

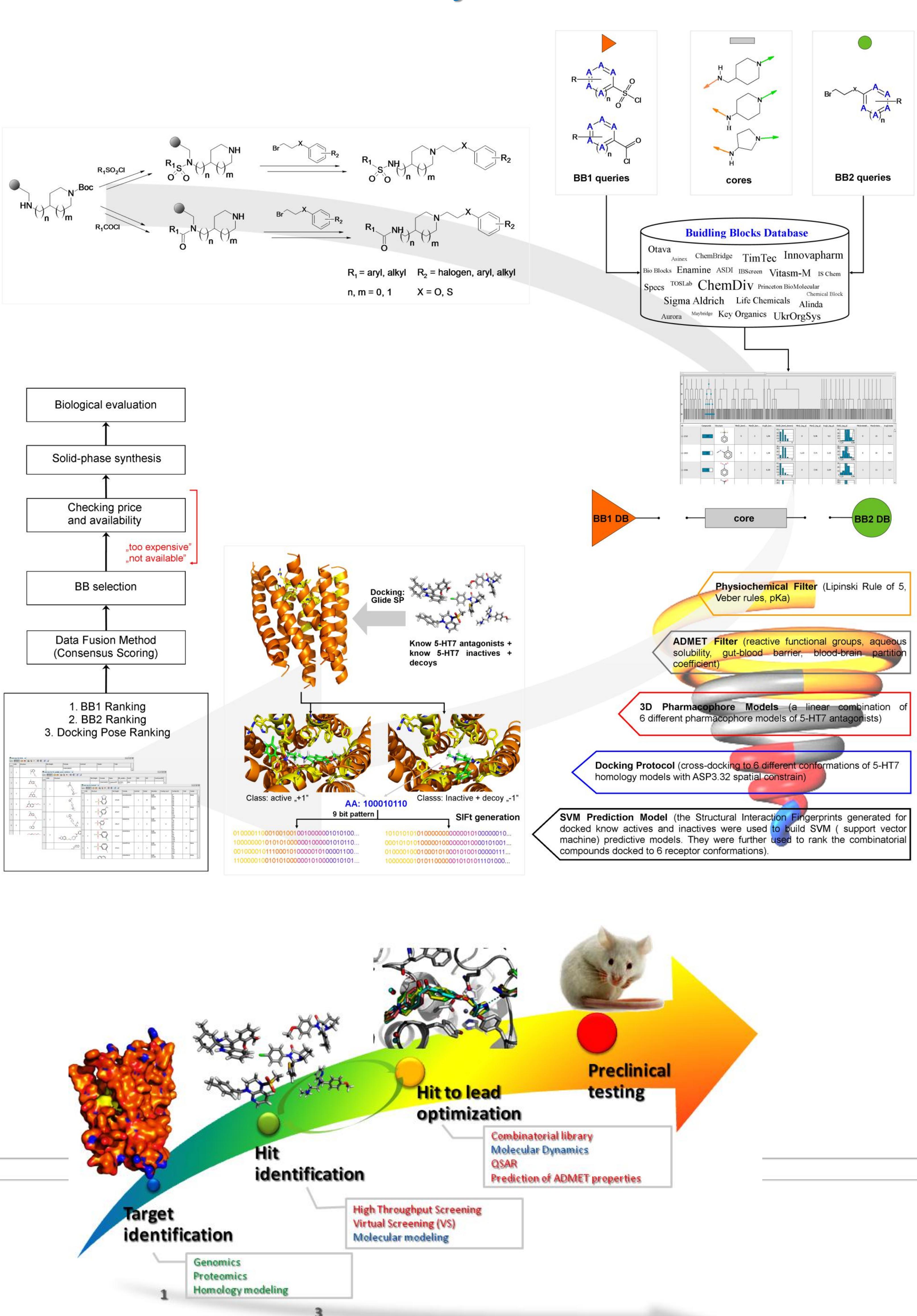
# An application of chem- and bioinformatic methods in modern drug design process



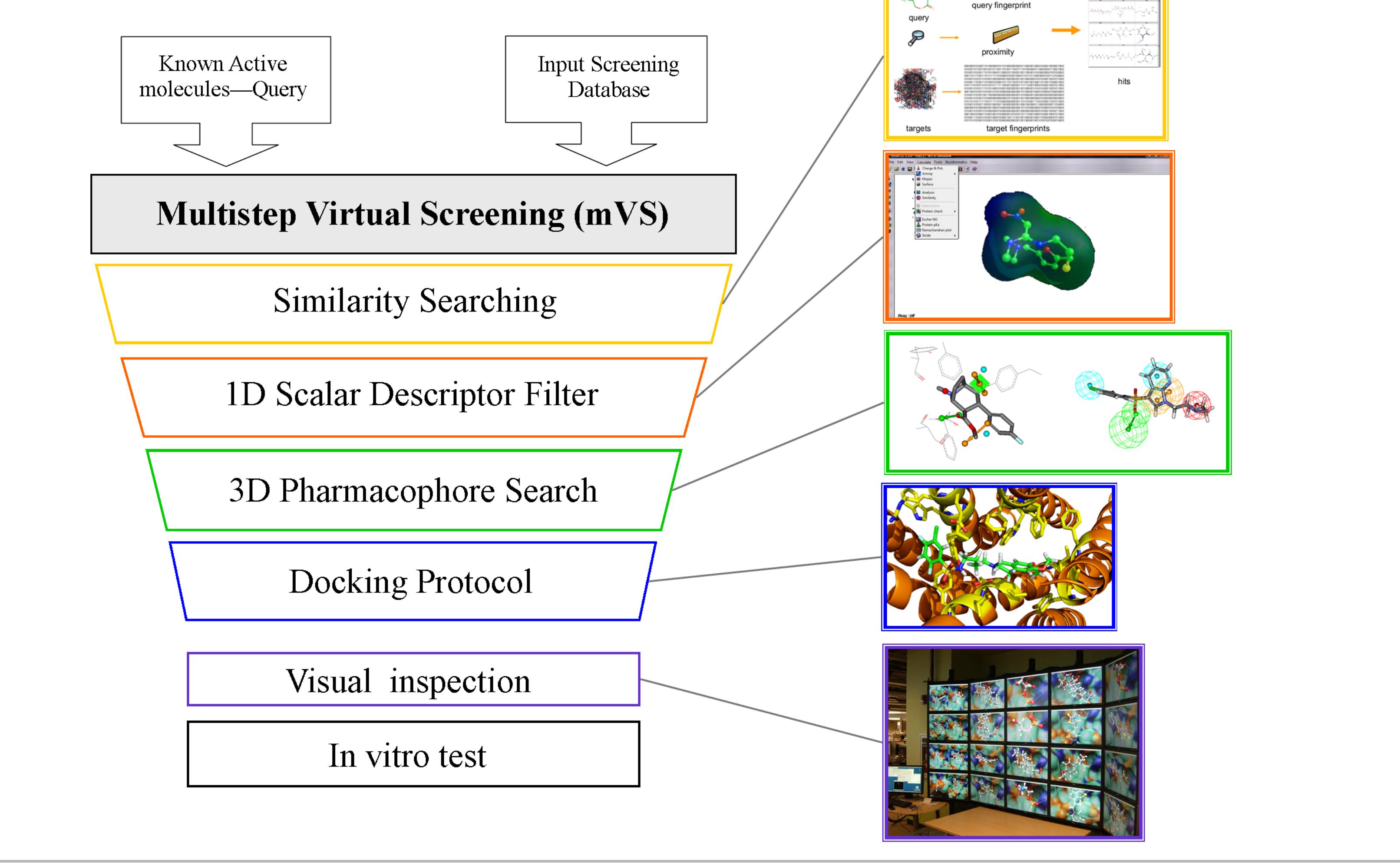
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## Virtual Combinatorial Library Protocol<sup>1</sup>



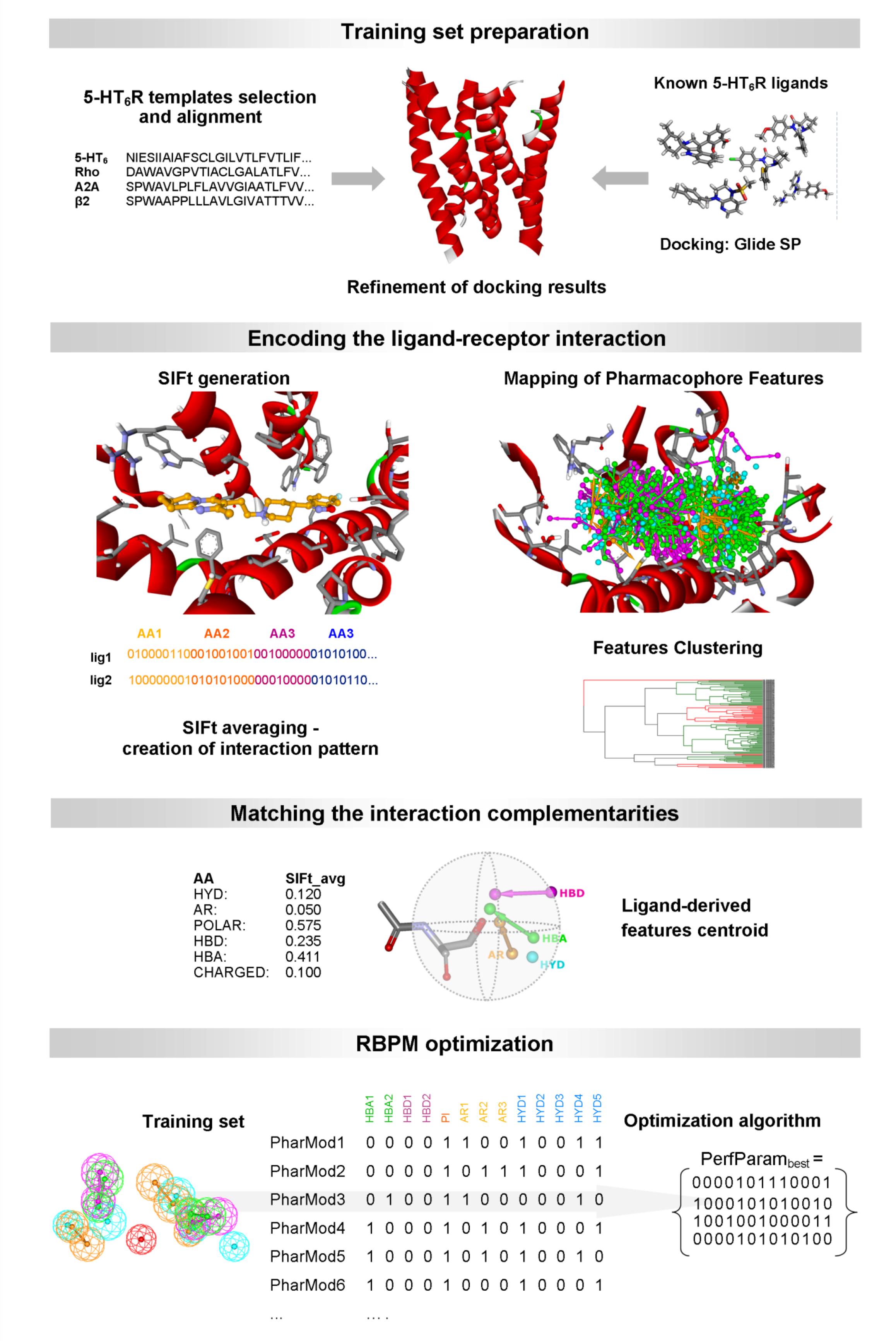
## Virtual Screening Protocol<sup>2</sup>



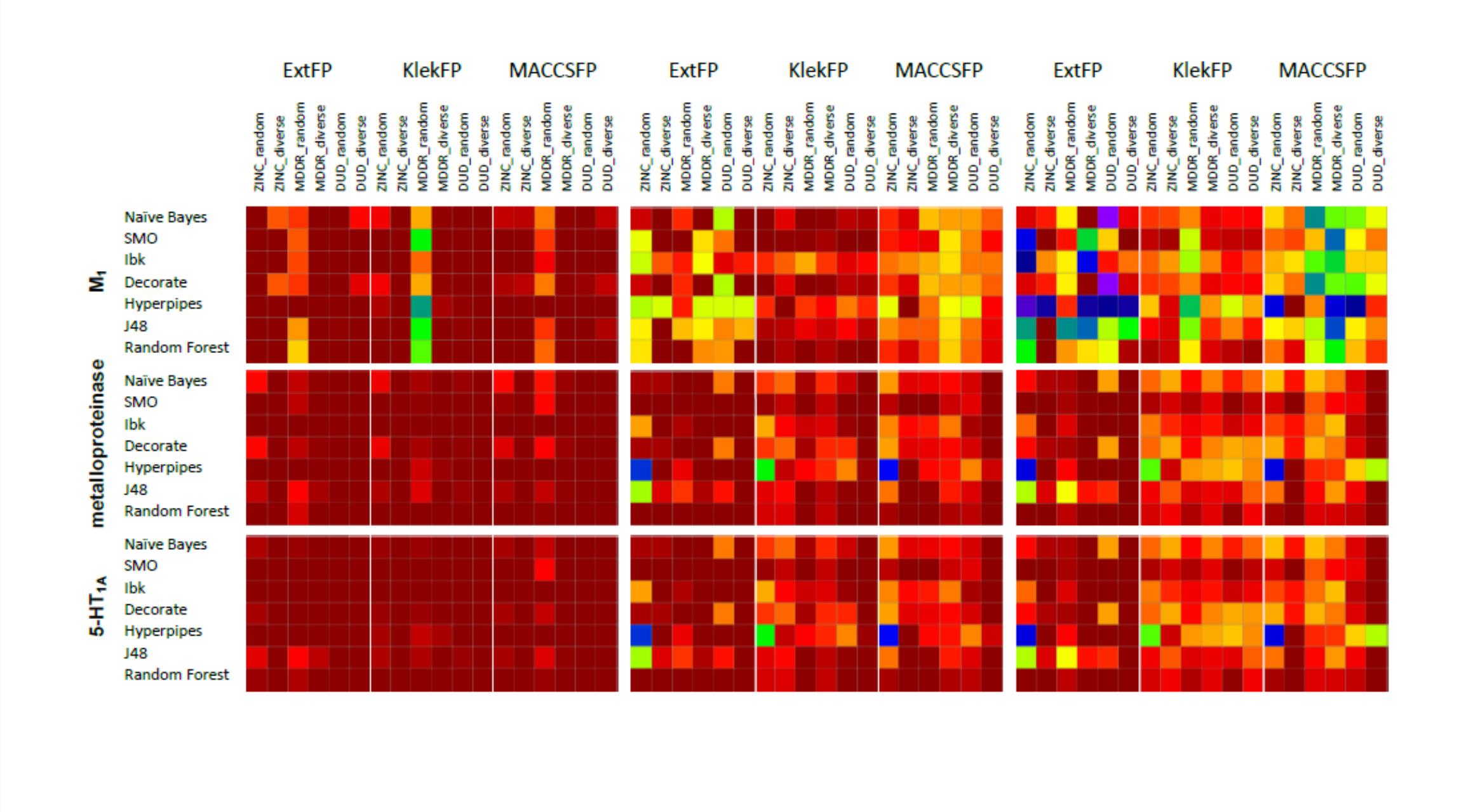
## References

- [1] Zajdel P, Kurczab R, Grychowska K, Satała G, Bojarski A. J., 2012, *Eur. J. Med. Chem.*, 56: 348-360,
- [2] Kurczab R, Nowak M, Chilmonczyk Z, Sylte I, Bojarski A. J., 2010, *Bioorg. & Med. Chem. Lett.*, 20: 2465-2468,
- [3] Kurczab R, Bojarski A. J., The new strategy in receptor-based pharmacophore query definition and its application to 5-HT<sub>7</sub>R, manuscript in preparation,
- [4] Smusz S, Kurczab R, Bojarski A. J., The influence of the inactives subset generation on the performance of machine learning methods *J. Cheminformatics*, in revision.

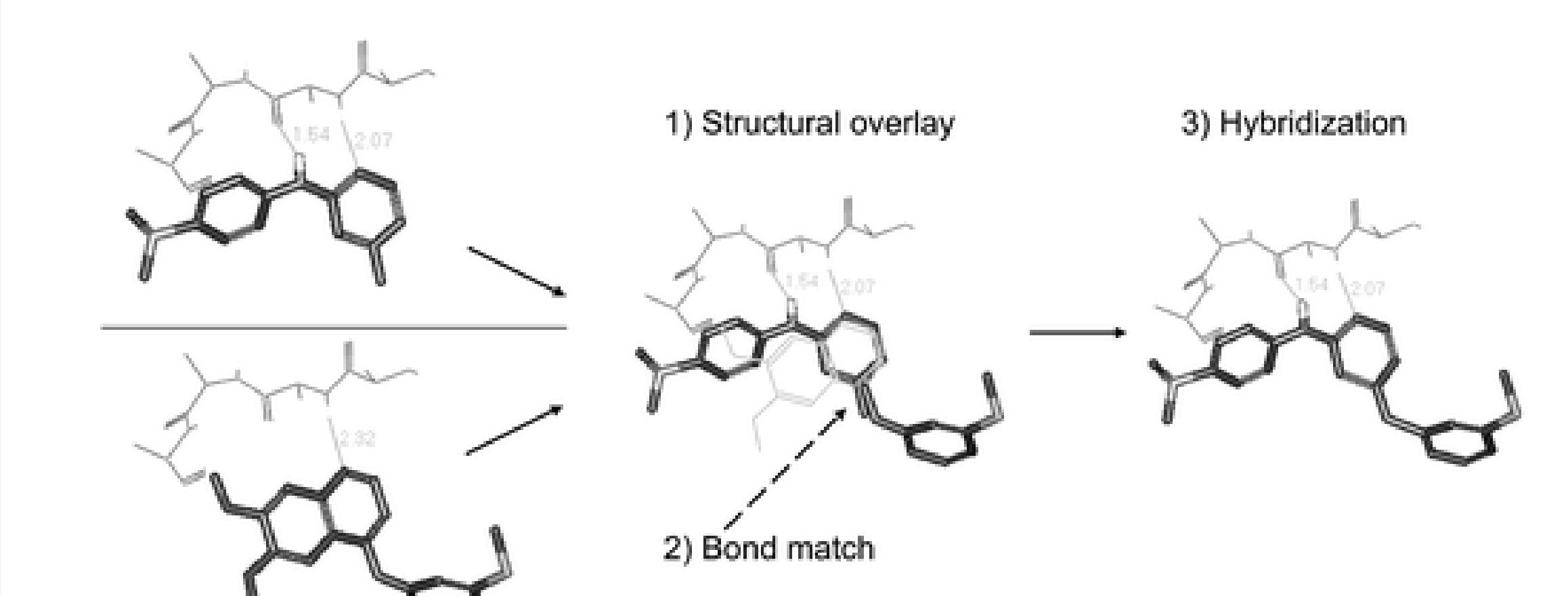
## Receptor-based pharmacophore query definition<sup>3</sup>



## Machine Learning<sup>4</sup>



## Hybridization of known ligands



## Scaffold-hopping

