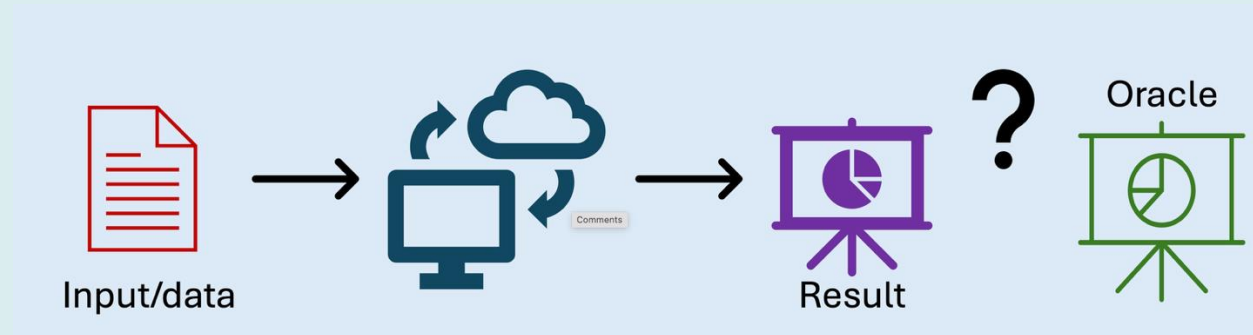


Mentor Introductions

SIMCODES 2025



Building better software testing techniques to improve the quality of software



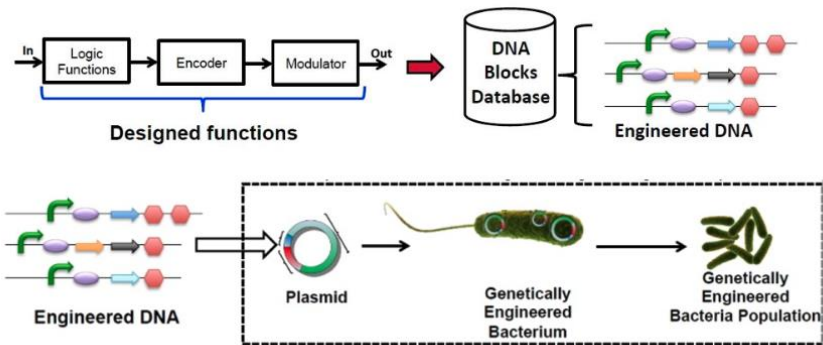
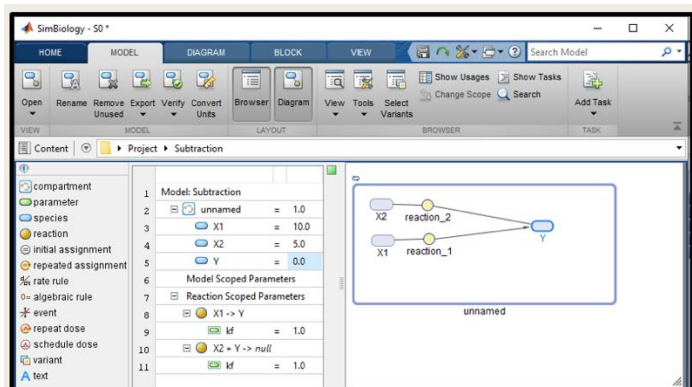
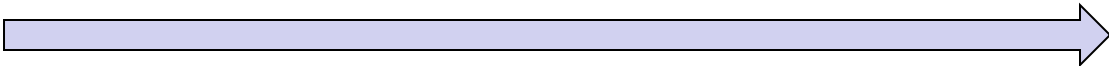
LaVA-Ops

Laboratory for Variability-Aware Assurance and Testing of Organic Programs

<https://faculty.sites.iastate.edu/mcohen/research>



Correct Scientific Software



1 Objective value 0.647004

Model iml_cdg_edit_correct

Media cdg_edit_correct_malts

2 Simulation Information

Method Name: rdidLP

Solution Type: OPTIMAL

Environmental Conditions: Env: CoreBiomol(4)

Objective Function: max R_BIOMASS_Fc_ML_1515_WT_Tp3TM -0.0

Biomass value: 0.0

Net Conversations:

Consumption		
Metabolite ID	Metabolite Name	Value
id: R3_0	run Fc3-1	1000.0
id: R3_0	D-Glucose	10.0
id: R3_0	H2O-H2O	10.0

3 Environment.from_compounds(["malt", "h2o", "o2", "pi", "nh4", "mn2", "zn2", "si

<

FBA(iml1515_model1)

Objective: 0.7665324382387810

Status: Optimal

4

```

medium["EX_co2_e"] = 0
medium["EX_malt_e"] = 10
medium["EX_nh4_e"] = 0
medium["EX_pi_e"] = 0
medium["EX_si_e"] = 0

iml1515_model1.medium = medium

iml1515_model1.slim_optimize()

1.7797995283262238

```

Chemical Reaction Networks

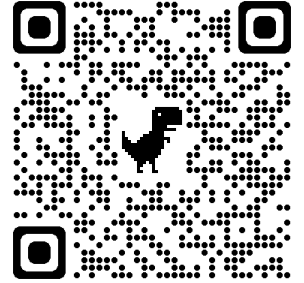
Synthetic Biology

Predictive Biology

++ Computational Chemistry and Quantum Computing

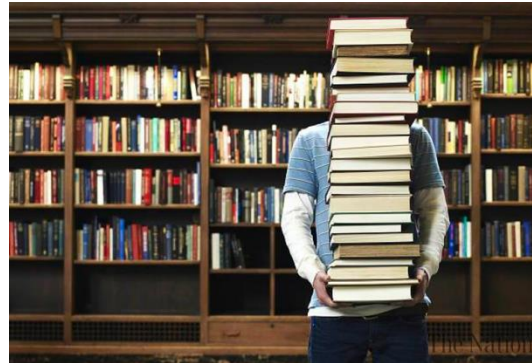
Qi Li

- My name
 - Name can tell a lot
 - Gender, Ethnicity, Age, Religion, ...
 - Important feature for many NLP tasks
- My background
 - BS. Math
 - MS. Stats
 - PhD. CS

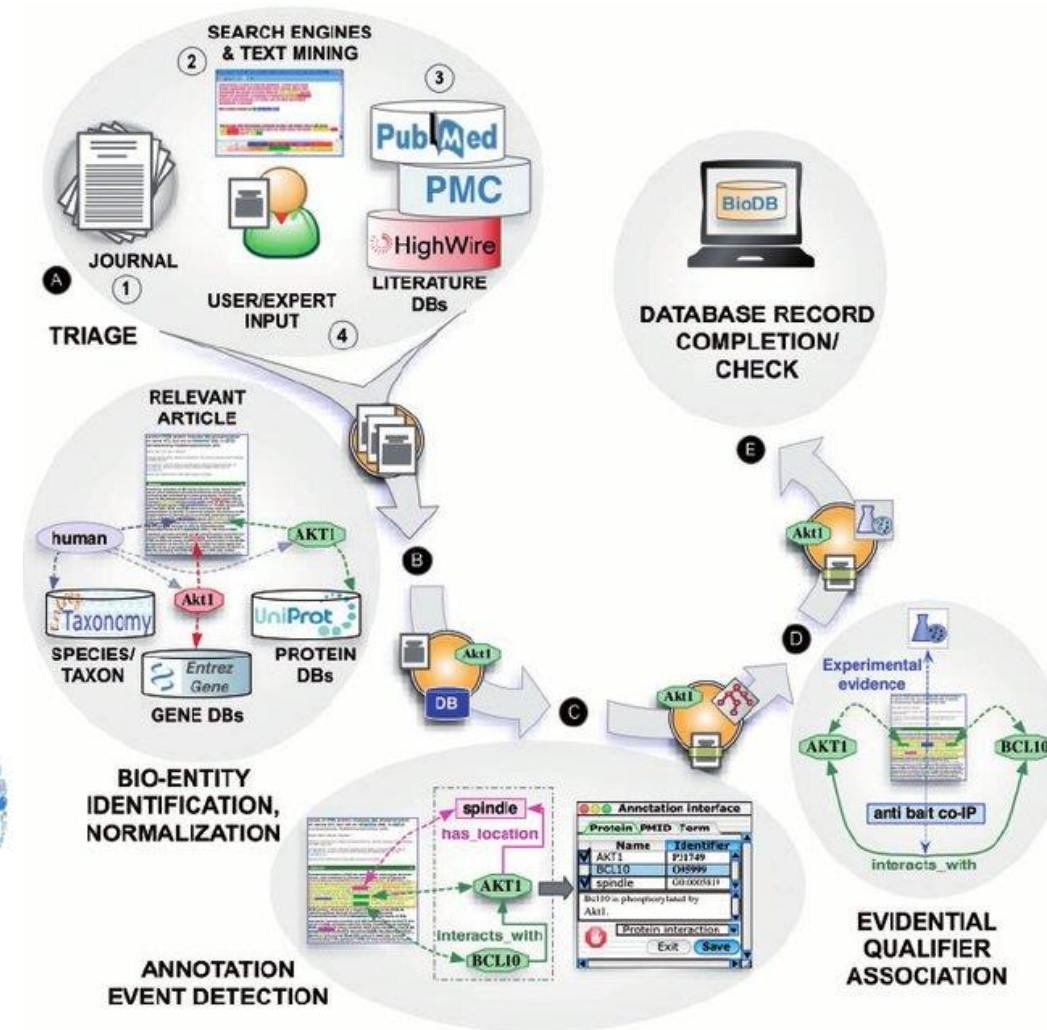


My research

- NLP -- Information extraction
- Generation models
- Applied ML



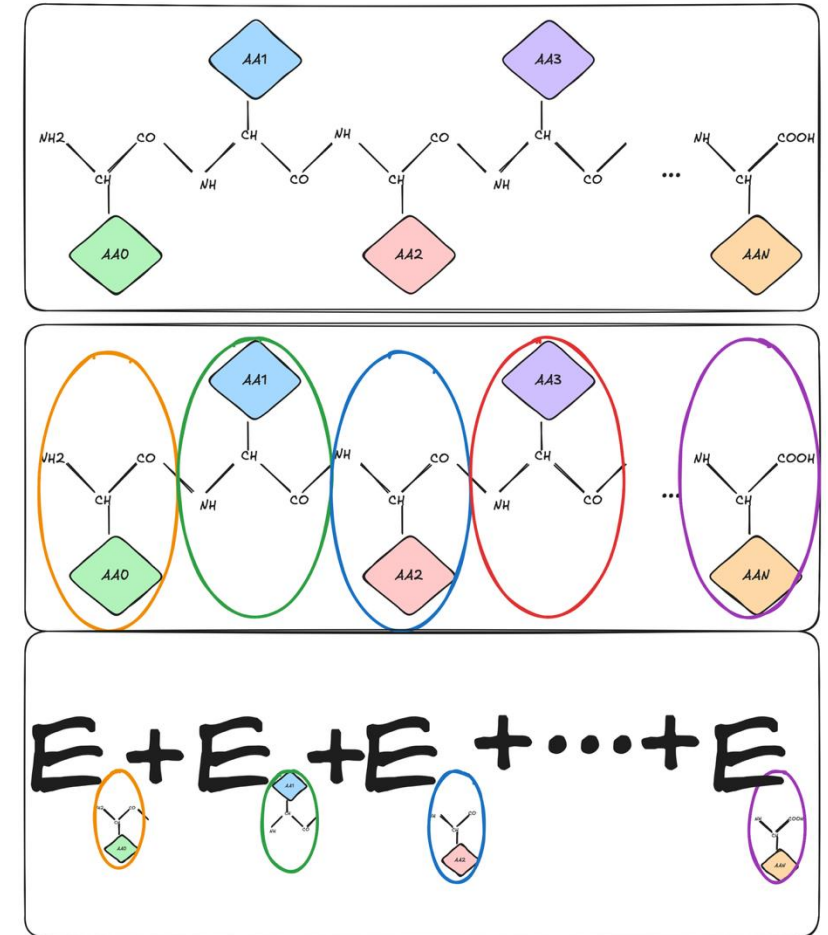
Knowledge
& Insights



SIMCODES Project:

Developing components for automating fragmentation

- Mentee: Gabriela and Devarsh
- Goals:
 - Function for identifying amino acids in a protein.
 - Function for splitting a protein by amino acid.
 - Idea: protein as an unnatural language

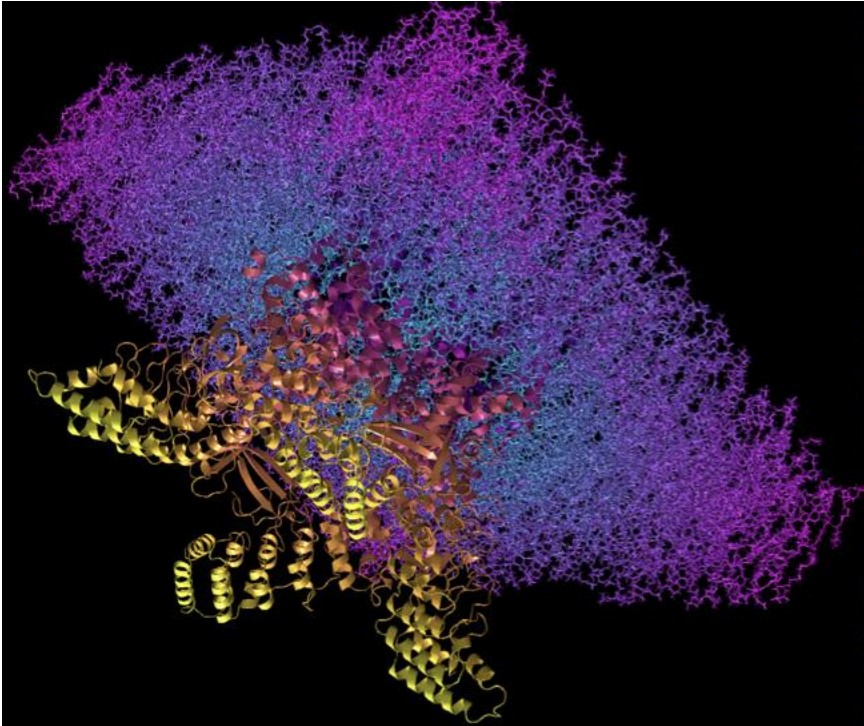


Computation, AI & Theory-Driven Biophysics (CATBio 🐱 🧬)

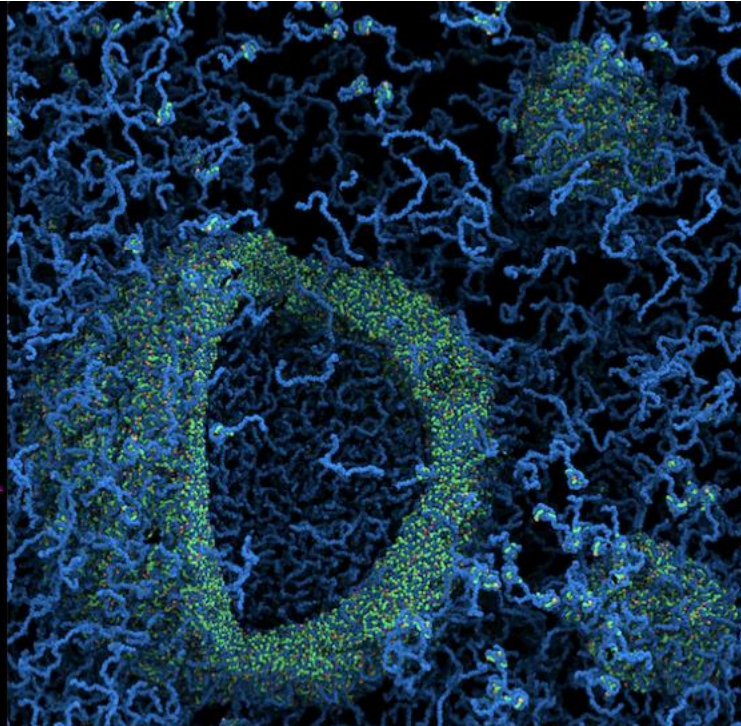
1. Allosteric, sensing and adaptation of protein dynamics to environments

2. Dynamics and thermodynamics of biomolecular condensates

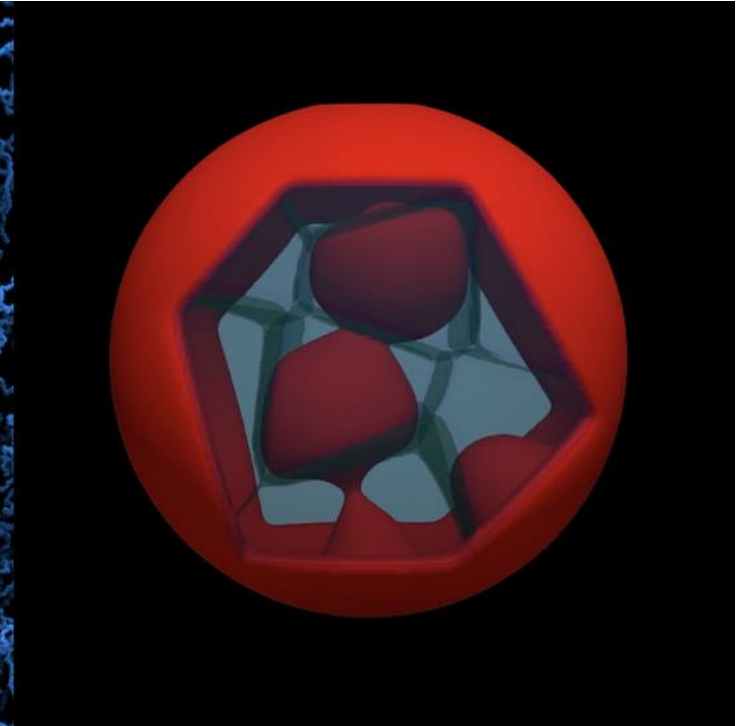
3. Mesoscale models of chromatin organization and dynamics



MD simulations,
bioinformatics, AI



MD simulations,
Soft-matter theory,
Model development



Phase-field and reaction-
diffusion simulations,
model development

PotoyanGroup



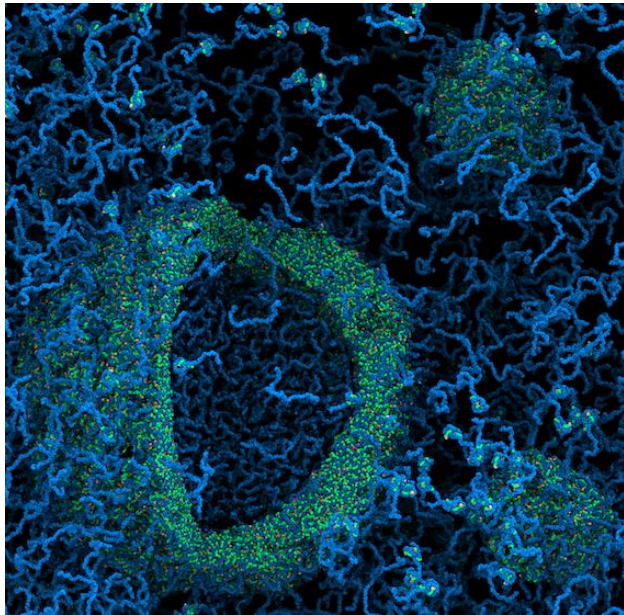
RESEARCH CORPORATION
for SCIENCE ADVANCEMENT



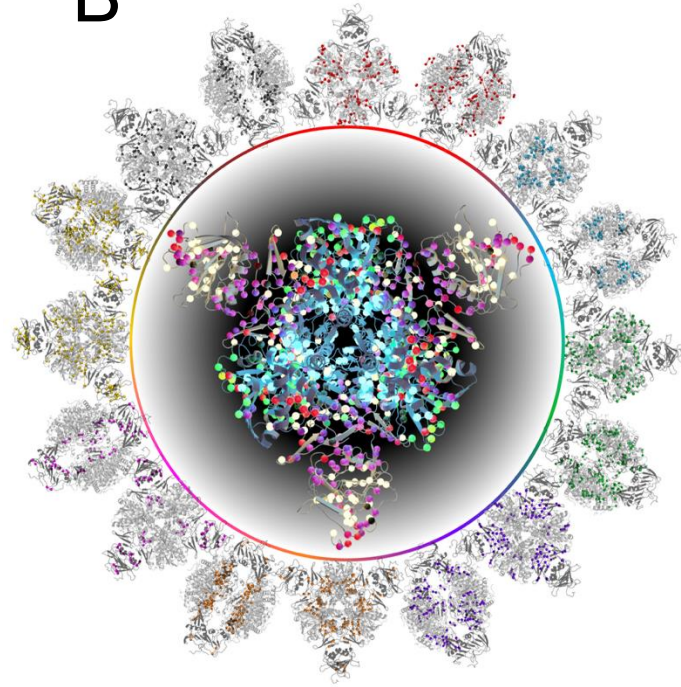
National Institutes
of Health

Iowa State University - Department of Chemistry

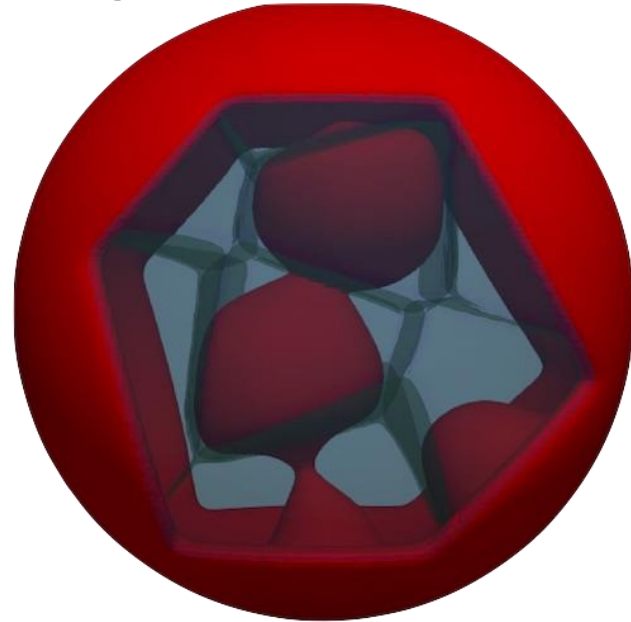
A



B



C





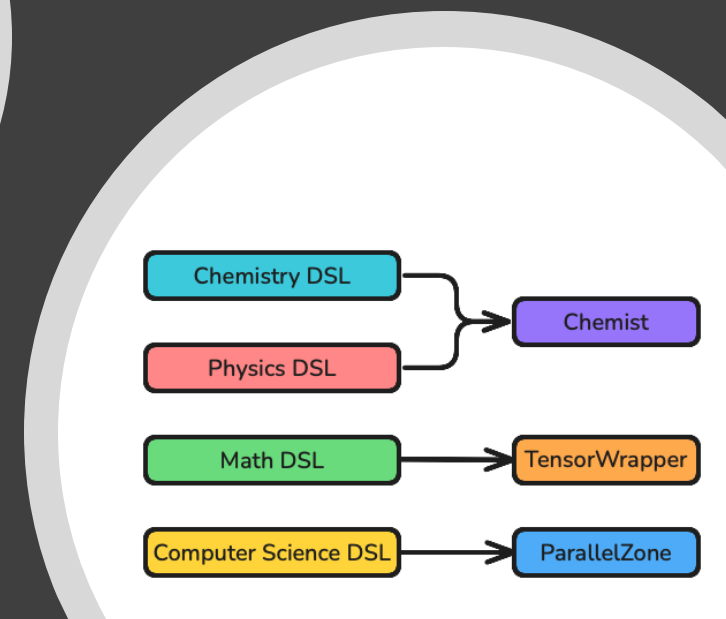
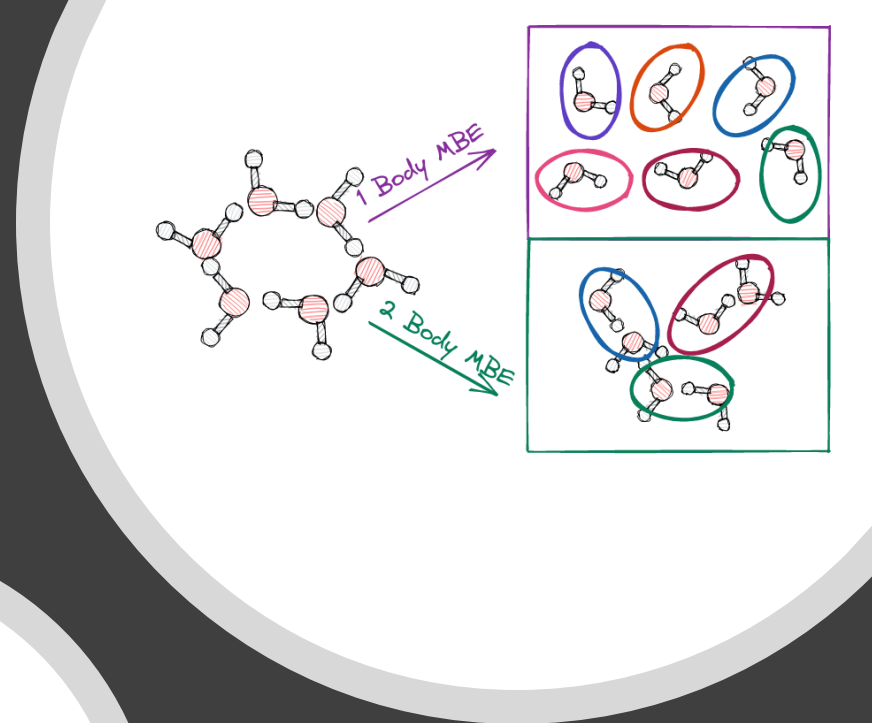
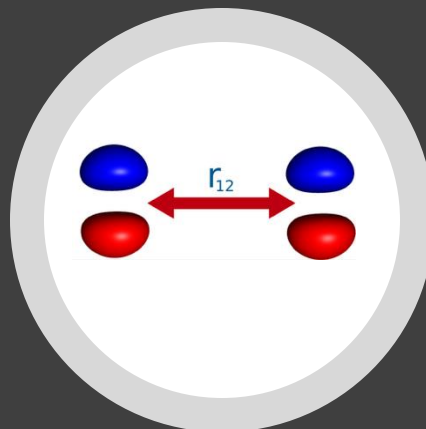
Ryan Richard

- Scientist at Ames National Laboratory and Adjunct Professor of Chemistry at Iowa State University.
- Background: Grew up in Ohio, PhD in chemistry from “The” Ohio State University.
- Interests: travel, technology, video games, exercise, being told what to do by my dogs.



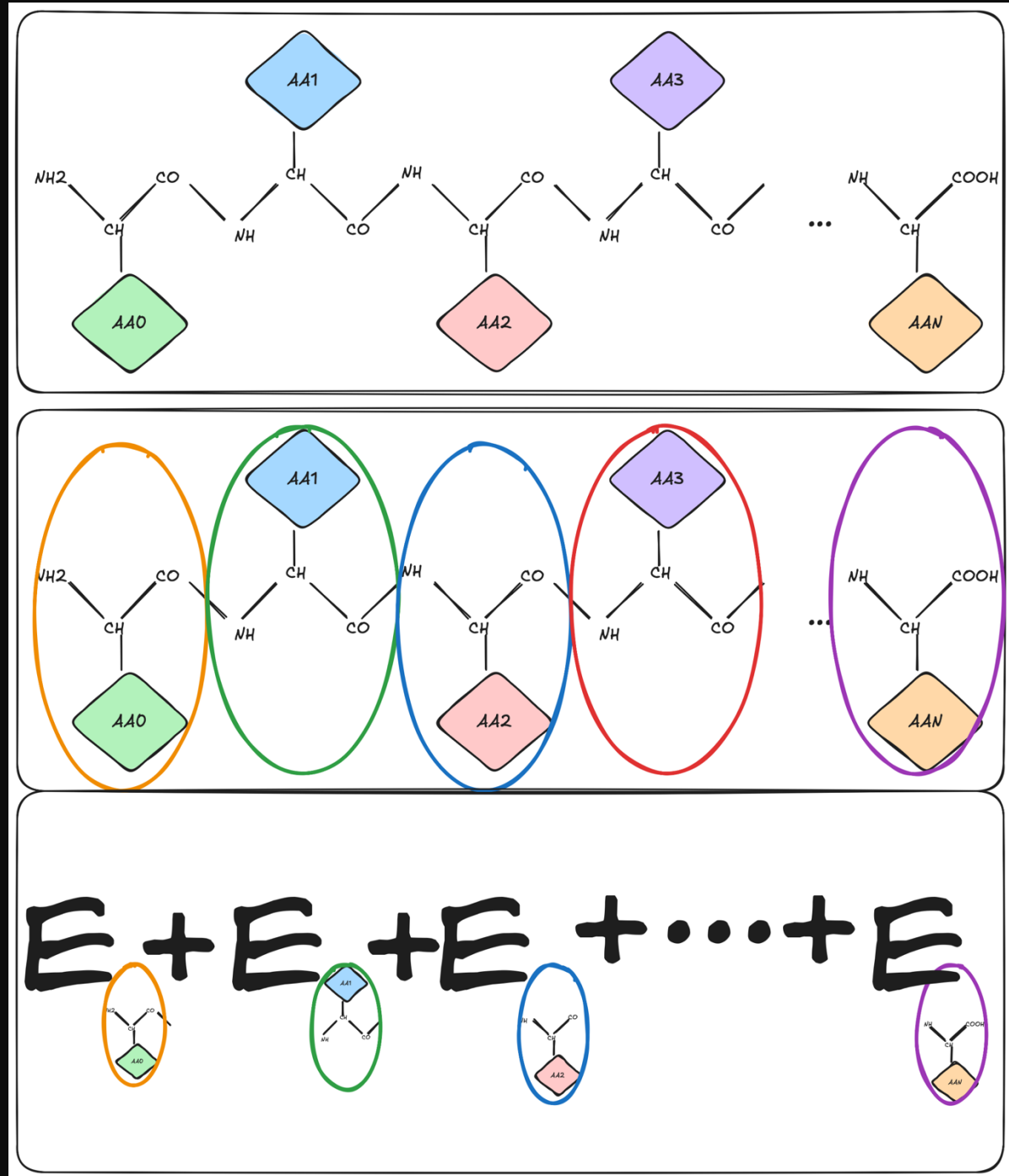
Research in the Richard Group

- Research interests: Sustainable scientific software development, high-performance computing, reduced scaling electronic structure theory.
- Develop for NWChemEx ecosystem.
- Strong emphasis on software engineering applied to theoretical chemistry.



SIMCODES Project: Developing components for automating fragmentation

- Mentee: Daniel Woodard
- Summary: Winner-take-all battle between Daniel's chemical intuition and the best AI/ML model Gabriela and Devarsh can make.
- Goals:
 - Develop tools for reading PDF files.
 - Function for identifying amino acids in a protein.
 - Function for splitting a protein by amino acid.





The Richard and Windus Groups

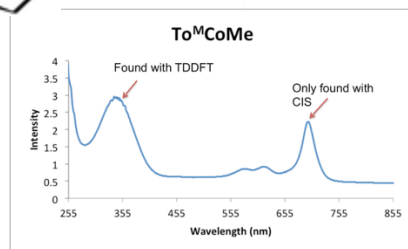
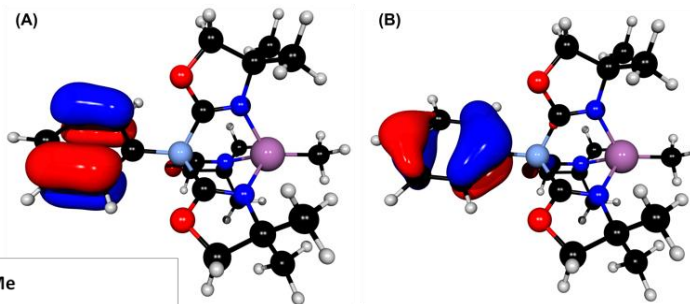
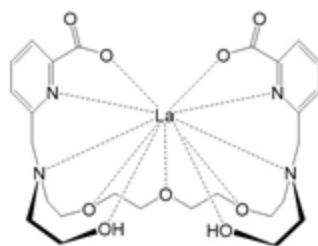
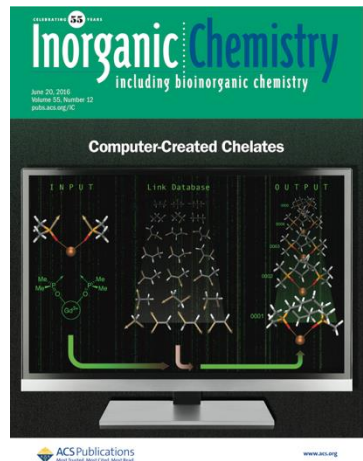
Research in Song's Group

Statistical Mechanics and Biophysics at Iowa State University

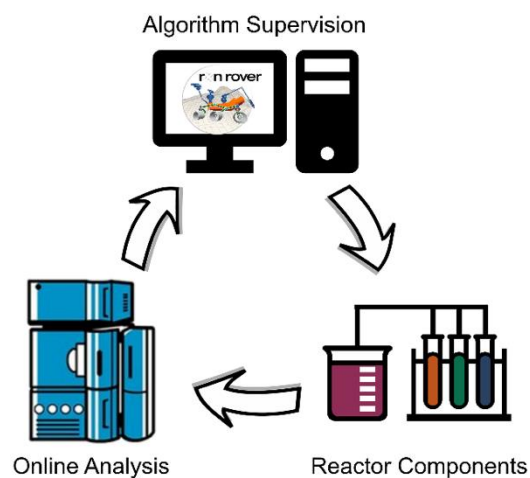
- Electron Transfer and Energy Transfer in Photosynthesis and Electric Cells, electron transport in DNA
- Solvation Dynamics and Dielectric Fluctuations in Inhomogeneous Dielectric Materials, such as Protein and Nucleic Acids
- Theory of Protein Crystallization and protein-protein interaction
- Molecular Debye-Huckel theory of ionic fluids
- Phase behaviors and nucleation
- Sub-diffraction imaging using heterodyne and entangled photons
- Studies of nano-domains of supercooled liquids using machine learning

The Windus Group

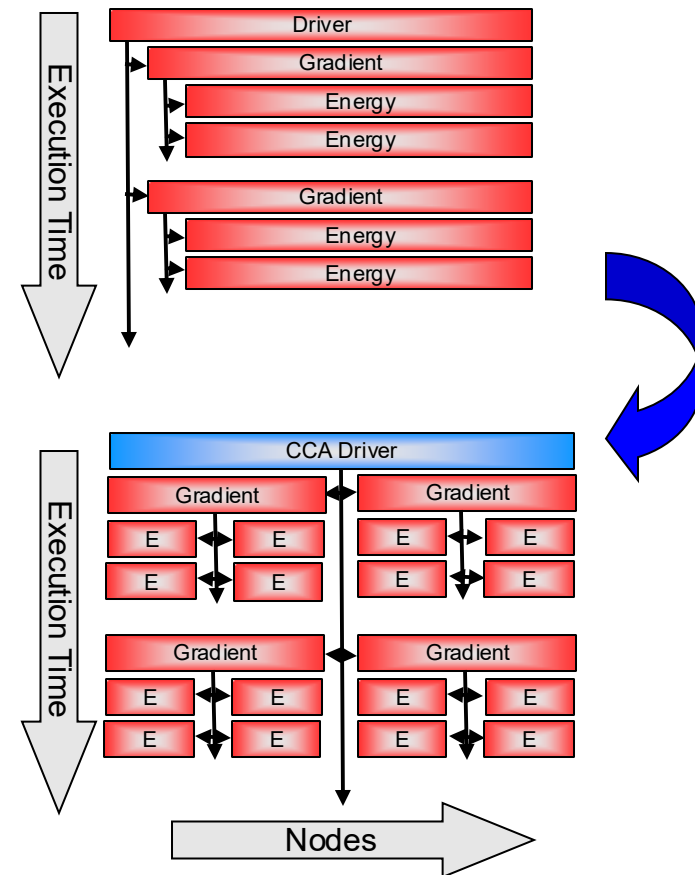
Separations and catalysis challenges



Reaction optimization and lab automation



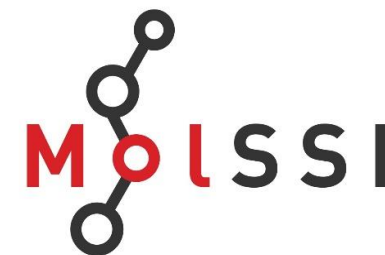
High performance, exascale computing



The Windus Group and Funding



NWChemEx





- What is your area of research?
 - Theoretical and Computational Chemistry
 - developer of General Atomic and Molecular Electronic Structure System (GAMESS) package
 - Reduced scaling electronic structure theory
 - Heterogeneous high-performance computing and energy efficient computing
 - Research Interests
 - Protein-ligand interactions; drug design
 - Heterogeneous catalysis modeling
 - Radical chemistry
 - Quantum computing and machine learning
 - How do you relax at the end of the day?
 - Play with my kid; play games
- Scientist II, Ames National Laboratory
 - Adjunct assistant professor, Department of Chemistry
 - pxu@iastate.edu
 - pxu@ameslab.gov