







# Eugen Hruska, Ph.D.








 euhruska.github.io  
 Emory University

 0000-0001-5679-8419  
 @HruskaEugen

 eugen-hruska  
 eugen.hruska@emory.edu

## Publications

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- 1 **Hruska, E.**, Gale, A., Huang, X., & Liu, F. (2022). Autosolvate: A toolkit for automating quantum chemistry design and discovery of solvated molecules. *submitted*.  
 <https://doi.org/10.26434/chemrxiv-2022-px3r8>
- 2 **Hruska, E.**, Gale, A., & Liu, F. (2022). Bridging the experiment-calculation divide: Machine learning corrections to redox potential calculations in implicit and explicit solvent models. *J. Chem. Theory Comput.*  <https://doi.org/10.1021/acs.jctc.1c01040>
- 3 Gale, A., **Hruska, E.**, & Liu, F. (2021). Quantum chemistry for molecules at extreme pressure on graphical processing units: Implementation of extreme-pressure polarizable continuum model. *J. Chem. Phys.* 154, 244103.  <https://doi.org/10.1063/5.0056480>
- 4 **Hruska, E.** (2020). *Adaptive sampling of conformational dynamics* (Doctoral dissertation). Rice University.  <https://scholarship.rice.edu/handle/1911/108744>
- 5 **Hruska, E.**, Balasubramanian, V., Lee, H., Jha, S., & Clementi, C. (2020). Extensible and scalable adaptive sampling on supercomputers. *J. Chem. Theory Comput.*  
 <https://doi.org/10.1021/acs.jctc.0c00991>
- 6 **Hruska, E.**, Abella, J. R., Nüske, F., Kavraki, L. E., & Clementi, C. (2018). Quantitative comparison of adaptive sampling methods for protein dynamics. *J. Chem. Phys.*, 149(24), 244119.  <https://doi.org/10.1063/1.5053582>
- 7 Balasubramanian, V., Bethune, I., Shkurti, A., Breitmoser, E., **Hruska, E.**, Clementi, C., Laughton, C., & Jha, S. (2016). Extasy: Scalable and flexible coupling of md simulations and advanced sampling techniques, 361–370.  
 <https://doi.org/10.1109/eScience.2016.7870921>