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Taylor Smith | Scientific Software Engineer

Scientific software engineer specializing in biophysics and bioinformatics, with experience in microscopy and spectroscopy data analysis, computational modeling, and open-source scientific software. Skilled in mentoring teams, leading collaborative projects, and developing robust, reproducible pipelines for biological research.

Education

Ph.D., Biophysics  2011–2017

Example University
📍 Example City

B.S., Biochemistry  2008–2011

Example University
📍 Example City

Skills

Programming Languages

 Python  C++  TypeScript
 JavaScript  Rust

Scientific & Bioinformatics Tools

Microscopy Image Processing
Spectroscopy Data Analysis
Bioinformatics Pipelines
NumPy / SciPy / Pandas
Matplotlib / Seaborn / Plotly
Jupyter

Data Engineering & Software

Apache Airflow / Dask
Docker / Kubernetes
Cloud platforms (AWS / GCP)
CI/CD (GitHub Actions, Jenkins)
Git / GitHub

Professional Experience

Lead Scientific Software Engineer

 Example Bioinformatics Lab | Remote

 2020–Present

- Led the end-to-end development of robust, open-source Python libraries for microscopy and spectroscopy data analysis, spanning raw data ingestion, signal processing, statistical modeling, and visualization, with a strong emphasis on reproducibility, performance optimization, and maintainability to support large-scale, high-throughput experimental workflows used daily by interdisciplinary research teams.
- Lead development of open-source Python libraries for microscopy and spectroscopy analysis.
- Designed reproducible pipelines for high-throughput experimental datasets.
- Mentored a team of junior developers and research interns in computational biology tools.
- Collaborated with biologists to integrate software pipelines into laboratory workflows.
- Established coding standards, testing, and CI/CD for bioinformatics projects.

Scientific Software Engineer

 Advanced Imaging and Bioanalytics Company | Example City

 2017–2020

- Built software for real-time microscopy image acquisition and spectral analysis.
- Developed interactive dashboards for visualization of bioinformatics and experimental data.
- Contributed to open-source scientific analysis projects, including bioimaging libraries.
- Provided workshops and training sessions on computational tools for researchers.

Research Associate

 National Institute for Molecular Imaging | Example City

 2012–2017

- Automated processing and analysis of microscopy and spectroscopy datasets.
- Assisted in integration of computational models with experimental workflows.
- Contributed example notebooks and documentation to internal bioinformatics libraries.