

# Thomas Murphy

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## EDUCATION

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### Oregon State University: B.S. Biology | March 2022

Major & Option: Biology with Genetics, GPA: 3.68

Minor: Chemistry, GPA: 3.82

Capstone Project: "Phosphorylation on residue serine-88 of DYNLL1 plays central role in functional regulation"

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## RESEARCH EXPERIENCE

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### Oregon Health & Science University: Div. of Infectious Diseases | Jun. 2024 - Present

*Research Assistant & Lab Manager*

- Helped design, execute, and troubleshoot complex methodologies to investigate *Pseudomonas aeruginosa* heteroresistance, including:
  - Serial passaging evolution experiments
  - Population analysis profiling with area under the curve calculations
  - Custom efflux pump activity assays
- Spearheaded the operational setup of a new BSL2 laboratory, writing all EHS-required compliance manuals (BSL2-lab specific, SDS, PSDS) and 15+ standard operating protocols.
- Mastered common microbiological techniques, including bacterial culturing, aseptic manipulation, and species isolation from diverse clinical samples.
- Managed all laboratory operations, including equipment maintenance, purchasing, and inventory control.
- Designed and implemented a centralized database for experimental results and laboratory management, improving team-wide data accessibility.
- Assisted adjacent studies on Gram-negative antimicrobial resistance, novel bacterium characterization, and transplant-related infections.

### Oregon State University: Department of Statistics | Jan. 2022 - March 2022

*Graduate Class Project: RNA-Seq Statistical Analysis*

- Coordinated with senior faculty to define project goals and objectives.
- Acquired, cleaned, and pre-processed raw sequencing data from the NCBI Sequence Read Archive (SRA) to prepare a dataset for downstream analysis.

- Analyzed and visualized differential gene expression data using a combination of Linux Shell scripting, the open-source Galaxy suite, and R/Python programming.
- Examined the biological and statistical backgrounds of ribonucleic acid sequencing.

## Oregon State University: Dept. of Biochem. and Biophysics | Sep. 2021 - Jan. 2022

### Senior Capstone Project

- Designed and co-led a collaborative study to investigate the effects of DYNLL1 phosphorylation on cellular regulation and transportation.
- Identified key amino acid residues as phosphorylation targets and engineered novel recombinant DYNLL1 proteins using non-canonical amino acid insertion techniques in *E. coli* TOP10ΔserB cell lines.
- Purified and verified successful phosphoserine incorporation in recombinant DYNLL1<sub>pSer-88</sub> using Immobilized Metal Affinity Chromatography (IMAC), Discontinuous Native PAGE, and Phos-Tag™ PAGE.
- Analyzed protein conformational signatures and binding affinity by performing Circular Dichroism (CD) Spectroscopy to confirm the structural impact of phosphorylation.

## PUBLICATIONS & PRESENTATIONS

Egge, **Murphy**, Strasfeld, Lewis, Lu, dePaula Baptisita, Miller, Hakki “Mechanisms and adaptive resistance in dual cefepime tazobactam heteroresistant *Pseudomonas aeruginosa* bacteremias in hematologic malignancy patients” (Manuscript in preparation; anticipated submission Dec. 2025)

Egge, **Murphy**, van der Zee, Stuckey, Strasfeld, Lewis, Lu, dePaula Baptisita, Miller, Hakki “Clinical impact and genomic features of β-lactam heteroresistance among hematologic malignancy patients with *Pseudomonas aeruginosa* bacteraemia” (Jan. 2026) Interdisciplinary Meeting on Antimicrobial Resistance and Innovation 2026, Las Vegas NV

**Thomas P. Murphy**, Lynne Strasfeld MD, William R. Miller MD, Morgan Hakki MD, Rodrigo de Baptista PhD, Stephanie L. Egge MD “Efflux activity drives cefepime heteroresistance in *Pseudomonas aeruginosa* from hematologic malignancy patients” (Nov. 2025) American Society for Microbiology Northwest Branch Meeting 2025, Portland OR, Poster and Presentation

Stephanie L Egge, **Thomas P Murphy**, William R Miller, James S Lewis, Morgan Hakki “Prevalence and clinical features of cefepime heteroresistance among bloodstream isolates of *Pseudomonas aeruginosa* in patients with hematologic malignancy” (Sept. 2024) OHSU Department of Medicine Research Retreat, Portland OR, Poster

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## TECHNICAL SKILLS

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### Bioinformatics

- **Programming & Command Line:** Unix/Linux Command Line | Python (Biopython) | R
- **Analysis Suites & Platforms:** Galaxy-suite | Geneious Prime | SnapGene | NCBI/BLAST | UCSC Genome Browser | Illumina Basespace | BV-BRC | PANTHER | Ensembl
- **Genomic & Transcriptomic Analysis:**
  - *Workflows:* read processing | genome assembly | gene prediction | read mapping | sequence alignment | motif finding | GO enrichment analysis | functional annotation.
  - *Databases:* GenBank/RefSeq (NCBI) | GEO | ENCODE | REACTOME
  - *Data Types:* BAM | SAM | FASTA/Q | BED | CSV
- **Protein Structure & Systems Modeling:**
  - *Prediction & Interaction:* AlphaFold | EMBL-EBI | STRING | DOMINE
  - *Structural Visualization:* PyMOL | UCSF Chimera
  - *Systems Modeling:* COPASI
- **Data Management & Collaboration:**
  - *Version Control:* Git/GitHub
  - *Database:* SQL

### Statistical Analysis

- **Experimental Design & Theory:** experimental & sampling design | power & confidence analysis | control considerations (negative/positive) | error management (Type I/II, FDR, FWER)
- **Statistical Modeling & Analysis:**
  - *Regression:* single/multivariate regression | Poisson regression | regression trees | variable selection (Ridge, Lasso)
  - *Classification & Testing:* T-testing | ANOVA | model selection (AIC/BIC) | multiple comparison corrections (Bonferroni, Benjamini–Hochberg)
  - *Multivariate/ML Methods:* principal component analysis (PCA) | principal component regression (PCR) | cluster analysis (Partitional, Hierarchical) | finite mixture models (FMMs)
- **Data Visualization (R & Python):** ggplot | matplotlib | seaborn | plotly

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## LEADERSHIP EXPERIENCE

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### Oregon Health & Science University: Div. of Infectious Diseases | Jun. 2024 - Present

#### *Research Assistant & Lab Manager*

- Trained and mentored a summer undergraduate intern in advanced microbiological techniques, experimental design, and data analysis.
- Help supervise their 10-week research project from protocol development to final data presentation.

### Doernbecher Children's Hospital: Assistive Technology Clinic | May. 2025 - Present

#### *Volunteer Liaison*

- Established and cultivated relationships with community partners to build broader patient outreach networks for the clinic.
- Participated in meetings with government officials, educating them on the clinic's mission and available services.
- Helped guide and support patient families by informing them of assistive technology resources and helping them navigate clinic services.

### Murphy and Co. Painting | Jun. 2016 - Jun. 2024

#### *Owner*

- Built strong relationships with clients with practiced interpersonal communication skills.
- Led and supervised painting crews, delegating tasks and managing project schedules to ensure adherence to timelines and quality standards
- Fostered a collaborative work environment and directed all client communications, resulting in high team productivity and strong client retention.

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## RELEVANT COURSEWORK

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### **Molecular Biology:**

Cell and Molecular Biology (BB 314) | General Microbiology + Laboratory (MB 302,303) | Molecular Biology Laboratory (BB 315) | Bacterial Molecular Genetics (MB 310) | Embryology and Development (Z 425) | Functional Genomics (BOT 460) | Comparative Genomics (BOT 475) | Advanced Molecular Genetics (BB 486) | Microbial Biotechnology (MB 456)

**Chemistry:**

General Chemistry Series + Laboratory (CH 231,232,233) | Organic Chemistry Series + Laboratory (CH 331,332,337) | Biochemistry Series + Advanced Biochemistry Laboratory (BB 450,451,494) | Environmental Chemistry (CH 390)

**Ecology:**

Ecology (BI 370) | Evolution (BI 445) | Invertebrate Biology + Laboratory (Z 361,362) | Population Biology (BI 483)

**Bioinformatics:**

Introduction to Sequence Analysis (BB 345) | Introduction to Genome Biology (BDS 474) | Applied Bioinformatics (BB 485)

**Mathematics:**

Differential + Integral Calculus (MTH 251, 252) | Infinite Series and Sequences (MTH 253) | Calculus through Data & Modeling: Vector Calculus (Coursera; John Hopkins University) | Linear Algebra from Elementary to Advanced Specialization (Coursera; John Hopkins University)

**Statistics:**

Introduction to Statistical Methods (ST 351) | Methods of Data Analysis Series (ST411,412,413) | Statistical Genomics (ST 592)

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**AWARDS & HONORS**

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Best Poster Award, American Society for Microbiology Northwest Branch Meeting 2025

Abstract Selected for Oral Presentation, American Society for Microbiology Northwest Branch Meeting 2025