Amy Glen

glena@oregonstate.edu github.com/amykglen amykglen.github.io

EDUCATION

Oregon State University

Corvallis, OR

Ph.D. in Computer Science, GPA: 3.97

Sept. 2020-June 2024

Thesis: Computational methods for integrating and querying biomedical knowledge to facilitate translational science

University of Vermont B.A. in Biology, GPA: 3.95

Burlington, VT

2009-2012

Experience

Oregon State University

Remote

Researcher, Ramsey Lab

2019-present

- Serve as an architect and primary technical lead for our multi-institutional team within the NIH's NCATS
 Biomedical Data Translator; engineer key software components of our biomedical knowledge graph and
 reasoning engine
- Designed and implemented an automated system for building a version of the RTX-KG2 biomedical knowledge graph in which equivalent entities are resolved (<u>RTX-KG2c</u>); became the primary version of the graph and underlies the production RTX-KG2 API
- Engineered two of the <u>ARAX</u> reasoning engine's five core modules: Expand, which answers complex queries by reaching out to dynamically-selected knowledge provider ReST APIs, and Resultify, which identifies intricate result subgraphs from an answer knowledge graph
- Designed and built an in-memory, ontology-integrated graph database platform (<u>PloverDB</u>) that served as the back-end database and reasoning service for the RTX-KG2 API for four years; reduced query times to ≈8% of what they were with the prior Neo4j-based platform
- Expanded PloverDB to serve as a standalone platform (<u>Plover2.0</u>) for hosting and serving biomedical knowledge graphs via standards-compliant web APIs; sped up the <u>RTX-KG2</u> API by more than 10-times and was also adopted by the Multiomics team within Translator
- Overhauled our team's <u>node synonymization service</u> to use a semi-supervised machine learning graph clustering approach (Label Propagation); greatly reduced Type I errors in entity resolution
- Created interactive visualizations of the Biolink Model hierarchies, used daily across the consortium
- Oversee ongoing development of all of the above tools
- Mentored/guided undergraduate, post-baccalaureate, and graduate students contributing to Translator

PloverAI Remote

Co-Founder, Principal, Engineering & Product

2023-present

- Co-founded a company that provides a knowledge graph-powered biomedical literature search tool, focused on integrating clinical trial information and visualizing results
- Designed, built, and deployed the prototype software product and user interface in summer 2023
- Led initial funding efforts through strategic grant applications in 2024

DTxIntelligence Remote

Co-Founder, Principal, Engineering & Product

2018-present

 Co-founded a company that provides an evidence-focused <u>competitive intelligence data service</u> for digital therapeutics and connected devices, initially focused on diabetes

- Designed and implemented the core database and web application during the first year; wore many hats, including product manager, software engineer, and UI/UX designer
- Moved to a consulting role once the company was up and running

cvPipeline Remote

Cardiovascular Product Manager

2015 - 2018

- Managed and edited two of six sectors (Heart Valves and Hypertension) of an R&D intelligence database service for cardiovascular medical devices
- Responsible for keeping the service up to date on the latest clinical trials, regulatory approvals, patents, publications, scientific presentations, and more

VR Analytical
Analyst

Bend, OR
2014-2015

- Designed and developed a scientific knowledge database and query system to aid analytical chemists in identifying unknown compounds extracted/leached from pharmaceutical packaging
- Created a financial projections system using VBA that allowed the company to plan ahead more effectively

Publications

- [1] S. L. Goetz, A. K. Glen, and G. Glusman, "MicrobiomeKG: Bridging microbiome research and host health through knowledge graphs", bioRxiv, 2024. DOI: 10.1101/2024.10.10.617697.
- [2] G. Qin, K. Narsinh, Q. Wei, J. C. Roach, A. Joshi, S. L. Goetz, S. T. Moxon, M. H. Brush, C. Xu, Y. Yao, A. K. Glen, E. D. Morris, A. Ralevski, R. Roper, B. Belhu, Y. Zhang, I. Shmulevich, J. Hadlock, and G. Glusman, "Generating biomedical knowledge graphs from knowledge bases, registries, and multiomic data", bioRxiv, 2024. DOI: 10.1101/2024.11.14.623648.
- [3] A. K. Glen*, C. Ma*, L. Mendoza, F. Womack, E. C. Wood, M. Sinha, L. Acevedo, L. G. Kvarfordt, R. C. Peene, S. Liu, A. S. Hoffman, J. C. Roach, E. W. Deutsch, S. A. Ramsey, and D. Koslicki, "ARAX: A graph-based modular reasoning tool for translational biomedicine", *Bioinformatics*, vol. 39, no. 3, btad082, 2023, ISSN: 1367-4811. DOI: 10.1093/bioinformatics/btad082.
- [4] K. Fecho, A. E. Thessen, S. E. Baranzini, C. Bizon, J. J. Hadlock, S. Huang, R. T. Roper, N. Southall, C. Ta, P. B. Watkins, M. D. Williams, H. Xu, W. Byrd, V. Dančík, M. P. Duby, M. Dumontier, G. Glusman, N. L. Harris, E. W. Hinderer, G. Hyde, A. Johs, A. I. Su, G. Qin, Q. Zhu, and The Biomedical Data Translator Consortium, "Progress toward a universal biomedical data translator", Clinical and Translational Science, vol. 15, no. 8, pp. 1838–1847, 2022. DOI: https://doi.org/10.1111/cts.13301.
- [5] D. R. Unni, S. A. T. Moxon, M. Bada, M. Brush, R. Bruskiewich, J. H. Caufield, P. A. Clemons, V. Dancik, M. Dumontier, K. Fecho, G. Glusman, J. J. Hadlock, N. L. Harris, A. Joshi, T. Putman, G. Qin, S. A. Ramsey, K. A. Shefchek, H. Solbrig, K. Soman, A. E. Thessen, M. A. Haendel, C. Bizon, C. J. Mungall, and The Biomedical Data Translator Consortium, "Biolink model: A universal schema for knowledge graphs in clinical, biomedical, and translational science", Clinical and Translational Science, vol. 15, no. 8, pp. 1848–1855, 2022. DOI: https://doi.org/10.1111/cts.13302.
- [6] E. C. Wood*, A. K. Glen*, L. G. Kvarfordt, F. Womack, L. Acevedo, T. S. Yoon, C. Ma, V. Flores, M. Sinha, Y. Chodpathumwan, A. Termehchy, J. C. Roach, L. Mendoza, A. S. Hoffman, E. W. Deutsch, D. Koslicki, and S. A. Ramsey, "RTX-KG2: a system for building a semantically standardized knowledge graph for translational biomedicine", BMC Bioinformatics, vol. 23, p. 400, 2022. DOI: 10.1186/s12859-022-04932-3.

[7] Y. Chodpathumwan, A. Termehchy, S. A. Ramsey, A. Shrestha, A. Glen, and Z. Liu, "Structural generalizability: The case of similarity search", in *Proceedings of the 2021 International Conference on Management of Data*, ser. SIGMOD '21, Virtual Event, China: Association for Computing Machinery, 2021, pp. 326–338. DOI: 10.1145/3448016.3457316.

Presentations

- 1. Co-presenter, "Working with the Clinical Trials and Drug Approvals KPs, or: How I learned to stop worrying and love the Plover", NCATS Biomedical Data Translator Relay Conference, Washington, D.C., September 2024
- 2. Presenter, "RTX-KG2 canonicalization: A graph clustering approach to entity resolution", Every Cure Knowledge Sharing Series, virtual (hosted by Every Cure), August 2024
- 3. Poster, "ARAX: Graph-based modular reasoning for biomedicine", Outstanding Scholars Program Wine & Cheese Event, Oregon State University, Corvallis, OR, January 2023
- 4. Presenter, "RTX-KG2: A general-purpose biomedical knowledge provider", NCATS Biomedical Data Translator Knowledge Provider Info Working Group, virtual, November 2022
- 5. Poster, "PloverDB: A speedy, specialized graph database", ARCS Foundation Oregon Annual Scholars Event, Portland, OR, October 2022

Teaching

• **Teaching Assistant** at Oregon State University Analysis of Algorithms (CS325) Winter 2024, Spring 2024

• Teaching Assistant at Oregon State University Networks in Computational Biology (CS446/546)

Winter 2023, Winter 2024

Technical Skills

- Programming languages: Python, C/C++, Bash, Cypher, SQL, JavaScript, TypeScript, HTML/CSS
- Development tools: GitHub, git, Docker, AWS, Heroku, OpenAPI, uWSGI
- Web frameworks: Django, Angular, Flask
- Other: Neo4j, linux/unix, UI/UX design, data visualization, Pandas

SCHOLARSHIPS AND AWARDS

• ARCS Scholar, ARCS Foundation Oregon	2020-2023
• EECS Outstanding Scholars Fellowship, Oregon State University	2020-2022
• RELU Scholarship, Oregon State University	2018, 2019
• Joan M. Herbers Award in Biology, University of Vermont	2012
• Wasson Athletic Prize, University of Vermont	2012
• Elite 88 Award, National Collegiate Athletics Association	2011

Extracurricular Achievements

• Inducted to UVM Athletic Hall of Fame, cross-country skiing	2023
• NCAA Champion, cross-country skiing, team and individual	2012
• NCAA Division I athlete, cross-country skiing	2008-2012
• U.S. Junior National Champion, cross-country skiing	2006, 2008

^{*} Authors contributed equally to the work.