

SIMCODES Midterm Presentation

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Acknowledgements

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My Project

Title: Machine Learning-Enhanced Computational Modeling of Metal-Protein Interactions

Description: Focuses on enhancing computational models of metal-protein interactions by integrating quantum chemistry, fragmentation methods, and machine learning. We aim to validate a fragmentation-based strategy, develop machine learning algorithms to predict metal binding in biological systems, and create scalable workflows for future therapeutic screenings.

My Role in the Project

I am the chemistry student in my group. I mainly focus on generating data that the computer scientists use for their work. I do this by utilizing GAMESS and EFP to process systems of large molecules, which calculates the interaction energies between the fragments of molecules to find the total energy.

FRAGMENT-FRAGMENT INTERACTION ENERGIES

CHARGE-CHARGE = -0.0023757819CHARGE-DIPOLE -0.0043063981 CHARGE-OUADRUPOLE = -0.0047534661CHARGE-OCTUPOLE 0.0002179785 DIPOLE-DIPOLE -0.0008127373 DIPOLE-QUADRUPOLE = -0.0022544413QUADRUPOLE-QUADRUPOLE = -0.0014311137OVERLAP PEN. ENERGY = -0.0021016063

ELECTROSTATIC ENERGY = -0.0178175663

REPULSION ENERGY = 0.0153321026

POLARIZATION ENERGY = -0.0042998917

TOTAL DISPERSION ENERGY(E6+E7+E8) = -0.0025801538

E7 DISPERSION ENERGY = 0.0015726003 E6 DISPERSION ENERGY = -0.0023665564 E8 DISPERSION ENERGY = -0.0017861977 CHARGE TRANSFER ENRGY = -0.0012615909

FINAL EFP ENERGY = -0.0106271000

What I've Learned So Far:

Git/GitHub

I've been utilizing Git and GitHub throughout the entire process. I had never used it before.

GAMESS

I use GAMESS throughout the process of calculating energies.



EFP Method

I use EFP to calculate the energies of each fragment in a molecule.

Terminal

I use a Linux terminal everyday. I've learned basic commands and usage.



Progress

GAMESS and EFP

I've ran many databases through GAMESS using EFP. This data has been used by the computer science students to make a neural network.

Amino Acid Optimization

I've created a database of coordinates for every amino acid and all of their conformers. I also optimized these conformers. These will be used for metal-protein calculations.

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RESULTS OF DOM CALCULATIO

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FREE ENERGY IN SOLVENT = <PSI| H(0)+V/2 |PSI>
                                                           -322.9374758713 A.U.
INTERNAL ENERGY IN SOLVENT = <PSI| H(0) |PSI>
                                                           -322.9219772680 A.U.
DELTA INTERNAL ENERGY = <D-PSI| H(0) |D-PSI>
                                                              -0.0154986033 A.U.
                                                              0.0000000000 A.U.
TOTAL INTERACTION (DELTA + ES + CAV + DISP + REP)
                                                              -0.0154986033 A.U.
                                                           -322.9374758713 A.U.
FREE ENERGY IN SOLVENT
                                    -202646.35 KCAL /MOL
INTERNAL ENERGY IN SOLVENT
                                    -202636.62 KCAL/MOL
DELTA INTERNAL ENERGY
                                         -9.73 KCAL/MOL
TOTAL INTERACTION
                                         -9.73 KCAL/MOL
TOTAL FREE ENERGY IN SOLVENT =
                                   -202646.35 KCAL/MOL
```

Challenges

- What I've found most challenging so far is the more detailed computer related tasks.
 I've come across many different issues and error messages that I've needed help debugging.
- Finding a general understanding of the computer science and math focused aspects
 of my project has also been very difficult. I have no background in comp sci and my
 only math experience is Calculus I.

What I Aim To Accomplish

Data Generation

I want to develop a lot more data for the computer science side so that they can have more training data to refine their ML model.

Research Skills

I hope to learn more research-related skills; specifically, how to prepare for a symposium, how to make a poster, how to write an abstract, and how to present at a symposium.