

In silico Drug Discovery Workshop

hosted by the NCATS Assay Guidance Manual Program

Wednesday, October 23, 2024 – Thursday, October 24, 2024 ~ Virtual (All times are in ET)

AGENDA: Day 1

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11:00 AM	Introductory Comments Joni Rutter, National Center for Advancing Translational Sciences (NCATS), National Institutes of Health (NIH) Kimberly Sciarretta, Biomedical Advanced Research and Development Authority (BARDA)	
11:20 AM	Overview of Workshop Objectives Sarine Markossian and Alexey Zakharov, NCATS, NIH	
11:30 AM	Keynote: Computer-Aided Drug Discovery Then and Now Martha S. Head (Marti), Amgen Inc.	
12:10 PM	Lunch	
12:40 PM		ata sourcing for modeling and validation in in silico drug discovery Markossian, NCATS 15 years of ChEMBL: challenges & opportunities Barbara Zdrazil, ChEMBL Protein Data Bank: From Two Epidemics to the Global Pandemic to mRNA Vaccines and Paxlovid Stephen K. Burley, Rutgers
1:40 PM	Break	
2:00 PM		der Tropsha, UNC Chapel Hill Cartography-guided exploration of (ultra)large chemical spaces Alexandre Varnek, The University of Strasbourg Virtual Screening (TBD) Gisbert Schneider, ETH Zürich Synthesis-aware library design and optimal experimental design Connor W. Coley, Massachusetts Institute of Technology (MIT) Physics Based modeling (TBD) John D. Chodera, Memorial Sloan-Kettering Cancer Center
4:00 PM	Break	
4:15 PM	Session 3: Applications of in silico methodologies in drug discovery Chair: Morgan Sherer, BARDA 4:15 PM	



Summary of Day 1 and Adjourn

5:15 PM

AGENDA: Day 2

11:00 AM Session 3: Applications of in silico methodologies in drug discovery (continued)

Chair: Morgan Sherer, BARDA

11:00 AM Industrializing early drug discovery at scale

Marissa Saunders, Recursion Pharmaceuticals

11:30 AM TBD

Alex Zhavoronkov, Insilico Medicine

12:00 PM **Lunch**

12:30 PM Session 4: Emerging trends in in silico drug discovery

Chair: Alexey Zakharov, NCATS

12:30 PM Understanding the chameleonicity and permeability of PROTACs using Molecular Dynamics

simulations, Markov models, and deep learning Bryn Taylor, Johnson & Johnson Innovative Medicine

1:00 PM Data generation in support of computational drug discovery

Cheryl H Arrowsmith, University of Toronto

1:30 PM Human-Al-Robot Collaboration to Accelerate Materials and Molecular Discovery

Milad Abolhasani, North Carolina State University

2:00 PM Protein therapeutics design (TBD)

David Baker, University of Washington

2:30 PM Break

3:00 PM Panel Discussion: Bridging the gaps in translation

Chair and moderator: Rommie Amaro, University of California San Diego (USCD)

Panelists:

3:00 PM Dangers of poor data (TBD)

Pat Walters, Relay Therapeutics

3:20 PM Team environment for better translation (TBD)

Christopher Austin, GSK

3:40 PM Paying attention to attention in Al generative models

Russ B Altman, Stanford University

4:00 PM **TBD**

TBD

4:20 PM Discussion and Q&A

5:00 PM Closing Session: Summary of Discussions and Perspectives on the Challenges Ahead

Alexey Zakharov, NCATS; Alexander Tropsha, UNC; Rommie Amaro, USCD

5:15 PM Closing Statement and Adjourn