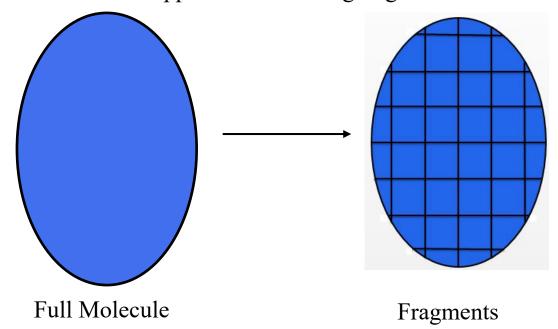
Sibali Debnath

Indiana University Bloomington

- Accurate QM and MD simulations to understand the fundamental interactions in supramolecular systems
- Fragmentation based approach for treating large molecules using accurate QM methods



• MD simulations to investigate the conformational space of complex molecules

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Challenges

- Post-trajectory analysis: Accelerating the analysis of output trajectories
- Visualizing complex (high dimensional) data

Things I want accomplish in the summer institute

- How python can be used for HPC?
- Different techniques to visualize the scientific data
- Learn about GPU computing and programming