

### **SKILLS**

- Coding: intermediate C++ & Python, (STL, object orienting programing, numpy, pandas, matplotlib). Familiar with Linux command line/ bash/ cloud computing, vim, git. [6+ years]
- Computational protein design/ engineering
  - o Extensive knowledge of the **protein design** (sequence prediction) problem. [6+ years]
  - Implementing an **entropy based statistical algorithm** in C++ to solve the design problem. [6+ years]
  - o Using molecular dynamics simulations (NAMD) to investigate protein interactions. [6+ years]
  - o Protein visualization and animation with VMD & Pymol (render materials, lighting, electrostatic surfaces, RMSD & heatmap analysis, movie making). [6+ years]
  - o Collaboration with protein experimentalists, applying theoretical models to interpret data (Mass, CD, melting temperature, PI, charge-PH relations). [5+ years]
- Bioinformatics/Genomics
  - Genomics concepts: blogs on NGS techniques & applications; genetics exon, intron, splicing; epigeneticstranscription factors, promoters, histone packing, transposons; GWAS, DNM; cell biology; etc.
  - o Algorithms for DNA sequences: blogs and coding projects of naïve exact match, Boyer-Moore, index assisted matching, edit distance and Smith-Waterman dynamic programming algorithms.
  - o Common software & data processing: blogs and coding projects regarding SAM/BAM, BED, GFF/GTF, VCF/BCF, FastA/Q files; alignment and variant callings with samtools, bfctools, bedtools.

#### **EDUCATION**

### University of Pennsylvania

Ph.D., Physical Chemistry - GPA: 3.90/4.0

Advisor: Dr. Jeffery Saven

Thesis: Computational engineering of protein features: charge variation, host-guest assembly, and structural motif.

## University of Science and Technology of China

**Expected Jun 2023** 

Advisor: Dr. Tongwen Xu

B.S., Chemistry – GPA: 3.63/4.3

Outstanding Student Scholarship for 4 years (2015 – grade 1 scholarship, top 3%; 2014 & 2013 – grade 2 scholarship top 10%; 2012 – grade 3 scholarship, top 25%)

### RESEARCH EXPERIENCE

## Graduate Research Associate - University of Pennsylvania

Jan 2017 - 2023

Focus: Applying a statistical model to solve the protein design problem, creating proteins that possess desired functionalities, such as high thermal stability, enzyme activity or host-guest assemblies.

- Computational Protein Design (2 projects 18 designed proteins 16 confirmed to fold correctly)
  - 1 Designed a wide variation of charges for a short peptide of 29 amino acid residues (~3600 Da).
    - 17 peptide sequences calculated. 15 of the 17 sequences folded as desired (88% success rate).
    - 13 of the 18 exterior residues (72%) in the peptide were modified without disturbing the folded structures.
  - 2) Designed a super-positive variant of human carbonic anhydrase II enzyme with net charge of +21 (natural enzyme has -1 charge).
    - This is the first example of a computationally designed super-positively charged enzyme.
    - 20 out of the enzyme's 70 exterior residues (28%) were mutated, without compromising the enzyme's folded structure or eliminating functionality (81% activity retained).
    - The super-positive enzyme variant can be directly encapsulated within a ferritin cage (negatively charged), without adding a super-positively charged fusion partner (previous encapsulation method).

- o The encapsulation leads to enhanced thermal stability and activity (63% enhanced at 40 °C).
- Molecular dynamics (MD) simulations (Uncovering unstable interactions in problematic sequences.)
  - o MD Simulations revealed unstable sidechain interactions near the peptides' N-terminus for 2 unsuccessful designs. MD supported redesigns which eliminated the found interactions folded correctly in experiments.

### Undergraduate Researcher, University of Science and Technology of China

Jul 2014 – Jun 2016

Synthesized 3 microporous anion exchange membranes (crucial component of fuel cells) with exceptional high
hydroxide conductivity and chemical stability. At the time of publication, one of the membranes demonstrated
the highest conductivity at 80 °C reported in the field.

#### **PUBLICATIONS**

## 5 papers | 4 oral presentations | 10 + posters

- 1. **Guo, R.**; Sinha, N. J.; Misra, R..; Tang, Y.; Langenstein, M.; Kim, K.; Fagan, J. A.; Kloxin, C. J.; Jensen, G. V.; Pochan, D. J.; Saven, J. G., "Computational Design of Homotetrameric Peptide Bundle Variants Spanning a Wide Range of Charge States", *Biomacromolecules* 2022, 23, 4, 1652–1661
- Sinha, N. J.; Guo, R.; Misra, R.; Fagan, J.; Faraone, A.; Kloxin, C. J.; Saven, J. G.; Jensen, G. V.; Pochan, D. J., "Colloid-Like Solution Behavior of Computationally Designed Coiled Coil Bundlemers" *Journal of Colloid and Interface Science*, 606 (2022) 1974–1982
- 3. Bulos, J. A.; **Guo, R.**; Wang, Z.; Delessio, M. A.; Saven, J. G.; Dmochowski, I. J., "Design of a Superpositively Charged Enzyme: Human Carbonic Anhydrase II Variant with Ferritin Encapsulation and Immobilization" *Biochemistry*, 2021, 60, 47, 3596–3609
- 4. \*Yang, Z.; \*Guo, R.; Malpass-Evans, R.; Carta, M.; McKeown, N. B.; Guiver, M. D.; Wu, L.; Xu, T., "Highly Conductive Anion-Exchange Membranes from Microporous Tröger's Base Polymers" *Angew. Chem. Int. Ed.* 2016, 55, 11499 –11502 [\*co-first authors]
- 5. Yang, Z.; Liu, Y.; **Guo, R.**; Hou, J.; Wu, L.; Xu, T., "Highly Hydroxide Conductive Ionomers with Fullerene Functionalities" *Chem. Commun.*, 2016, 52, 2788-2791
- 6. **Guo, R.** et. al.; "Computational Design of The Electrostatic Features of Polymer-forming Peptide Bundles" *ACS Srping 2023 talk*

## LEADERSHIP, TEACHING AND OUTREACH EXPERIENCE

### Computational Lab Manager, Saven Lab

2018 - 2023

- Mentored 4 graduate students in research knowledge & tools, including protein design principles & algorithms;
   the group developed C++ code base; protein visualization & MD simulation software.
- o Administrator of the group's Red Hat Linux based high-performance computing clusters and RAID arrays.

## XSEDE Supercomputer Resource Proposal Manager, Saven Lab

2019 - 202

- o Organizer and main contributor to the group's annual proposal for the Extreme Science and Engineering Discovery Environment (XSEDE) supercomputing resource allocation.
- o Requests granted rate: 90% for 2022, 100% for 2021 & 2020, averaging ~70K CPU & 600K GPU node hrs.

#### Graduate Teaching Assistant, UPenn

Fall 2016 - Spring 2017 & Fall 2019 - Spring 2020

- o Rewarded "Commendation for Excellence in Teaching" in 2017 (8 awarded out of 32 TA candidates).
- Teaching courses include General Chemistry I, General Chemistry Lab I & II.

# Philadelphia Area Girls Enjoying Science (PAGES) Outreach

Nov 2019

Contributed to the lecture design and instructed 60+ sixth-grade girls in the outreach, offering them hands-on experiences with science and fostering awareness of scientific career opportunities for women.

### Vice Chairman of the Chemistry Student Group, School of Chemistry & Material Science, USTC 2014 – 2015

o Leader of the technical department, producing posters (~20/year) and media contents for publicity purposes.

Rui Guo