# Adam Boudraa

**Phone:** (316)-300-7858 | **Email:** [adamboudraa@gmail.com](mailto:adamboudraa@gmail.com) | **Github:** **[github.com/adamboudruh](https://github.com/adamboudruh)**

## Education

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| **Cascadia College** | Associate's Degree in Integrated Studies DTA |
| GPA: **3.64 / 4.0** | Graduation: May 2024 |
| **University of Minnesota** | Full Stack Web Development Bootcamp |
| | Certified: April 2024 |
| **Oregon State University** | Bachelor of Science in Computer Science |
| GPA: **3.75 / 4.0** | Expected Graduation: May 2026 |

## Technical Skills

- *Languages: Java, Python, C, Javascript, TypeScript, HTML/CSS*
- *Frameworks and Tools: React, Node.js, Express.js, MongoDB, MySQL, Git, Figma, Docker, JWT, AWS Cognito, REST API, GraphQL,*

## Work Experience

- *Electronics Department Rep.        Target/Consumer Cellular        September 2022 - Present*
  - Maintain inventory accuracy and unload incoming merchandise shipments
  - Work with other team members to delegate tasks, ensure smooth operations, and foster effective collaboration
  - Sign customers up for cellular service plans, answer questions, and troubleshoot any technical issues that arise

## Projects

- *myPlanner*
  - Skills used: MERN, Apollo GraphQL, React Admin, Typescript, AWS Cognito
  - Scheduling application that allows users to create employees with customizable availabilities, make shifts for those employees
- *Fit-Connect*
  - Skills used: MERN, Apollo GraphQL, Javascript, JWT auth, Stripe, state management
  - Worked with a team of three other developers to produce a MERN stack e-commerce application displaying fitness courses from a variety of instructors
  - Users can leave reviews under courses, update and delete those reviews
- *CRUD Blog Page*
  - Skills used: MySQL, Sequelize.js, Javascript, Handlebars, REST API
  - A web-application with CRUD functionality that allows users to create accounts and post blog posts to the main feed

## Extra-Curricular Activities

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| ● Wichita East High School Game Development Club | 2019-2021 |
| ● Snohomish County Robotics Club | 2024 |
| ● Seattle Eastside Cancer Research | 2024 |

  - Group of other software developers in greater-seattle area researching how fine tuning methods for LLM models to be more accurate and applied to bioinformatics