Austin Clyde

Data Science and Learning Division Argonne National Laboratory 9700 South Cass Avenue, Bldg. 240 Lemont, IL 60439 Phone: (630) 252-3738 aclyde@anl.gov

3506 N Reta Ave, FL 1 Chicago, IL 60657 Phone: (585) 406-9146 aclyde11@gmail.com austinclyde.com

RESEARCH INTERESTS

AI for science, high-performance computing, large language models, science and technology studies (STS).

EMPLOYMENT

Assistant Computational Scientist, Argonne National Laboratory2022 to presentResearch Assistant, Argonne National Laboratory2019 to 2022Desk Developer, Cryptocurrencies, XR Trading2018 to 2019

EDUCATION

University of Chicago

Ph.D. Computer Science expected December 2022

Thesis: Artificial Intelligence and High-Performance Computing for Accelerating Structure-Based Drug Discovery

M.S. Computer Science June 2019

B.A. Mathematics June 2019

FELLOWSHIPS

Visiting Research Fellow, Science, Technology, and Society, Harvard Kennedy School

2021 to 2022

AWARDS AND RECOGNITION

• Pozen Family Center for Human Rights Graduate Lectureship '22

The Pozen Center awards up to three lectureships per academic year to advanced doctoral students, each of whom teaches one undergraduate Human Rights course of their design.

- Special recognition in Department of Energy Secretary's Honor Award for COVID-19 Research '21.

 Annual award recognizing DOE employees and contractors for their service and contributions to the Department's mission and the benefit of the Nation.
- ACM Gordon Bell Special Prize for High Performance Computing-Based COVID-19 Research '20 and finalist '21.

Gordon Bell Prize, commonly referred to as the 'Nobel Prize' of supercomputing, is awarded each year to recognize outstanding achievement in high-performance computing.

• Impact Argonne Award for Discovery '20 and for Innovation '20.

Internal Argonne Award for the first sighting of new knowledge which appears to lead the way to finding more knowledge (development of drug screening pipeline), and for the initial development of a new concept (reinforcement learning for drug discovery).

GRANTS

Active Grants (PI)

9/22 – 5/24 Title: "Finding Ligands Targeting the Conserved RNA Binding Site of SARS-CoV-2 NSP13"

Principal Investigator. Austin Clyde

Funder: CACHE (Critical Assessment of Computational Hit-finding Experiments) Challenge 2 Total Direct Costs: \$31,500

12/21 – 10/23 Title: "Predict Hits for the WDR Domain of LRRK2"

Principal Investigator. Austin Clyde

Funder. CACHE (Critical Assessment of Computational Hit-finding Experiments) Challenge 1 Total Direct Costs: \$31,500

Active Grants (Non-PI)

1/22 – 12/25 Title: "IMPROVE: Innovative Methodologies and New Data for Predictive Oncology Model

Evaluation"

Principal Investigators: Rick Stevens Funder: National Cancer Institute Total Annual Direct Costs: \$2,400,00

Role: Key personnel, author of milestones

4/22 – 9/23 *Title*: "Autonomous Discovery for Science"

Principal Investigators: Thomas Brettin

Funder: Argonne National Laboratory Directed Research and Development

Total Annual Direct Costs: \$1,700,000 Role: Key personnel, author of milestones

6/20 – 6/23 Title: "AI-driven Data Integration, Inference and Multi-scale Modeling Approaches to Low-

dose Radiation Effects Understanding (RadBio-AI-3)"

Principal Investigators: Rick Stevens, Anuj Kapadia, Francis Alexander

Funder. Office of Science, Department of Energy

Total Annual Direct Costs: \$5,000,000, Argonne: \$2,000,000

Role: Key personnel, author of Argonne milestones

PUBLICATIONS

Under review

4. ChemoGraph: Interactive Visual Exploration of the Chemical Space Bharat Kale, Austin Clyde, et al.

- 3. Reproducibility is not Usability: A Case Study in Cancer Drug Response Models Austin Clyde et al.
- 2. The Black Box Civic Epistemology: AI Senselessness and Democratic Participation Austin Clyde
- 1. Deep Surrogate Docking: Accelerating Automated Drug Discovery with Graph Neural Networks Ryien Hosseini, Filippo Simini, Austin Clyde, et al.

In Preparation

7. Large Language Models as Databases Austin Clyde et al.

- 6. Who does AI Need to be Explainable to, and Who is Doing the Explaining? Austin Clyde.
- 5. The Hermeneutic Philosophy of Jasanoffian Co-Production Austin Clyde and Aishani Aatresh.
- 4. Epistemic Human Rights in the AI Age: Expertise, Interpretation, and Explainability Austin Clyde.
- 3. Protein-Ligand Docking Surrogate Models: A SARS-CoV-2 Benchmark for Deep Learning Accelerated Virtual Screening Austin Clyde, et al.
- 2. Uncertainty Calibrated Cancer Drug-Response Prediction Austin Clyde, et al.
- 1. Human Rights and AI for Science Austin Clyde.

Peer-reviewed Journals

9. Scaffold-Induced Molecular Graph (SIMG): Effective Graph Sampling Methods for High-Throughput Computational Drug Discovery

Austin Clyde, et al.

BMC Bioinformatics.

8. Intelligent Resolution: Integrating Cryo-EM with AI-driven Multi-resolution Simulations to Observe the SARS-CoV-2 Replication-Transcription Machinery in Action

Anda Trifan, Defne Gorgun, Zongyi Li, Alexander Brace, Maxim Zvyagin, Heng Ma, **Austin Clyde**, et al. The International Journal of High Performance Computing 2022; Finalist for Gordon Bell Special Prize for HPC-Based COVID-19 Research '21.

7. #COVIDisAirborne: AI-Enabled Multiscale Computational Microscopy of Delta SARS-CoV-2 in a Respiratory Aerosol

Abigail Dommer, Lorenzo Casalino, Fiona Kearns, Mia Rosenfeld, Nicholas Wauer, Surl-Hee Ahn, John Russo, Sofia Oliveira, Clare Morris, Anthony Bogetti, Anda Trifan, Alexander Brace, Terra Sztain, **Austin Clyde**, et al.

The International Journal of High Performance Computing (to appear); Finalist for Gordon Bell Special Prize for HPC-Based COVID-19 Research '21.

6. High Throughput Virtual Screening and Validation of a SARS-CoV-2 Main Protease Non-Covalent Inhibitor

Austin Clyde, et al.

Journal of Chemical Informatics 2022 (front cover).

5. Structural, electronic and electrostatic determinants for inhibitor binding to subsites S1 and S2 in SARS-CoV-2 main protease

Daniel Kneller, Hui Li, Stephanie Galanie, Gwyndalyn Phillips, Audrey Labbe, Kevin Weiss, Qiu Zhang, Mark Arnould, **Austin Clyde**, et al.

Journal of Medicinal Chemistry 2021 (appeared as supplemental cover).

4. A cross-study analysis of drug response predictions in cancer cell lines

Fangfang Xia, Jonathan Allen, Prasanna Balaprakash, Thomas Brettin, Cristina Garcia-Cardona, **Austin Clyde**, et al.

Briefings in Bioinformatics 2021.

3. Pandemic Drugs at Pandemic Speed: Accelerating COVID-19 Drug Discovery with Hybrid Machine Learning-and Physics-based Simulations on High Performance Computers Agastya Bhati, Shunzhou Wan, Dario Alfè, Austin Clyde, et al. *Interface Focus 2021*.

2. AI-driven multiscale simulations illuminate mechanisms of SARS-CoV-2 spike dynamics.

Lorenzo Casalino, Abigail C Dommer, Zied Gaieb, Emilia P Barros, Terra Sztain, Surl-Hee Ahn, Anda Trifan, Alexander Brace, Anthony T Bogetti, **Austin Clyde**, et al.

The International Journal of High-Performance Computing Applications, Gordon Bell Special Prize for HPC-Based COVID-19 Research '20.

1. Learning Curves for Drug Response Prediction in Cancer Cell Lines

Alexander Partin, Thomas Brettin, Yvonne A Evrard, Yitan Zhu, Hyunseung Yoo, Fangfang Xia, Songhao Jiang, **Austin Clyde**, et al.

BMC Bioinformatics 2021.

Conference Proceedings

8. Spatial Graph Attention and Curiosity-driven Policy for Antiviral Drug Discovery

Yulun Wu, Nicholas Choma, Andrew Chen, Mikaela Cashman, Érica T Prates, Manesh Shah, Verónica G Melesse Vergara, **Austin Clyde**, et al.

The 10th International Conference on Learning Representations (ICLR) 2022.

7. Autodock on Enamine Hit Locator Library and Drugbank

Hubertus van Dam, Martin Purschke, Dean Hidas, Oleg Tchoubar, Maksim Rakitin, Xiaohui Qu, **Austin Clyde**, et al.

IEEE Healthcare Summit Data Hackathon 2021.

6. **IMPECCABLE:** Integrated Modeling PipelinE for COVID Cure by Assessing Better LEads Aymen Al Saadi, Austin Clyde, et al.

50th International Conference on Parallel Processing (ICPP 21).

5. Stream-AI-MD: Streaming AI-driven Adaptive Molecular Simulations for Heterogeneous Computing Platforms

Alexander Brace, Michael Salim, Vishal Subbiah, Heng Ma, Murali Emani, Anda Trifan, **Austin Clyde**, et al.

Platform for Advanced Scientific Computing (PASC '21).

4. Scalable HPC and AI Infrastructure for COVID-19 Therapeutics" in the Platform for Advanced Scientific Computing

Hyungro Lee, Andre Merzky, Li Tan, Mikhail Titov, Matteo Turilli, Dario Alfe, Agastya Bhati, Alex Brace, **Austin Clyde**, et al.

Platform for Advanced Scientific Computing (PASC '21).

3. Targeting SARS-CoV-2 with AI-and HPC-enabled lead generation: a first data release Yadu Babuji, Ben Blaiszik, Tom Brettin, Kyle Chard, Ryan Chard, Austin Clyde, et al. *Platform for Advanced Scientific Computing (PASC '21)*.

2. Benchmarking Machine Learning Workloads in Structural Bioinformatics Applications Heng Ma, Austin Clyde, et al.

First International Workshop on Benchmarking Machine Learning Workloads on Emerging Hardware '20.

1. Virtual screening with deep learning using cancer cell line dose-response data Austin Clyde, et al.

American Association for Cancer Research '20.

Workshops

5. Scaffold embeddings: Learning the structure spanned by chemical fragments, scaffolds and compounds

Austin Clyde, et al.

NeurIPS Workshop on Learning Meaningful Representation of Life '21.

4. Created in our Likeness: Is Open-Source AI at Odds with Animal Rights? Austin Clyde

MANCEPT Workshop on Politics, Animals, and Technology '21.

3. Scaffold-Induced Molecular Graph (SIMG): Effective Graph Sampling Methods for High-Throughput Computational Drug Discovery

Austin Clyde, et al.

Sixth Computational Approaches to Cancer Workshop at Supercomputing '20.

2. Integrating High-Performance Simulations and Learning toward Improved Cancer Therapy Austin Clyde, et al.

Fifth Computational Approaches to Cancer Workshop at Supercomputing '19.

1. Combining molecular simulation and machine learning to INSPIRE improved cancer therapy David Wright, Adrian Devitt-Lee, Austin Clyde, et al. CompBioMed Conference '19.

Book Chapters

3. Accelerating COVID-19 Drug Discovery with High-Performance Computing Austin Clyde, to appear in High Performance Computing (HPC) for Drug Discovery and Biomedicine (ed. Heifetz).

2. Large Language Models for Science

Austin Clyde, Arvind Ramanathan, Rick Stevens

Forthcoming in AI for Science.

1. Ultrahigh Throughput Protein-Ligand Docking with Deep Learning Austin Clyde

Artificial Intelligence in Drug Design, Methods in Molecular Biology. Springer, New York, NY, 2022.

Inventions and Patents

1. RLMM: Reinforcement Learning for Molecular Mechanics

Austin Clyde, Arvind Ramanathan, Rick Stevens ANL-IN-20-123, patent filed.

Pre-prints

5. enSLMs: Genome-scale language models reveal SARS-CoV-2

Maxim Zvyagim, Alexander Brace, Kyle Hippe, Yuntian Deng, Bin Zhang, Cindy Orozco Bohorquez, **Austin Clyde**, et al. (*Gordan-Bell Prize '22 submission*)

4. Open Science Discovery of Oral Non-Covalent SARS-CoV-2 Main Protease Inhibitor Therapeutics

COVID Moonshot Consortium.

3. COVID Moonshot: Open Science Discovery of SARS-CoV-2 Main Protease Inhibitors by Combining Crowdsourcing, High-Throughput Experiments, Computational Simulations, and Machine Learning

COVID Moonshot Consortium.

- 2. Regression enrichment surfaces: a simple analysis technique for virtual drug screening models. Austin Clyde, et al.
- 1. A Systematic Approach to Featurization for Cancer Drug Sensitivity Predictions with Deep Learning

Austin Clyde, et al.

Public Opinion Pieces

- 3. Algorithmic Systems Designed to Reduce Polarization Could Hurt Democracy, Not Help It *Tech Policy Press* '22.
- 2. AI for science and global citizens

Patterns, Cell Press '22.

1. Human-in-the-Loop Systems Are No Panacea for AI Accountability

Tech Policy Press '21.

Public Comments

1. Equitable Data Engagement and Accountability (87 FR 54269)

For the White House Office of Science and Technology Policy, '22.

Short Stories

1. MvMuse

Future Humans: An Anthology (read at Harvard Kennedy School's STS Program's 20th Anniversy)

Presentations and Invited Talks

23. AI for Science: Drug Design

ALCF Intro to AI-Driven Science Training, Chicago, 2022.

22. Safeguarding Information Integrity

Harvard Data Science Institute Trust in Science Workshop, Boston, 2022.

21. Realigning Deep Learning with Virtual Ligand Screening

AI-Driven Drug Discovery Summit, Boston, 2022.

- 20. AI and Biology: A 100 Year History & AI-Driven Drug Discovery as an Emerging Technology Non-Proliferation Emerging Technology Seminar Series, Depts. of Commence and Energy, 2022.
- 19. The Boundaries of AI Ethics in American Undergraduate Computer Science Education Science and Democracy Network, Harvard Kennedy School, Boston, 2022.
- 18. Sharing What's Possible, Not Just What's Done

Democratizing HPC for Science, Platform for Advanced Scientific Computing, Basel, 2022.

17. End-to-End Computational Drug Design for COVID-19: From Screening to Series and Back Again

CompBioMed2 Workshop, Rome, 2022.

16. Automating Biological Discoveries at Scale

ISC High Performance, Hamburg, 2022

15. The Demos and the Expert: On Techniques of Self-Rule

Night of Ideas, French Embassy and Harvard University, 2022.

- 14. Throwing Away the Mirror: Why Algorithmic Disgorgement is an Epistemic Rights Violation A Right to Truth? Information, Communication, and Democracy in the 21st Century, HKS, 2022.
- 13. **How Does AI Change the Ethics of Synthetic Biology?**Convergence of SynBio and Data Science: *Prospects, Promise and Perils*, Northwestern University, 2022.
- 12. AI and Democracy

University of Chicago Department of Computer Science Student Seminar 2021.

11. The Human-in-the-Loop Learning Fallacy

Harvard Kennedy School STS Fellows 2021.

- 10. Virtual Screening with Deep Learning: Understanding Speed and Accuracy in Context Frontline Genomics Webinar 2021.
- 9. All for Drug Discovery

ATOM Consortium, June 2021.

8. **Panelist**

Intellectual Property Management: A Seminar Series, Argonne National Laboratory 2021.

- 7. Ensemble Models and Consensus Scoring for Computational Docking National Virtual Biotechnology Laboratory Seminar 2021.
- 6. **Integrating Simulations and Learning Towards Improved Cancer Therapy** CompBioMed Webinar 2020.
- 5. Tiered BFE Estimation Workflow

CompBioMed Conference 2020.

4. Accelerating Virtual Docking Screens with Deep Learning

Janssen Pharmaceuticals Seminar 2020.

3. Virtual Drug Screening with Deep Learning

NIH.AI Workshop on Applications of Machine Learning for NGS and Drug Data 2019.

2. Computational Drug Discovery at Scale

ComBioMed Kick-off meeting 2019.

1. Molecule Generation, Search, and Optimization

ATOM Technical Meeting 2019.

TEACHING

1. AI, Algorithms, and Human Rights (HMRT 23450, CMSC 10450, MAAD 13450)

Instructor, Fall 2022

Awarded Pozen Family Center for Human Rights Graduate Lectureship, University of Chicago

Cross-listed in Computer Science and Media Arts and Design.

2. Introduction to Computer Science II (C/C++) (CMSC 15200)

Instructor, Summer 2021

Department of Computer Science, University of Chicago.

3. Reinforcement Learning for Drug Discovery Practicum (MPCS 57010)

Instructor, Spring 2020

University of Chicago Master's Program in Computer Science.

4. **Computational Thinking** (CAAP)

Instructor, Summer 2022, 2021 & 2020

University of Chicago Academic Achievement Program.

5. Machine Learning in Biology and Medicine (CMSC 35440)

Graduate Teaching Assistant, Fall 2019

University of Chicago.

TRAINING, MENTORING, AND ADVISING

- 1. Max Zvyagin, post-bac student at ANL, deep learning for discovery.
- 2. Ryien Hosseini, master's student, deep learning for discovery.
- 3. Bharat Kale, Ph.D. student, chemical informatics, and visualization.

SERVICE AND OUTREACH

Outreach

- Organizer, South by Southwest Panel on the Future Faces of AI, on DEI and equity in AI.
- Organizer, panel, AI Across America with first-generation low-income students along with members of the congressional AI Caucus and technology company executives.
- Co-host of Science Technology and Society Podcast for New Books Network
- Panelist, Socioeconomic Diversity Summit, University of Chicago, 2018.
- Volunteer, Hour of Code, December 2018.
- Advisor, SEED AI.
- Member, Association for Computing Machinery (ACM).

Service

- Organizer, AI and human rights workshop series between University of Chicago's Pozen Family Center for Human Rights and Harvard Kennedy School's Program on Science, Technology, and Society.
- Organizer, minisymposium titled "Democratizing AI" at Platform for Advanced Scientific Computing (PASC '22).
- · Committee member, AI for Science and Security Initiative, U.S. National Laboratory System
- Organizer, Harvard STS Writing Group for graduate students, fellows, and visitors.

Chairing and Reviewing

- · Area chair (chemistry and small molecules), Learning Meaningful Representations of Life at NeurIPS 2022.
- Reviewer, Cell Patterns (2021, 2022)
- Reviewer, NeurIPS workshop on Machine Learning and the Physical Sciences (2021, 2022),
- Reviewer, NeurIPS dataset and benchmark track (2022),
- Reviewer, ICML (2022)