Organizer: Co-Organizer:

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3RD INTERNATIONAL MINI-SYMPOSIUM

ON

MOLECULAR MACHINE LEARNING

APRIL 29[™] 2021 3:00 PM (UTC +1)

Nadine Schneider Novartis, CHE Ola Engkvist AstraZeneca, SWE Nils Weskamp Boehringer Ingelheim, GER

Patrick Riley Google, USA

INDUSTRY **EDITION**

Chair:

Frank Glorius University of Münster, Germany







Nadine Schneider is a researcher in the Computer-Aided Drug Design team in Global Discovery Chemistry at the Novartis Institutes for BioMedical Research (NIBR), Basel, Switzerland. She obtained her PhD in Molecular Modeling in the group of Prof. Matthias Rarey at the University of Hamburg, Germany after receiving a BSc and MSc in Bioinformatics from the Saarland University. In 2014 she joined NIBR for a postdoc focusing on Cheminformatics and Data Science under supervision of Gregory Landrum and Nikolaus Stiefl.



Ola Engkvist is head of Molecular AI in Discovery Sciences. AstraZeneca R&D. He did his PhD in computational chemistry at Lund University followed by a postdoc at Cambridge University. After working for two biotech companies he joined AstraZeneca in 2004. His main research interests are deep learning based molecular de novo design, synthetic route prediction and large scale molecular property predictions. He has published over 100 peer-reviewed scientific publications.



Nils Weskamp is heading a team of data scientists in the Computation onal Chemistry group at Boehringer Ingelheim in Biberach, Germany. Together with his colleagues, he applies a broad range of smart digital tools that support chemists in their daily work. His current research interests focus on explainable machine learning and the user experience of Al. Before joining Bl. Nils obtained a PhD in Computer Science from University of Marburg, Germany.



Patrick Riley is a principal engineer and a senior researcher of the Applied Science team at Google Research. His team collaborates with scientists across organizations to apply Google's knowledge and experience in machine learning and data science to important problems in the natural sciences. Much of his recent work has focussed on machine learning applications for the chemistry of small molecules but has worked in fields as varied as materials science, protein design, and nuclear fusion.

SCHEDULE: THURSDAY, APRIL 29TH 2021

3:00 pm Introduction

3:10 pm Nadine Schneider

Real-World Application of Machine Learning in Drug Discovery

Machine learning models are becoming more and more relevant for drug discovery. There is a big hope that novel algorithms and models can remedy the challenge of finding new drugs more efficiently. A very important prerequisite for this is better understanding the data before applying new models. Another critical point are the users, to make machine learning models effective and education of the users is important to deepen their knowledge and create the right expectations on machine learning models. In this presentation, several of these aspects will be discussed using

3:40 pm **Ola Engkvist**

Applying Artificial Intelligence for Drug Design

lachine learning (ML) and Artificial Intelligence (AI) have impacted many industrial sectors including mpany is drug design. By applying ML/Al together with automation there are opportunities to applying ML/AI to small molecule drug design. Focus will be on deep learning based molecular novo design and synthetic route prediction and the in-house developed platforms REINVENT for novo design and AiZynth for synthetic route prediction.

4:10 pm Break

4:30 pm Nils Weskamp

Smart Digital Assistants to Support Molecule Design:

Data, Algorithms & User Experience

key ingredients are needed to build smart digital tools that support scientists in their daily foundation of well-connected high quality data, smart applications of modern data science s and excellence in user experience. This talk will discuss the state of the art and remaining challenges in these fields.

5:00 pm Patrick Riley

DNA Encoded Small Molecule Libraries Meet Machine Learning

AI/ML methods is often discussed in the drug discovery community. Many modern ML rely on large data volumes, which are often not available. DNA Encoded Small Molecules ELs) can provide so hewhat noisy binding information for up to 10 - molectures, provious in beginning for ML rhodels. In this work, we train graph convolutional neural networks on the property control of the provious provides and used to the pr what noisy binding information for up to 1012 molecules, providing ng data, make predictions on commercial or easily synthesizable libraries, and use automatable filters to choose compounds. In a large prospective study, we show that h is effective with an overall hit rate of ~30% at 30 μM and discovery of potent





Starting Times

Beijing 9:00 pm New Delhi 6:30 pm 3:00 pm Paris 2:00 pm London New York 9:00 am

Please find the registration and Zoom invitation using the following link:

www.uni-muenster.de/Chemie.oc/glorius/symposium_mml.html