

# William Hogan

📞 510.362.1060 | ✉️ whogan@ucsd.edu | 🔗 linkedin.com/in/wphogan | 📍 San Diego, CA

## SUMMARY

Computer Science Ph.D. Candidate specializing in machine learning and natural language processing. Published novel information extraction methodologies. Interested in the intersection of LLMs, unsupervised methods, and biomedical applications. Self-starter and problem-solver with excellent communication skills. Passionate about honing expertise in the service of solving real-world problems with cutting-edge technologies.

## EDUCATION

### University of California, San Diego

*Doctor of Philosophy in Computer Science, GPA: 4.0*

2021 – March 2025 (expected)

San Diego, CA

- Ph.D. Candidate specializing in Natural Language Processing
- Advisor: Prof. Jingbo Shang

### University of California, San Diego

*Master of Science in Computer Science*

2019 – 2021

San Diego, CA

- Specialization in Machine Learning and Natural Language Processing

### University of California, Santa Cruz

*Bachelor of Science in Electrical Engineering, graduated with honors*

2003 – 2008

Santa Cruz, CA

## PROFESSIONAL EXPERIENCE

### Graduate Student Researcher

*Joan & Irwin Jacobs Center for Health Innovation, UCSD*

Jan. 2024 – Present

San Diego, CA

- Lead researcher on patient safety and infection prevention project
- Develop LLMs and automated methods to process EHR text to uncover insights on patient safety issues that were previously unknown or difficult to understand

### Research Data Scientist Graduate Intern

*Dell Technologies*

Summer 2023

Round Rock, TX

- Developed a cutting-edge text-to-SQL model to enhance user experience by enabling complex data retrieval using natural language queries

### Research Data Scientist Graduate Intern

*Dell Technologies*

Summer 2022

Round Rock, TX

- Applied state-of-the-art NLP and computer vision methods to identify fraudulent purchase orders
- Designed novel algorithm that prevented up to \$2.1M in company losses

### Graduate Student Researcher

*Center for Microbiome Innovation, UCSD*

Sept. 2019 – June 2022

San Diego, CA

- Researcher within the Artificial Intelligence for Healthy Living program funded by IBM
- Co-developed end-to-end NLP pipeline to extract information from large amounts of raw biomedical texts
- Developed high-performing models for relationship extraction, acronym resolution, and bacteria normalization
- Co-created and maintained web-based annotator tool to develop NLP models

### Co-owner, Full-stack Developer

*Design Action Collective*

2015 – 2019

Oakland, CA

- Lead developer on over 30 websites and apps while also co-managing a web development company
- Improved department-wide workflow to create cleaner, more efficient code
- Improved internal standards for code commenting, git usage, pair programming, and website accessibility

## PUBLICATIONS

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| “Open-world Semi-supervised Generalized Relation Discovery Aligned in a Real-world Setting”<br><i>Hogan, Li, Shang</i>                             | EMNLP, 2023<br><a href="#">See Publication</a> |
| “Fine-grained Contrastive Learning for Relation Extraction”<br><i>Hogan, Li, Shang</i>   | EMNLP, 2022<br><a href="#">See Publication</a> |
| “An Overview of Distant Supervision for RE with a Focus on Denoising and Pre-training Methods”<br><i>Hogan</i>                                     | arXiv<br><a href="#">See Publication</a>       |
| “Abstractified Multi-instance Learning (AMIL) for Biomedical Relation Extraction”<br><i>Hogan, Huang, Katsis, Baldwin, Kim, Baeza, Bartko, Hsu</i> | AKBC, 2021<br><a href="#">See Publication</a>  |
| “BLAR: Biomedical Local Acronym Resolver”<br><i>Hogan, Baeza, Katsis, Baldwin, Kim, Hsu</i>  | ACL, 2021<br><a href="#">See Publication</a>   |
| “Normalization of Predominant and Long-tail Bacterial Entities with a Hybrid CNN-LSTM”<br><i>Hogan, Mehta, Baeza, Katsis, Kim, Bartko, Hsu</i>     | AKBC, 2020<br><a href="#">See Publication</a>  |
| “MDAD: An Annotated Corpus for Disease-Bacterium Association” †<br><i>Huang, Hogan, Katsis, Baldwin, Kim, Baeza, Bartko, Hsu</i>                   | TBD  |
| †: work in progress  |  |

## TEACHING, SERVICE, & VOLUNTEERING

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| Teaching Assistant, <i>Introduction to Data Mining</i> at UCSD   | Winter, 2024   |
| Teaching Assistant, <i>Advanced Data-driven Text Mining</i> at UCSD                                    | Spring, 2023   |
| Program Committee Member, EMNLP  | 2022           |
| Participated as a PC member for the <i>Unsupervised and Weakly-Supervised Methods in NLP</i> workshop. |                |
| Program Committee Member, BioNLP   | 2021 – Present |
| Participated as a program committee member for the 21st BioNLP workshop, co-located with ACL, 2022.    |                |
| GradPal Mentor, UCSD   | 2021 – Present |
| Welcomed incoming students to campus and the Computer Science and Engineering program.                 |                |
| Developer Mentor, Design Action Collective   | 2016 – 2019    |
| Mentored junior web developers on coding best practices.   |                |

## AWARDS

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| First place of 38 participants in UCSD NLP Text-mining Kaggle Competition                | 2020 |
| First place in American Society of Civil Engineers National Student Robotics Competition | 2008 |
| Chancellor's Award for Outstanding Achievement   | 2008 |
| Dean's Award for Outstanding Achievement   | 2008 |

## SIDE PROJECTS

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|---|-----------|
| “Generating Position-specific Scoring Matrices for Protein Secondary Structure Prediction”  | Dec. 2020 |
| Designed and built a transformer to generate position-specific scoring matrices for protein sequences. See <a href="#">report</a> and <a href="#">repository</a> .  |           |
| “Expanding News Timeline Summarization”   | Dec. 2020 |
| Improved on existing state-of-the-art date-wise and clustering news timeline summarization (TLS) approaches, introduced more representative evaluation metrics, and expanded the available datasets to train news TLS models. See <a href="#">report</a> and <a href="#">repository</a> . |           |
| “8-state Protein Secondary Structure Prediction”  | June 2020 |
| Built a convolutional, residual, and recurrent neural network (CRRNN) that uses protein sequences and corresponding position-specific scoring matrices to predict protein secondary structures. See <a href="#">report</a> and <a href="#">repository</a> .                               |           |
| “Deep Photo Style Transfer”   | Mar. 2020 |
| Reproduced results from recent works in image style transfer using convolutional neural networks. See <a href="#">report</a> and <a href="#">repository</a> .   |           |