

# William Kumler

PH.D. CANDIDATE

Seattle, WA

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## PROFESSIONAL SUMMARY

Ph.D. candidate in chemical oceanography focusing on marine microbial metabolomics under Dr. Anitra Ingalls. Expertise in automated and untargeted liquid chromatography mass spectrometry, with significant experience developing computational tools for mass spectrometry data analysis.

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## EDUCATION

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| <b>Ph.D. in Chemical Oceanography</b> , University of Washington                        | 2019 - 2025 |
| <b>M.Sc. in Oceanography</b> , University of Washington                                 | 2019 - 2023 |
| <b>B.A. in Marine Science</b> , University of California, Berkeley                      | 2014 - 2018 |
| <b>B.S. in Molecular and Environmental Biology</b> , University of California, Berkeley | 2014 - 2018 |

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## WORK EXPERIENCE

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| <b>Ph.D. student</b> , Ingalls Lab, University of Washington, Seattle  | 2019 - 2025 |
| <ul style="list-style-type: none"><li>Published papers on marine microbial metabolomics with a focus on untargeted methodologies</li><li>Developed existing and novel mass spectrometry tools for analysis and visualization</li><li>Fieldwork in the North Pacific Subtropical Gyre and California Current</li><li>Mentored graduate and undergraduate students in metabolomics</li></ul> |             |
| <b>Lab Manager</b> , Koehl Lab, University of California, Berkeley   | 2018 - 2019 |
| <ul style="list-style-type: none"><li>Maintained live cultures of choanoflagellates and protozoa</li><li>Managed and organized lab members, materials, and safety protocols</li><li>Assisted visiting researchers with statistical analyses</li><li>Tutored undergraduates in ImageJ and R programming</li></ul>   |             |
| <b>Lab Manager</b> , Edwards Lab, University of California, Berkeley   | 2018 - 2019 |
| <ul style="list-style-type: none"><li>Performed HPLC-MS sample preparation and analysis using Thermo Fusion Lumos ID-X</li><li>Developed and maintained a lipidomics pipeline for data handling and statistical analysis</li><li>Educated undergraduates in R programming and SLURM cluster usage</li></ul>  |             |

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## PUBLICATIONS

- **W. Kumler**, W. Qin, R.A. Lundeen, B. Barone, L.T. Carlson & A.E. Ingalls (2024). Metabolites reflect variability introduced by mesoscale eddies in the North Pacific Subtropical Gyre. *Frontiers in Marine Science*, 11:1481409. DOI:10.3389/fmars.2024.1481409
- **Kumler, W.**, Hazelton, B.J. & Ingalls, A.E. (2023). Picky with peakpicking: assessing chromatographic peak quality with simple metrics in metabolomics. *BMC Bioinformatics*, 24:404. DOI:10.1186/s12859-023-05533-4
- **Kumler, W.** & Ingalls, A.E. (2022). Tidy data neatly resolves mass-spectrometry's ragged arrays. *R Journal*, 14(3), 193-202. DOI:10.32614/RJ-2022-050
- Boysen, A.K., Durham, B.P., **Kumler, W.**, et al. (2022). Glycine betaine uptake and metabolism in marine microbial communities. *Environmental Microbiology*, 24(5), 2380-2403. DOI:10.1111/1462-2920.16020
- **Kumler, W.E.**, Jorge, J., Kim, P.M., et al. (2020). Does formation of multicellular colonies by choanoflagellates affect their susceptibility to capture by passive protozoan predators? *Journal of Eukaryotic Microbiology*, 67(5), 555-565. DOI:10.1111/jeu.12808

In review:

- E.A. Seelen, S.J. Gleich, **W. Kumler**, H.S. Anderson, X. Bian, K.M. Bjorkman, D.A. Caron, S.T. Dyhrman, S. Ferrón, Z.V. Finkel, S.T. Haley, Y. Hy, A.E. Ingalls, A.J. Irwin, D.M. Karl, K.P. Kong, D. Lowenstein, A. Salazar Estrada, E. Townsend, J.C. Tracey, K. Turk-Kubo, B.A.S. Van Mooy & S.G. John (in review with *Nature Communications*). A tale of two nutrients: how nitrogen and phosphorus differentially control marine biomass production and stoichiometry.

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## CONFERENCE PROCEEDINGS

- **W. Kumler** & A.E. Ingalls. Databases are a speedy, small, and simple solution for mass-spectrometry data storage and access. Upcoming presentation at the *2025 Annual International Conference of the Metabolomics Society*, Prague, Czechia.
- P. Louail, **W. Kumler**, P. Vangeenderhuysen, C. Brunius, M. Witting, S. Neumann, RforMassSpectrometry contributors & J. Rainer. xcms at 20 and still in Peak form: Now anchoring a complete ecosystem for metabolomics data preprocessing and analysis. Upcoming presentation at the *2025 Annual International Conference of the Metabolomics Society*, Prague, Czechia.
- **W. Kumler**, L.T. Carlson & A.E. Ingalls plus the PARAGON team. Differences in <sup>15</sup>N use by source during the PARAGON cruise in the NPSG. Presented as a poster at the *2024 Simons Collaboration on Ocean Processes and Ecology*, New York, NY.
- J. Rainer, P. Louail, A. Vicini, R. Gine, J. Badia, M. Stravs, M. Garcia-Aloy, C. Huber, L. Salzer, J. Stanstrup, N. Shahaf, C. Panse, T. Naake, **W. Kumler**, P. Vangeenderhuysen, C. Brunius, H. Hecht, S. Neumann, M. Witting, S. Gibb, & L. Gatto (2024). An Open Software Development-based Ecosystem of R Packages for Metabolomics Data Analysis. Presented as a poster at the *2024 Annual International Conference of the Metabolomics Society*, Osaka, Japan. DOI: 10.5281/zenodo.13347220.
- S. Garcia, F.X. Ferrer-González, J.S. Sacks, **W. Kumler**, L.T. Carlson, & A.E. Ingalls (2024). Characterizing the quality and quantity of metabolite production in phytoplankton cultures and the environment through endo- and exometabolomics. Presented as a poster at the Gordon Research Conference on Marine Microbes, Les Diablerets, Switzerland. Poster #8.
- Y. Wang, **W. Kumler**, I. Kern, E. Seelen, S.G. John, & A.E. Ingalls (2023). Community metabolomes respond to nutrient supply in a mesocosm study in the North Pacific Subtropical Gyre. Presented as a poster at the *2023 Ocean Sciences Meeting*, New Orleans, LA. Poster #OB14B-0682.
- **W. Kumler**, L.T. Carlson, & A.E. Ingalls plus the PARAGON team (2023). Metabolic fate of dissolved nitrogen in the NPSG. Presented as a poster at the *2023 Ocean Sciences Meeting*, New Orleans, LA. Poster #OB14B-0702
- **W. Kumler**, L.T. Carlson, & A.E. Ingalls plus the PARAGON team (2023). Metabolic fate of dissolved nitrogen during PARAGON. Presented as a poster at the *2023 Simons Collaboration on Ocean Processes and Ecology*, New York, NY.
- **W. Kumler** (2022). Unmasking the POC/Ness Monster: Depth and Mesoscale Features Drive Variability in Particulate Metabolite Profiles of the MESOSCOPE transect. Presented as a virtual poster at the *2022 Simons Collaboration on Ocean Processes and Ecology*, New York, NY.
- **W. Kumler** (2022). Depth, diel, and eddy direction: The effect of three environmental factors on metabolite composition in the North Pacific Subtropical Gyre. Presented as a virtual poster at the *2021 Simons Collaboration on Ocean Processes and Ecology*, New York, NY.
- **W. Kumler** & A.E. Ingalls (2021). RaMS: R-based Access to Mass-Spectrometry Data. Presented as a poster at the *2021 Annual Conference of the Metabolomics Society (virtual)*. Poster #298.
- **W. Kumler**, H. Fredricks, J. Ossolinski, A. Allen, K. Thamatrakoln, K. Bidle, B. Van Mooy, & B. Edwards (2019). Sign of the times: the lipid signature of a collapsing phytoplankton bloom. Presented as a poster at the *2019 Aquatic Sciences Meeting*, San Juan, Puerto Rico. Poster #411.
- **W. Kumler** & M.A.R. Koehl (2018). Evolution of multicellularity: Capture of unicellular vs colonial choanoflagellates by a protozoan predator. Presented as a poster at the *2018 Society of Integrative and Comparative Biology Annual Meeting*, San Francisco, CA. Poster #P1-115.

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## TALKS GIVEN

- **Building a robust model of peak quality for untargeted mass-spectrometry** (2023)
  - Seminar given at UW's eScience Seminar (watch here)
  - Seminar given at UW's Quantitative Science Seminar
- **Data Visualization** (2022)

- Workshop led at the Graduate Climate Conference
  - **Profiling R code and identifying performance bottlenecks** (2019)
    - Invited speaker at FSH 507 (Super-advanced R)
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## TECHNICAL PROJECTS

- **RaMS**: R package for reading and tidying mass-spectrometry data. Available on CRAN and GitHub.
  - **pylgrams**: Python language gloss of the RaMS package. Available on PyPI and GitHub.
  - **squallms**: R package for interactive multi-file annotation of chromatographic feature quality. Available on Bioconductor and GitHub.
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## AWARDS

- Achievement Rewards for College Scientists (ARCS)
  - Mary C. Landsteiner Student Award
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## REFERENCES

Available upon request.