

## **Internal Webinar**

### **Simulations of ligand binding processes in proteins**

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The presentation will discuss the application, assessment and development of high-throughput simulation protocols to investigate ligand transport processes in proteins. The challenges of rare event sampling for studying ligand binding events in haloalkane dehalogenases and the importance of knowledge-based seeding aided by Markov State Models will be discussed. In addition, the transport tools package will be introduced to address issues related to assessing large datasets of protein-ligand binding processes. Finally, the impact of hydrotropic solvents on cytochrome c dynamics will be explored, offering insights into structure-based drug design and rational protein engineering.

***Thursday, Nov 2<sup>nd</sup> 2023***

***12:00 PM***

