

Generative AI for Drug Design: A powerful tool to support medicinal chemists

Ruel Cedeno*, Yaelle Fischer, Dhoha Triki, Philippe Schambel

Department of Computational Chemistry, Novalix, 67000 Strasbourg, France

Abstract

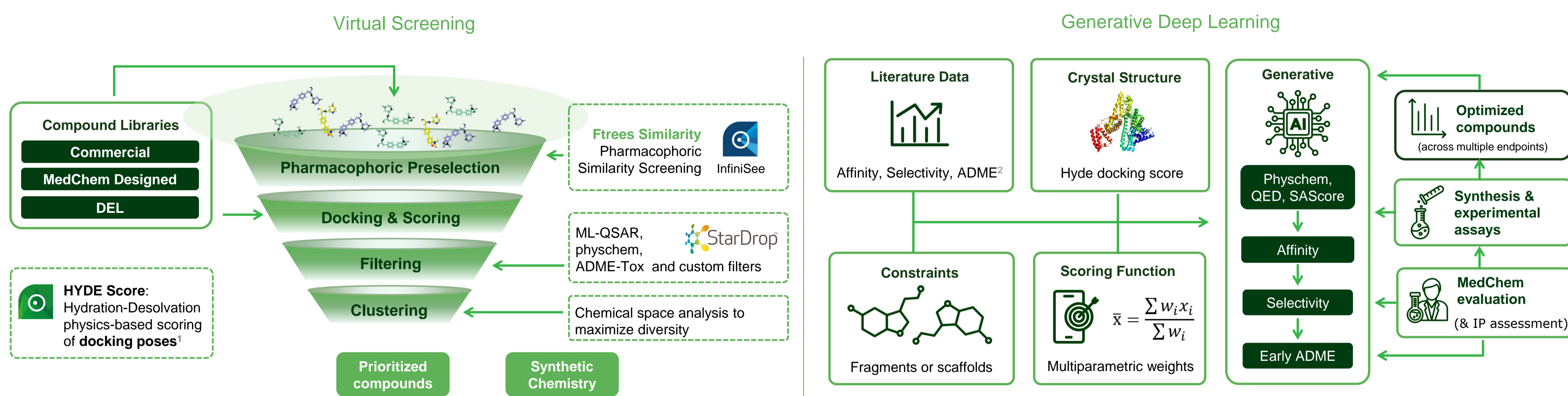
One major challenge in medicinal chemistry is to design compounds that satisfy multiple parameters simultaneously (potency, efficacy, physicochemical and ADME-PK properties, synthesizability, patentability etc). Generative artificial intelligence (AI) has emerged as a promising tool to address this with the help of *de novo* design guided by multiparameter optimization (MPO) algorithms. In this work, by combining the leading opensource packages¹⁻³ and proprietary softwares⁴, we built an automated *in silico* workflow which takes our

project data and structural constraints as input and optimized molecules as output. After performing extensive benchmark tests, we demonstrate that such integrated workflow offers outstanding performance in terms of (1) ability to generate known inhibitors in preclinic and clinical trials (2) druglikeness (3) synthesizability (4) structural diversity, using CCR5 inhibitors as a test case study.

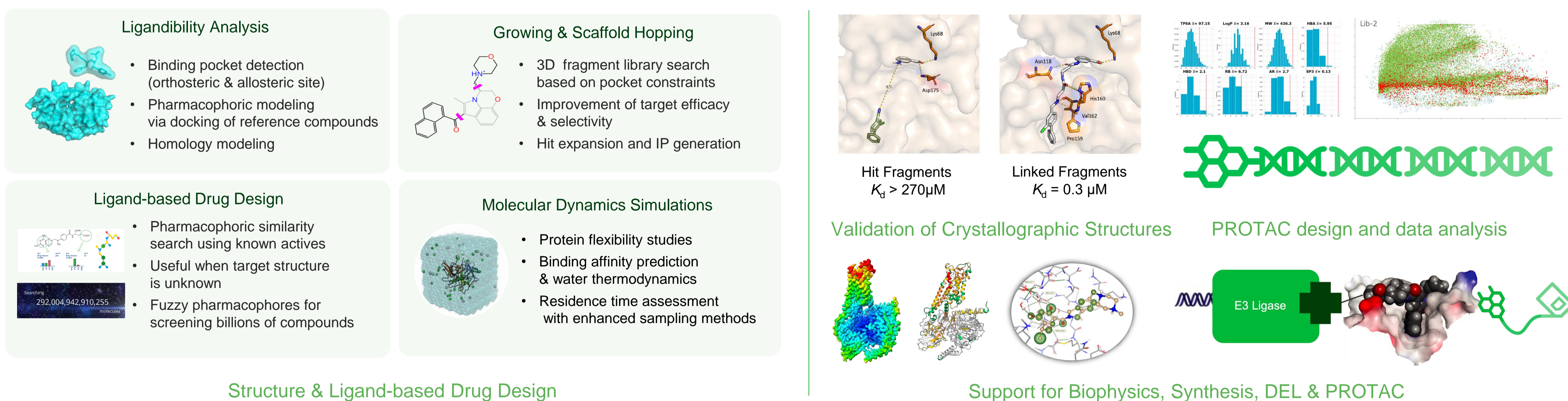
Combining the above results, we integrated our generative AI platform in synergy with virtual screening as well as the classical structure-

based and ligand-based drug design. Furthermore, with internal access to a wide range of biophysics/biochemical assays, augmented synthetic chemistry, DNA-encoded library (DEL) and PROTAC development systems which can generate the important experimental data, such an AI-integrated platform represents a powerful tool for supporting medicinal chemists in accelerating projects from hit identification to lead optimization, enabling the discovery of new medicines.

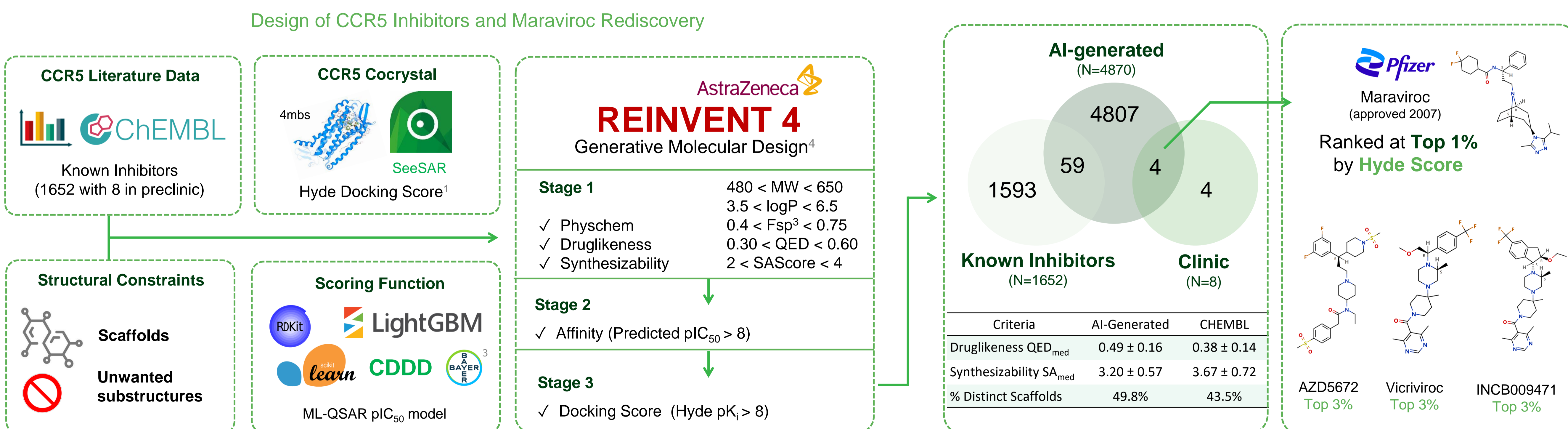
Chem-informatics workflow at Novalix Drug discovery



Hit Identification → Hit to Lead → Lead Optimization



Test case study using generative deep learning



Conclusions & Perspectives

- ✓ We described the implementation of deep learning for molecular design and multiparameter optimization as part of the cheminformatics tools at Novalix.
- ✓ By leveraging a unique combination of leading opensource packages and proprietary softwares, we built an *in silico* workflow which demonstrated excellent performance.
- ✓ This AI-integrated workflow can be coupled with diverse research services at Novalix which are customized to the requirements of our clients.

References

- Schneider, N.; Lange, G.; Hindle, S.; Klein, R.; Rarey, M. A consistent description of Hydrogen bond and Dehydration energies in protein-ligand complexes: methods behind the HYDE scoring function. *Journal of Computer-Aided Molecular Design*, 2012, 27, 15–29.
- Fang, C.; Wang, Y.; Grater, R.; Kapadnis, S.; Black, et al. Prospective Validation of Machine Learning Algorithms for Absorption, Distribution, Metabolism, and Excretion Prediction: An Industrial Perspective. *Journal of Chemical Information and Modeling*, 2023, 63, 3263–3274.
- Winter, R.; Montanari, F.; Noé, F.; Clevert, D.-A. Learning continuous and data-driven molecular descriptors by translating equivalent chemical representations. *Chemical Science*, 2019, 10, 1692–1701.
- Loeffler, H. H.; He, J.; Tibo, A.; Janet, J. P.; Voronov, A.; Mervin, L. H.; et al. Reinvent 4: Modern AI-driven generative molecule design. *Journal of Cheminformatics*, 2024, 16.
- Guo, J.; Knuth, F.; Margreiter, C.; Janet, J. P.; Papadopoulos, K.; Engkvist, O.; et al. Link-INVENT: generative linker design with reinforcement learning. *Digital Discovery*, 2023, 2, 392–408.