Express, and Discover card payments will be accepted only at the following website: http://pharmacy.buffalo.edu/ under Quick Links. Contact course secretary: Suzette Mis, (716) 645-4831; mis@buffalo.edu, if you need further assistance.





