

Inverse design of chemical reagents for fast and sensitive detection of pesticides in the environment using mass spectrometry

Henrik Hupatz, Berit Olofsson, Anneli Krueve

henrik.hupatz@mmk.su.se

Stockholm University Center of Circular and Sustainable Systems (SUCCeSS), Svante Arrhenius väg 16C, 114 16 Stockholm, Sweden

BACKGROUND

- Accumulation of polar pesticides in the environment is threatening water quality
- A fast and sensitive analysis is required (Limit of quantification: 0.1 µg/l)
- Mass spectrometry (MS) coupled with chromatography is a sensitive analytical method for a wide range of analytes
- Derivatization can facilitate the analysis of challenging highly polar small molecules

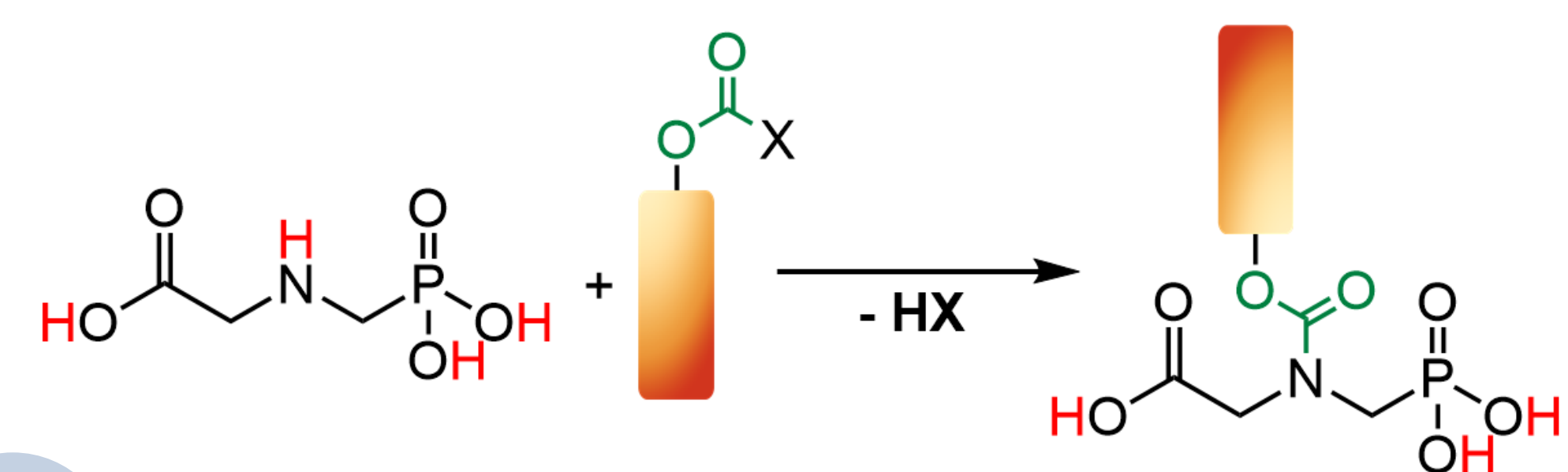


AIM

- Development of a machine learning model for the inverse molecular design of easily accessible derivatizing reagents for glyphosate with high sensitivity and improved sustainability of chemical analysis.
- Verify predicted properties of reagents by synthesis and practical analysis of water samples.

PRELIMINARY RESULTS

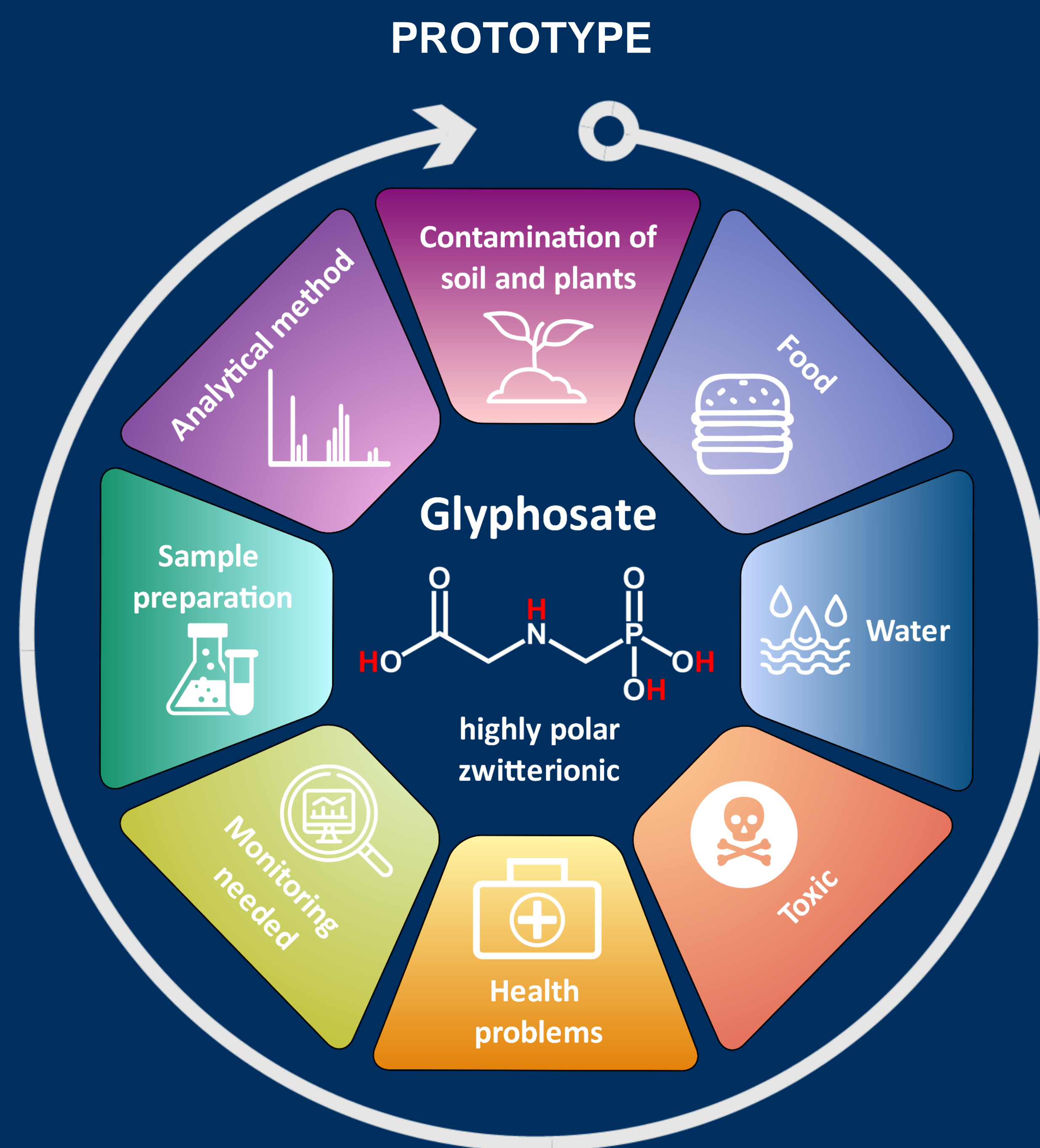
- Sensitivity of an analyte in electrospray ionization (ESI-MS) is expressed with the ionization efficiency (IE)
- Our machine learning algorithm predicts logIE for given structures based on molecular fingerprints or PaDEL descriptors
- Monte Carlo tree search (MCTS) generates structures (orange) with improved range for logIE, logp and synthetic accessibility (SAScore) compared to established reagents



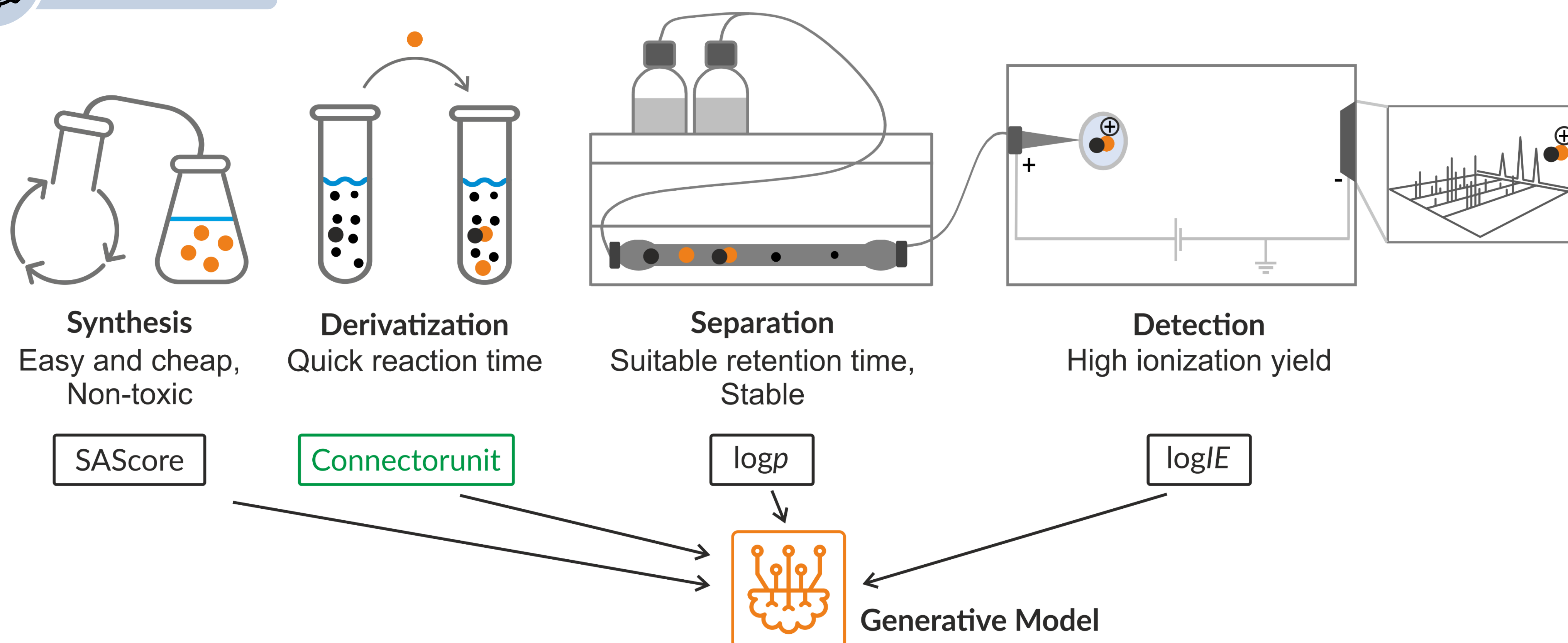
SUGGESTIONS APPRECIATED

Neural network based property-guided generative models might yield higher quality structures and cover a wider chemical space. Which machine learning methods might be well suited?

Can predicted ionization efficiencies facilitate the inverse design of derivatizing reagents for the sensitive detection of polar pesticides by mass spectrometry?

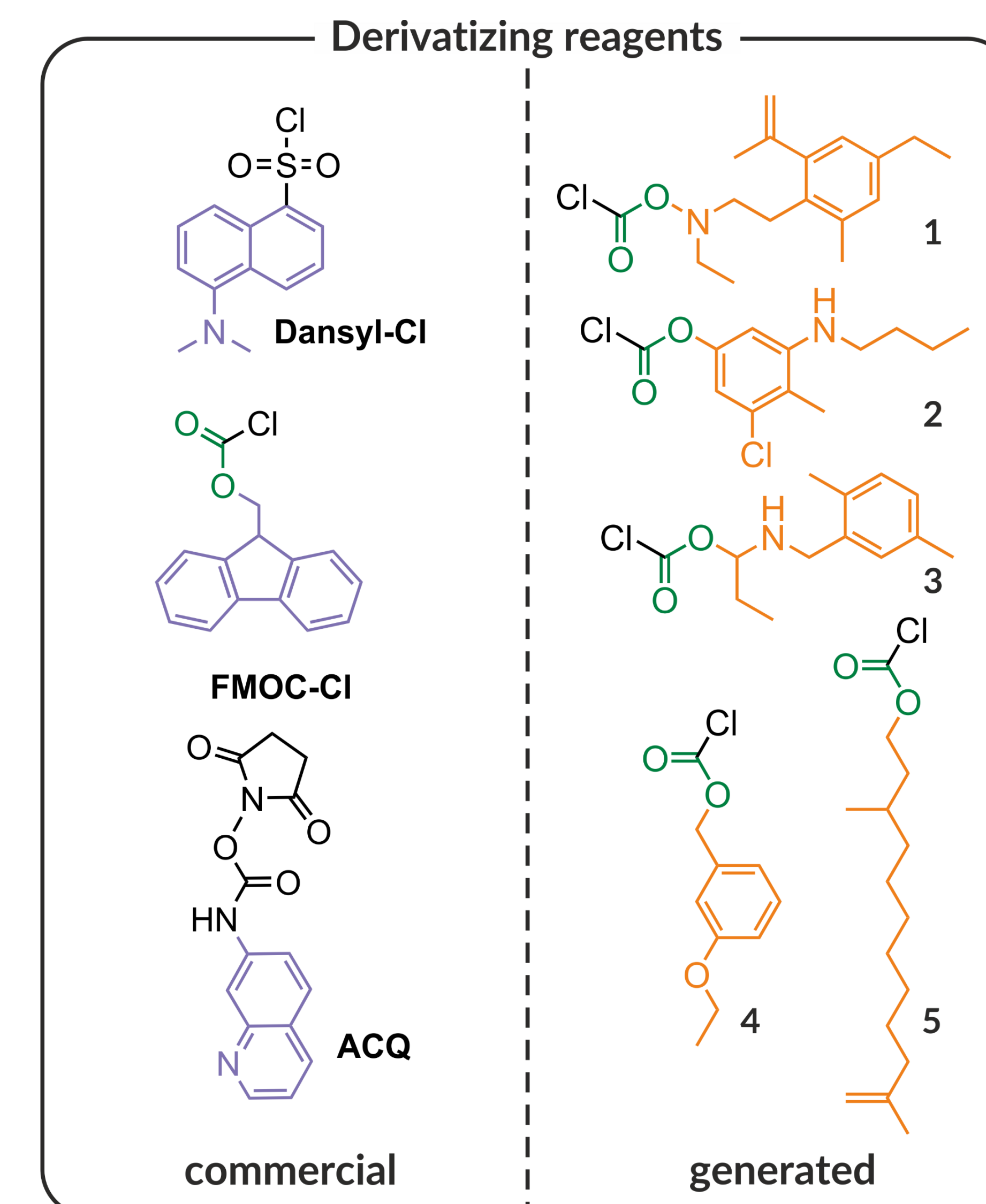
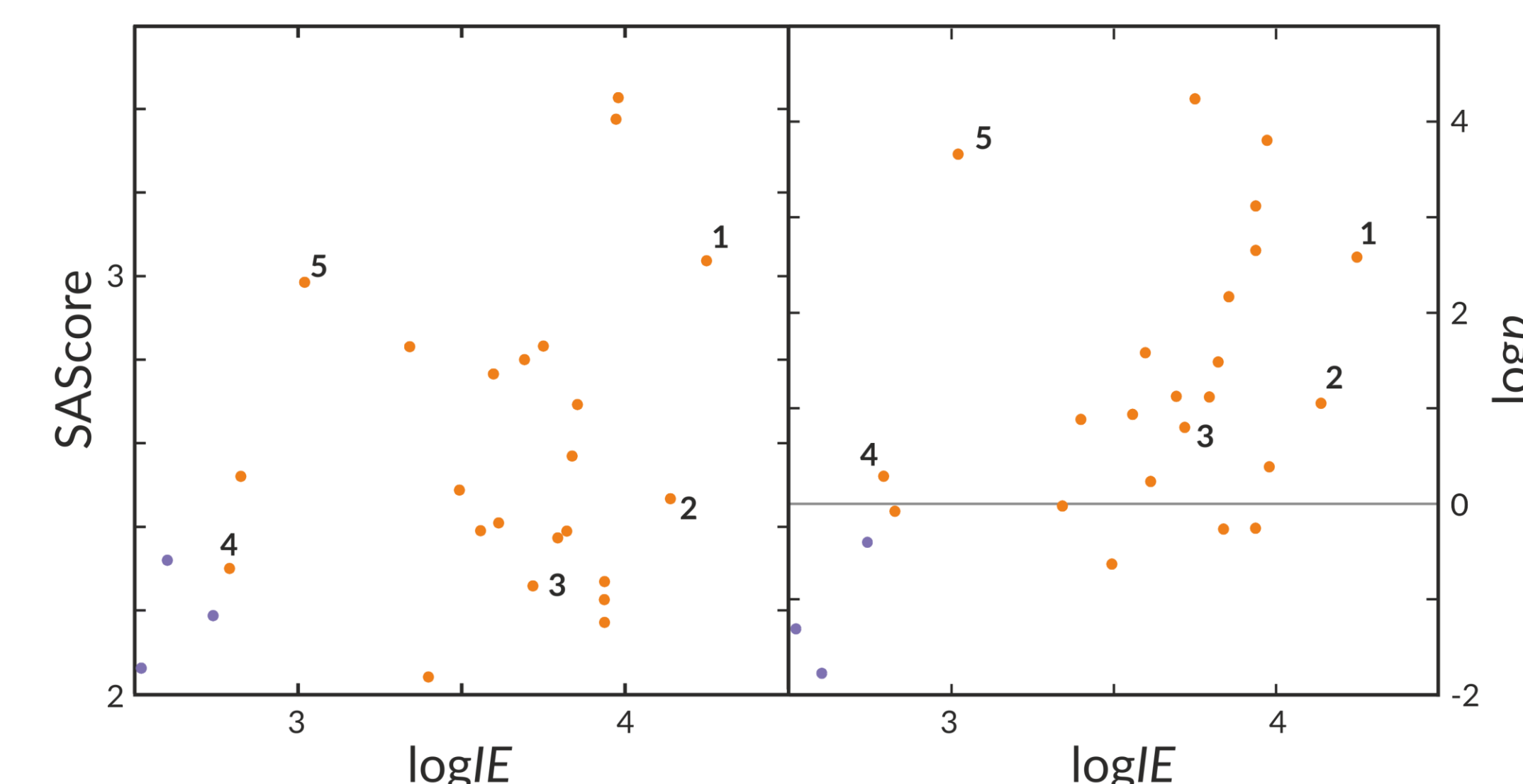


METHODS



MONTE CARLO TREE SEARCH

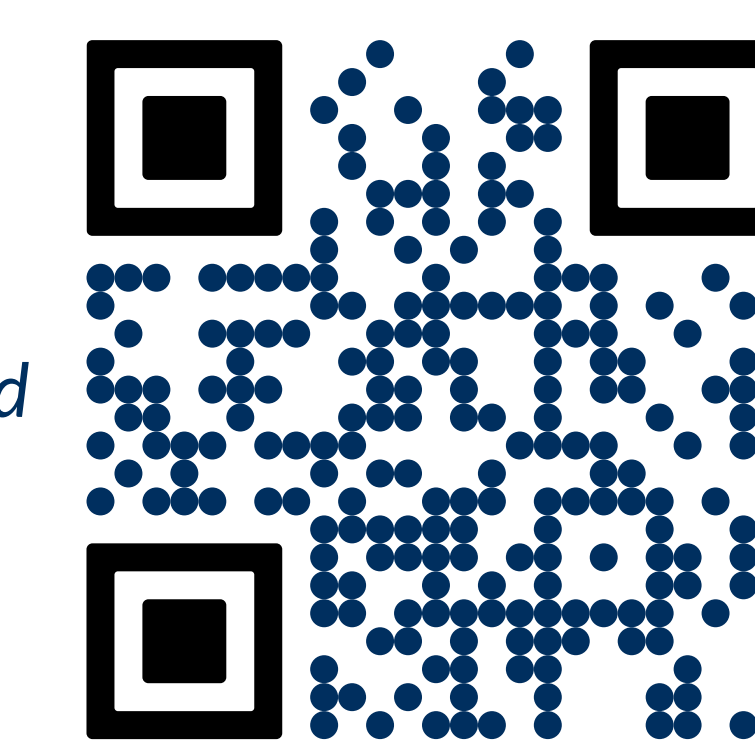
- Molecular graphs generated by MCTS from carbon seed to molecular weight of 400 with MCTS depth of 2 and width of 12
- Evaluation function: $E_v = \log IE - SAScore$
- For each MCTS the structure with highest E_v was picked



GENERATIVE MODEL

- SMILES or molecular graphs?
- Which architecture? Variational autoencoders or recurrent neural networks?
- How to train the model efficiently? Reinforcement learning.

For references, additional information and a poster copy:



Krueve lab



SUCCeSS