

Joshua S. Sharp, Ph.D.
Curriculum Vitae

Triplett-Behrakis Endowed Professor of Pharmacology	P.O. Box 1848
Department of BioMolecular Sciences	University, MS 38677-1848
Associate Professor of Chemistry and Biochemistry	Phone: 662-915-1758
Research Associate Professor	Fax: 662-915-5638
Research Institute of Pharmaceutical Sciences	e-mail: jsharp@olemiss.edu
The University of Mississippi	Webpage: http://massspec.olemiss.edu

Education

Institution	Degree	Year
Oak Ridge National Laboratory and the University of Tennessee Graduate School of Genome Science and Technology Knoxville, Tennessee 37996 Dissertation Advisor: Dr. Robert L. Hettich; Dissertation entitled “Development of Hydroxyl Free Radical Chemistry for the Surface Mapping of Proteins”	Ph.D. Life Sciences, emphasis in Genome Science and Technology	2003
The University of Tennessee Department of Microbiology Knoxville, Tennessee 37996	B.S. Microbiology <i>Cum laude</i>	1999

Postdoctoral Training

Institution	Position	Dates
Laboratory of Structural Biology, National Institute of Environmental Health Sciences, National Institutes of Health Research Triangle Park, North Carolina 27709	Intramural Research Training Award (IRTA) Fellow	Nov. 2003 – April 2007

Employment and Training

October 2024 – present: Acting Associate Dean of Research, School of Pharmacy, University of Mississippi.
Responsibilities include:

- Administered the research activities of the School of Pharmacy
- Led emergency laboratory relocation efforts during a major, widespread infrastructure problem. Found relocation space and moved labs in as short as 24 hours to accommodate emergency infrastructure remediation efforts.
- Led development of a School of Pharmacy RIPS Seed Grant Program, projected to commence in 2025
- Directed strategic development of School of Pharmacy Research Core facilities
- Led School of Pharmacy participation in University of Mississippi Research Master Planning
- Directed preparation of reports and bulletins on School of Pharmacy research
- Reviewed and approved School of Pharmacy grant transmittals
- Represented the School of Pharmacy in campus research-oriented working groups for a wide variety of purposes, including programming and strategic planning
- Worked with the Vice Chancellor for Research and Economic Development on the creation and expansion of numerous campus-wide research enhancement programs

July 2022 – present: Triplett-Behrakis Endowed Professor, Division of Pharmacology, Department of BioMolecular Sciences, School of Pharmacy, University of Mississippi. Responsibilities include:

- May 15, 2020 – present: Director and Principal Investigator, Glycoscience Center of Research Excellence (GlyCORE), an NIH-funded P20 COBRE center focusing on the development of research infrastructure and faculty expertise in glycoscience at the University of Mississippi. Administered approx. \$2.23 million in funding annually. Responsible for faculty professional development, outreach for Research Cores, strategic planning, administration, budgeting and reporting to funders and external advisors. Also serve as Core Director for the Analytical and Biophysical Chemistry Research Core, leading sample intake, analysis, and reporting for several hundred analyses for external users per year. Approx. 50% total effort.
- Establishment and leadership of an independent research group focusing on the development and application of novel mass spectrometry-based techniques for the structural and conformational analysis of protein three-dimensional structure, protein-protein complexes, protein-carbohydrate and protein-ligand complexes; glycosaminoglycan analysis and sequencing; and determination of glycosaminoglycan structures of potential biomedical and pharmacological interest. Obtained department-leading extramural funding to support independent research group. Established an international reputation of research excellence for structural proteomics as well as glycosaminoglycan analysis.
- Lead instruction in undergraduate, professional and graduate programs within the School of Pharmacy focusing on biochemistry, protein and carbohydrate chemistry, and organic and biological mass spectrometry, including organization of the following courses:
 - PHCL 343 – Biochemical Foundations of Therapeutics
 - PHCL 541 – Problems in Pharmacology (Analysis of Glycosaminoglycans)
 - PHCL 669 – Physiological Chemistry
 - BMS 610 – Carbohydrates and Glycoconjugates
 - BMS 725/CHEM 716 – Mass Spectrometry: Fundamentals and Applications
- Formal and informal mentorship of postdoctoral trainees, graduate students, and undergraduate research assistants
- Formal and informal mentorship of junior faculty
- Participation in collaborative projects and grants within the department, the University of Mississippi, and other institutions worldwide
- Service to the department, the University of Mississippi, professional societies and the broader community.

September 2023 – February 2024: Visiting Scholar at the University of California – San Diego. Trained in the laboratory of Prof. Lisa Jones in the development and evaluation of instrumentation and methodology for *in vivo* radical footprinting from mammalian tissue. Position was funded by sabbatical from the University of Mississippi and NIH grant funds for a joint project with the Jones lab, and did not constitute employment by UCSD.

2021 – present: Member, Board of Directors of GenNext Technologies, Inc. Responsible for oversight of the company's officers, ensuring proper maintenance and stewardship of company resources, and determination of the company's strategic direction.

2017 – present: Chief Technology Officer of GenNext Technologies, Inc. and Chair of the Scientific Advisory Board. My responsibilities include directing the scientific program of GenNext Technologies as they develop and commercialize new solutions in protein photochemistry and protein pharmaceutical analysis; successfully bringing product lines to market and supporting customers in their use; provide scientific and grantsmanship leadership in SBIR and STTR applications; advise our engineering team in the implementation of integrated solutions for photochemistry; advising our software team on the design and implementation of instrument control and data analysis software for our product lines; forming and chairing a Scientific Advisory Board of

preeminent experts in the field to give strategic guidance to GenNext Technologies; leading scientific seminars and presentations to educate potential client base on hydroxyl radical protein footprinting; advising instrumental and service arms of the company on scientific issues in support of customer projects and sales; and advising customers on experimental design, data collection and analysis for biotherapeutic higher order structure analysis. Have successfully sold instruments and services to over twenty of the top pharmaceutical companies, government agencies and R1 universities. Participated in a leadership role in the invention, development and roll-out of three hardware product lines (Pioneer® system, the Fox® Protein Footprinting System, and the AutoFox® system) and one software package (FoxWare® software) with ongoing development of new product lines underway.

August 2019 – present: Associate Professor of Chemistry and Biochemistry, University of Mississippi. Mentored department graduate students, served on graduate student committees, served on faculty search committees, assisted with student instruction in selected classes, taught cross-listed course in mass spectrometry.

July 2019 – present: Associate Professor of Pharmacology, Department of BioMolecular Sciences, University of Mississippi.

July 2019 – present: Research Associate Professor, Research Institute of Pharmaceutical Sciences

2015 – June 2019: Assistant Professor of Pharmacology, Department of BioMolecular Sciences, University of Mississippi.

2015 – June 2019: Research Assistant Professor, Research Institute of Pharmaceutical Sciences

2015 – 2020: Adjunct Professor, Department of Chemistry, The University of Georgia

2012 – 2017: Founding Partner and Chief Executive Officer of Photochem Technologies, LLC. This start-up business was developed to design and market structural biology solutions based on new technologies for the protein pharmaceutical industry. Responsible for the development of the core technology behind the company, direction of the scientific output, contact with potential clients and strategic partners, and the overall strategic direction of the start-up company.

2013 – 2015: Associate Research Scientist at the Complex Carbohydrate Research Center of the University of Georgia; Faculty of the University of Georgia Graduate School.

2009 – 2015: Adjunct Research Scientist, Department of Chemistry, University of Georgia.

2009 – 2019: Faculty Member, Graduate School of the University of Georgia

2007 – 2013: Assistant Research Scientist at the Complex Carbohydrate Research Center at the University of Georgia. As a Research Faculty member, developed and maintained an independent research group focusing on the development and application of novel mass spectrometry-based techniques for the analysis of protein three-dimensional structure, protein-protein and protein-ligand binding, and the determination of glycosaminoglycan sequences of biomedical and pharmaceutical interest. Secured NIH funding to advance my research program, both independently as Principal Investigator on an NIGMS R01 grant, and as part of integrated interdisciplinary teams as a Co-Investigator on of a P41 grant and an SBIR grant. Directed the training and education of M.S. and Ph.D. students in the Department of Chemistry. Directed research training of undergraduate students in Chemistry. Provided collaborative support to various research efforts within the Complex Carbohydrate Research Center, the wider University of Georgia, and in the external academic and business communities. Taught segments in undergraduate and graduate coursework on Physical Biochemistry (BCMB 4110/6110 and CHEM 4110), Advanced Proteomics (BCMB 9300), Mass Spectrometry (CHEM 8810), Advanced Physical

Biochemistry (BCMB 8040), and external CCRC training coursework on characterization of complex glycans by chromatography and mass spectrometry. Disseminated results of research through lectures and presentations to various departments and organizations, both within the University of Georgia system and in the broader scientific community. Served as peer reviewer for various journals and funding agencies, both nationally and internationally. Served as organizer and chair for meetings and workshops in the scientific community.

2003 – 2007: Intramural Research Training Award (IRTA) fellow with Dr. Kenneth Tomer in the Laboratory of Structural Biology, National Institute of Environmental Health Sciences, National Institutes of Health. Developed and applied novel methods for the structural analysis of oxidative damage-induced conformational change. Independently developed and applied methods for the quantitative and qualitative structural analysis of proteins and large protein complexes by narrow- and broad-range specificity surface modification and mass spectrometry. Developed and pursued research in peptide-hydroxyl radical chemistry. Developed and applied methods for determining structure of gas-phase rearrangement ions by formation of solution-phase analogs and MSⁿ. Directly mentored an undergraduate student researcher in research leading to a publication. Provided analytical and protein chemistry expertise in various collaborative projects examining protein structure and protein-protein interactions. Independently wrote and submitted NIH grant applications under the K22 and K99/R00 funding mechanisms.

2001 – 2003: Graduate Research Assistant for Dr. Robert Hettich at Oak Ridge National Laboratory and Dr. Jeffrey Becker at the University of Tennessee. Developed a method for the analysis of protein structure in solution by peroxide-mediated hydroxyl radical protein footprinting surface modification and mass spectrometry; collaborated with computational biologists and an X-ray crystallography lab to use surface modification data to model structures of novel proteins. Supervised the work of two junior graduate students and an undergraduate student on research I initiated. Assisted in securing independent funding for the project I initiated from the Department of Energy. Assisted in the preparation of an R01 NIH grant based on my graduate research, which was funded in 2004. Assisted in the teaching of a biological mass spectrometry survey course.

2000 – 2001: Teaching Assistant for the Genome Science and Technology Program at the University of Tennessee, Knoxville. Assisted in instruction of an upper-level undergraduate biochemistry course for majors and a graduate survey course in bioinformatics and analytical biochemistry. Responsible for student discussion sessions, course web sites, demonstrations, student evaluation, course evaluation, and some lectures.

1999: Graduate Fellow in the Biotechnology Program at Rutgers University and the University of Medicine and Dentistry, New Jersey. Worked on heterologous expression of the beta-amyloid precursor protein.

Student and Trainee Supervision and Advising

Dissertation/Thesis Committee Chair or Co-Chair

University of Mississippi, Department of BioMolecular Sciences

- Zhi Selina Cheng (Ph.D. conferred 2022)
- Niloofar Abolhasani Khaje (Ph.D. conferred 2021)
- Hao Liu (Ph.D. conferred 2021)
- Mohammad Riaz (M.S. conferred 2019)
- Darrienne Martin (2021 – present)
- Sidney Stuckett (2021 – 2022; left program)
- Anter Shami (2023 – present)

University of Mississippi, Department of Chemistry and Biochemistry

- Quadrat Yusuph (2023 – present)
- Godson Orachor (2024 – present)

- Sayed Mobarak (2024 – present)

University of Georgia, Department of Chemistry

- Qi Gao (Ph.D. conferred 2017)
- Boer Xie (Ph.D. conferred 2016)
- Zixuan Li (Ph.D. conferred 2016)
- Yulun Chiu (Ph.D. conferred 2015, Institute of Bioinformatics, University of Georgia)
- Rongrong Huang (Ph.D. conferred 2014)
- Caroline Watson (Ph.D. conferred 2012)
- Jessica Saladino (M.S. conferred, 2011)
- ViLinh Tran (M.S. conferred, 2010)
- David Fischler (2012 – 2015; transferred when I left University of Georgia)
- Tong Zhang (2014 – 2015; transferred when I left University of Georgia)

Dissertation Committee Member

University of Mississippi

- Ousman Boye, BioMolecular Sciences (2024 – present)
- Marjan Farsi, Biology (2024 – present)
- Boluwatife Olubusoye, Chemistry (2023 – present)
- Tahir Ali, BioMolecular Sciences (2022 – present)
- Marwa Farrag, BioMolecular Sciences (2022 – present)
- Christine Hamadani, Chemistry (2021 – present)
- Divyansh Prakash, Chemistry (2020 – present)
- Mohammed Ahmed, BioMolecular Sciences (Ph.D. conferred, 2024)
- Matthew Saucier, Chemistry (Ph.D. conferred, 2024)
- Hoda Ahmed, BioMolecular Sciences (Ph.D. conferred, 2024)
- Maali Alshammari, BioMolecular Sciences (Ph.D. conferred, 2022)
- Hanan Albataineh, BioMolecular Sciences (Ph.D. conferred, 2022)
- Alysia Gonzales, BioMolecular Sciences (M.S. conferred, 2022)
- Zhiquang Gao, Chemistry (Ph.D. conferred, 2022)
- William Vignovich, BioMolecular Sciences (M.S. conferred, 2020)
- Disha Prabhu, BioMolecular Sciences (2017 – 2020)
- Taisen Hao, BioMolecular Sciences (Ph.D. conferred 2017)

University of Mississippi Medical Center

- Poonam Sharma, Microbiology and Immunology (Ph.D. conferred 2023)

University of Georgia

- Morgan Stickney, Department of Chemistry (Ph.D. conferred 2020)
- David Fischler, Department of Chemistry (Ph.D. conferred 2019)
- Jiana Duan, Department of Chemistry (Ph.D. conferred 2018)
- Majors Badgett, Department of Chemistry (Ph.D. conferred 2017)
- Emily Betchy, Department of Chemistry (Ph.D. conferred 2017)
- Joshua Driver, Department of Chemistry (Ph.D. conferred 2017)
- Isaac Agyekum, Department of Chemistry (Ph.D. conferred 2017)
- Yuejie Zhao, Department of Chemistry (Ph.D. conferred 2017)
- Amika Sood, Institute of Bioinformatics (Ph.D. conferred 2016)
- Josette Wilkes, Department of Chemistry (Ph.D. conferred 2016)
- Chengli Zong, Department of Chemistry (Ph.D. conferred 2016)

Other Post-Baccalaureate Mentorship

- Chelsea Pratt and Brad Vanderwielen (visiting application scientists from Bio-Rad for training on SEC-MALS) 2019
- Emily Hart (visiting graduate student, Department of Pharmaceutical Sciences, University of Maryland at Baltimore) 2018
- Christoph Schröder (visiting graduate student, Institute of Pharmacy, Martin Luther University, Halle-Wittenberg, Germany) 2015
- Dandan Zhou (visiting postdoctoral scholar) 2013

Postdoctoral Research Associates and Staff

- Mingming Zhao, Ph.D. (2024 – present)
- Ajay Sharma, Ph.D. (2024 – present)
- Suman Choudhary, Ph.D. (2020 – 2022)
- Anter Shami, B.S. (2019 – present)
- Surendar Tadi, Ph.D. (2017 – 2020)
- Quntao Liang, Ph.D. (2016 – 2018)
- Charles Mobley, Ph.D. (2015 – 2021)
- Sandeep Misra, Ph.D. (2015 – present)
- Xiaoyan Li, Ph.D. (2013 – 2015)

Undergraduate Research Assistants

University of Mississippi

- Madison Allenbrand (2025 – present)
- Juliette Locklar (2025 – present)
- Dalton Shults (2024 – present)
- Lyle Tobin (2021 – present); Goldwater Scholarship recipient
- Himani Patel (2021)
- Andie Udziela (2021)
- Edward Puckett (2021)
- Helen Claire McNulty (2020 – 2021)
- Elaine Morrow (2020 – 2022)
- Martin Biggs (2018 – 2021)
- Jontae Warren (2017 – 2018)
- Olivia Buquoi (2017 – 2019)
- Addison Roush (2017 – 2020); Goldwater Scholarship recipient
- Ariel Wilson (2016)
- Tam Dinh (2016)
- Sydney Watson (2016 – 2017)
- Joseph Mason (2016 – 2019)
- Lindsey Miller (2016 – 2018); Taylor Medal recipient
- Chelsea Suppinger (2015 – 2017)

University of Georgia

- David Reehl (2014 – 2015)
- Aaron Purser (2013 – 2014)

Honors and Awards

Ron Hites Award, American Society for Mass Spectrometry, 2023

Part of the GenNext Technologies team that won the coveted 2022 Grand Prize Award in the BioTools Innovator competition, from a field of 160 emerging biotechnology-focused companies

Inaugural Triplett-Behrakis Endowed Professor for the School of Pharmacy, University of Mississippi, 2022 – present

Cumberland Pharmaceuticals Inc. Faculty Research Award, 2021

The Journal of Biomolecular Techniques Outstanding Manuscript Award, 2019

University of Mississippi nominee, 2019 Blavatnik National Award for Young Scientists, Life Sciences division

New Investigator Research Award, University of Mississippi School of Pharmacy, 2018

Rho Chi Society, 2018

Intramural Research Training Award Recipient for the National Institute of Environmental Health Sciences, National Institutes of Health.

Delegate for Oak Ridge National Laboratory to the 53rd Annual Meeting of Nobel Laureates in Lindau, Germany.

Service to Scientific Community

2024: Reviewer, National Institutes of Health Small Business Innovation Research Topic 023 Phase II Panel

2024 – present: Advisory Committee member for NSF Biofoundries: Glycoscience Resources Education and Training (BIO:GREAT) center at the Complex Carbohydrate Research Center, University of Georgia

2024: Reviewer, National Institute of General Medical Sciences study section 2024/10 ZGM1 RCB-X (C1) COBRE Phase 1 Panel

2024: Reviewer, Swiss National Science Foundation

2024: Reviewer, Czech Science Foundation grant program

2023 – 2024: Reviewer, Society for Glycobiology Annual Meeting Poster Awards

2023: Meeting Chair and Chair of Organizing Committee, Mid-South Glycoscience Meeting, June 29, 2023, Oxford MS

2023: Reviewer, National Institutes of Health Small Business Innovation Research PHS-2023-1 Topic 023 Panel

2022: Meeting Chair and Chair of Organizing Committee, Mid-South Glycoscience Meeting, June 22, 2022, Oxford MS

2022-2023: Reviewer for the Netherlands Organization for Health Research and Development grant program

2019-2023: Member, MBIE College of Assessors for the Endeavour Fund, New Zealand Ministry of Business, Innovation & Employment

2022: Reviewer for National Institutes of Health, ZRG1 OBT-Y (55) Special Emphasis Panel for Basic Mechanisms in Cancer Health Disparities

2021: Meeting Chair and Chair of Organizing Committee, Mid-South Glycoscience Meeting, July 19, 2021, Oxford MS

2021: *Ad hoc* reviewer for National Science Foundation, SBIR/STTR review panel

2021-present: Periodic reviewer of tenure and promotion dossiers (details withheld for confidentiality)

2020: *Ad hoc* reviewer for National Institutes of Health, IMST (02) review panel

2019: Roundtable Facilitator for “Basics of Biophysical Tools in the Assessment of Protein HOS”, CASSS Higher Order Structure 2019, April 8 – 10, San Mateo, CA

2019: Breakout Discussion Moderator for “Repeatability and Reproducibility in Higher Order Structure Analysis”, 10th Annual Biotherapeutics Analytical Summit, March 4 – 8, Alexandria, VA

2018-2019: Member, American Society for Mass Spectrometry 2019 Research Awards Committee

2018: *Ad hoc* Reviewer for National Institutes of Health, Biomedical Technology Research Resources

2018: Scientific Reviewer, SLAC National Accelerator Laboratory

2018: Member, NSF Division of Chemistry Review Panel

2017-2018: Member, American Society for Mass Spectrometry 2018 Research Awards Committee

2017: Session Chair for “Biomolecular Dynamics: From Solution to the Gas Phase,” 1st Advancing Mass Spectrometry for Biophysics and Structural Biology Meeting, July 28th – August 1, Ann Arbor, Michigan

2015 – 2019: Referee for Mississippi Region VII Science Fair: Lower Division

2014: *Ad hoc* reviewer for National Institutes of Health, Biological Chemistry and Macromolecular Biophysics IRG

2014: *Ad hoc* Reviewer for National Science Foundation Chemistry Division

2014: Reviewer for the National Fellowships Committee of Sigma Delta Epsilon, Graduate Women in Science

2013 – 2017: Extramural Peer Reviewer, Institut National de la Santé et de la Recherche Médicale (National Institute of Health and Medical Research), France

2013 – 2016: Hydrogen-Deuterium Exchange and Covalent Labeling Interest Group Chair, American Society for Mass Spectrometry

2013: Session Chair for “Proteomics and Protein Measurements”, ISPPP 33rd Symposium on the Separation and Characterization of Biologically Important Molecules, July 17 – 19, Boston, Massachusetts.

2012: Session Chair for “Carbohydrate Analysis: New Developments”, 2012 ASMS Conference on Mass Spectrometry and Allied Topics, May 23rd, Vancouver, Canada.

2011: Reviewer for the National Center for Research Resources, National Institutes of Health

2010 – 2012: FTMS Interest Group Chair, American Society for Mass Spectrometry

Invited journal referee

- ACS Biomaterials Science & Engineering
- ACS Medicinal Chemistry Letters
- Analyst
- Analytical and Bioanalytical Chemistry
- Analytical Chemistry
- Analytical Methods
- Angewandte Chemie
- Biochemistry
- Biochimica et Biophysica Acta - Proteins and Proteomics
- Bioconjugate Chemistry
- Biotechnology and Bioengineering
- Chemical Science
- Glycobiology
- International Journal of Biological Macromolecules
- Journal of the American Chemical Society
- Journal of the American Society for Mass Spectrometry
- Journal of Biological Chemistry
- Journal of Chromatography A
- Journal of Natural Products
- Journal of Physical Chemistry
- Journal of Proteome Research
- Journal of Visualized Experiments
- Methods
- Molecular and Cellular Proteomics
- Nature Biotechnology
- Nature Communications
- Nature Communications Biology
- Nature Methods
- Organic Letters
- Scientific Reports
- The FEBS Journal
- Toxicological Sciences

Service to University

University of Mississippi

- Chemistry and Biochemistry, Faculty Search Committee Member, Analytical/Physical Chemistry, 2024 – 2025
- BioMolecular Sciences, Faculty Search Committee Member, Pharmacology, 2024 – 2025
- Faculty Mentor for Prof. Hailiang Joshua Zhu, 2023 – present
- BioMolecular Sciences, Faculty Search Committee Member, Pharmacology, 2022 – 2023

- University Research Visioning Task Force, Member, 2022 – present
- Member, Continuing Education, Lecture, and Alumni Activities Committee, 2015 – 2016; 2022 – present
- Chair, BioMolecular Sciences, Faculty Search Committee, Medicinal Chemistry/Carbohydrate Chemistry, 2022 – 2023
- Vaccine Distribution and Administration Task Force, 2020 – 2022
- GlyCORE Research Mentor for Dr. Yu-Dong Zhou, 2021 – 2023
- Faculty Mentor for Prof. Sudeshna Roy, 2019 – 2024
- Faculty Mentor for Prof. Vitor Pomin, 2019 – present
- BioMolecular Sciences, Faculty Search Committee Member, Medicinal Chemistry/Computational Chemistry, 2019 – 2020
- BioMolecular Sciences, Mentoring Task Force Head, 2019
- Chancellor's Standing Committee University Research Board: School of Pharmacy representative, 2019 – 2022
- Member, Research and Graduate Affairs Committee, 2017 – 2022
- BioMolecular Sciences, Faculty Search Committee Member, Medicinal Chemistry/Computational Chemistry, 2017 – 2018
- Vice-Chair, Research and Graduate Affairs Committee, 2016 – 2017
- BioMolecular Sciences, Department Chair Search Committee Member, 2016
- BioMolecular Sciences, Faculty Search Committee Member, Pharmacognosy/NMR, 2016
- Pharmacy School Program Interviewer, 2016 – 2021
- BioMolecular Sciences, Faculty Search Committee Member, Pharmacology, 2015 – 2016
- Early-Entry Program Interviewer, 2015 – 2021

University of Georgia

- Elected delegate for the University of Georgia University Council representing Non-Academic Ranked Faculty within the Office of the Vice-President for Research, 2009 – 2012

Grants

Ongoing:

NIH P20GM130460-01A1 (Sharp, Ross) 05/15/20 – 03/31/26 5.4 calendar

Role: PI, Core Director NIH/NIGMS \$7.5 million total (direct)

GlyCORE: Glycoscience Center of Research Excellence

The major goal of this project is the establishment of a new Phase I COBRE center focused on glycoscience as related to human health. GlyCORE will support investigators in glycoscience through direct funding of selected research projects, establish mentors for early career investigators, support the recruitment of new faculty in glycoscience, and develop local and regional meetings for investigators to discuss their work. GlyCORE will also support the establishment of three central research cores to support these investigators with cutting-edge biomedical research tools.

NIH R01 GM127267 (Sharp) 06/01/18 – 08/31/27 2 calendar

Role: PI NIH/NIGMS \$3.5 million total (direct)

Molecular Structure Determination by Mass Spectrometry and Computational Modeling

The major goals of this project are to develop and test technologies for the structural analysis of individual protein conformers from conformationally heterogeneous samples; the development and application of hydroxyl radical protein footprinting and computational modeling to determine the structure of large protein:protein complexes; and the development and application of covalent topographical labeling technology for the structural analysis of complex carbohydrates.

Pending:

P20GM130460-06A1 (Sharp) 04/01/2025 – 03/31/2030 5.4 calendar

Role: PI, Core Director NIH/NIGMS \$7.5 million total (direct)

The major goal of this project is the development of the Glycoscience Center of Research Excellence under the COBRE Phase 2 mechanism. GlyCORE will continue to support investigators in glycoscience through direct funding of research projects, aid in recruitment of new faculty in glycoscience, development of research infrastructure supporting glycoscience, and expansion of the regional glycoscience community. Phase 2 also supports the transition of our central research infrastructure to a sustainable cost recovery model.

R01GM154816 (Lindert) 07/01/2024 – 06/30/2028 1 calendar

Role: Co-I NIH/NIGMS \$292,388 (Sharp direct total)

Software Development for Protein Structure Prediction from Covalent Labeling Mass Spectrometry

The major goals of this project are to create more accurate and widely applicable computational tools for protein structure prediction from covalent labeling data.

Completed:

NIH 3P20GM130460-04S1 (Sharp, Ross) 06/23/2023 0 calendar

Role: PI NIH/NIGMS \$143,467 total (direct)

GlyCORE Supplemental Award: Computational Chemistry and Bioinformatics Research Core

This award was a supplement to the previously awarded NIH P20 award entitled “GlyCORE: Glycoscience Center of Research Excellence.” The purpose of this award was to fund purchase of an updated and expanded GPU cluster for biomolecular modeling.

NIH 3P20GM130460-04S2 (Sharp, Ross) 04/2023 – 03/2024 0 calendar

Role: PI NIH/NIGMS \$474,864 (total)

Glycoscience Center of Research Excellence (GlyCORE) – Administrative Supplement – Colby

The purpose of this supplemental award is to support team-based research to investigate how glycosylation of CTHRC1 influences its binding to target cells and its role in regulating energy metabolism. We will also track the cellular localization of internalized CTHRC1 by chemically tagging different amino acid residues to elucidate its site of action.

NIH R43GM150353 (Weinberger) 04/03/2023 – 03/29/2024 0.24 calendar

Role: Consultant NIH/NIGMS \$350,000 (total)

Liquid Chromatography Flash Oxidation (LC-Fox) Protein Footprinting System

The major goals of this project are to prototype and test a chip-based inline LC-Fox to probe the higher order structure of heterogeneous mixtures of conformers, facilitating protein structure and protein pharmaceutical research and development.

NIH 2 R01 DK042667 (Dahms) 12/01/2019 – 11/30/2024 1 calendar

Role: Co-I NIH/NIDDK \$256,060 (Sharp direct total)

Structural Analysis of the Mannose 6-Phosphate Receptors

The major goals of this project are to characterize the structure of the cation-independent mannose 6-phosphate receptor extracellular region, both alone and complexed with various biologically-important ligands. We will also probe the pH-dependent conformational changes that drive the biological activity of the receptor in the lysosome.

R15 AI137996-01A1 (Stevens) 9/1/2019 – 8/31/2022 0.48 calendar

Role: Co-I NIH/NIAID \$88,537 (direct total year 1)

Myxobacterial Predatory Antimicrobial Production in Response to Quorum Sensing Signal Interception

The major goals of this project are to probe the production of new secondary metabolite chemical entities by myxobacterial cultivars in response to the induction of myxobacterial predation by exposure to prey quorum sensing molecules.

NIH R03NS110996 (Pomin) 04/1/2019 – 03/31/2022 0 calendar
Role: Co-I NIH/NINDS \$50,000 (direct total year 1)

Disruption of CS/KSPG-Mediated Neuronal Outgrowth Inhibition

The major goals of this project are to test the ability of various sulfated polysaccharides to reverse glycosaminoglycan-mediated inhibition of neuronal outgrowth that can occur at the glial scar, and to develop new technologies for sequencing sulfation patterns of keratan sulfate

NIH 3P20GM130460-02S1 (Sharp, Ross) 08/19/2021 0 calendar
Role: PI NIH/NIGMS \$109,814 total (direct)

GlyCORE Supplemental Award: FlexMap 3D

This award was a supplement to the previously awarded NIH P20 award entitled “GlyCORE: Glycoscience Center of Research Excellence.” The purpose of this award was to fund purchase of a FlexMap 3D multiplex analyzer to support glycoscience research and serve as a platform for developing custom glycan arrays.

University of Mississippi Disaster Resilience Constellation Seed Grant: COVID-19 Track (Sharp)
05/12/20 – 06/26/20 0 calendar

Role: PI Univ. of Miss. \$2,500 total

Safety of New COVID-19 Inhibitors for Nasal Prophylaxis

The major goal of this seed grant proposal is to test the toxicity of four heparin derivatives in a mouse model via intranasal administration. Compounds are tested for nasal irritation, bleeding, discharge, weight loss, anosmia, and serum anti-coagulation.

P41 GM103390 (Moremen) 02/01/2015 – 01/31/2020 2.0 calendar
NIH/NIGMS \$128,900/year (direct, Sharp only)

Research Resource for Integrated Glycotechnology

The major goals of this research resource are to develop and disseminate new technology for studying glycosaminoglycans and the structure and function of glycosaminoglycan-protein complexes.

Role: Co-Investigator, Projects 1 and 2 (subaward)

1608685 (Sharp) 09/01/16 – 08/31/20 0.72 calendar
NSF/CHE \$339,000 (total)

Measuring Protein Topology in Conformationally Heterogeneous Mixtures by Top-Down Hydroxyl Radical Protein Footprinting

The major goal of this project is the development and evaluation of high resolution hydroxyl radical protein footprinting using electron transfer dissociation of intact proteins to allow for the structural characterization of multiple conformations of a single protein sequence co-existing in solution.

Role: Principal Investigator

R43GM126617-01 (Weinberger) 01/01/18 – 12/31/18 Contractual
NIH/NIGMS \$223,704 (total)

Foxware™, an Advanced Data Analysis Package for Hydroxyl Radical Foot-Printing Higher Order Structural Analysis

The major goal of this SBIR project are to develop a alpha-version of a user-friendly software for analyzing covalent labeling data, annotating LC-MS peaks, and disseminating results into reports on structural changes

Role: Other Key Personnel (Analytical Support, CTO)

R43GM128486-01 (Persoff) 04/01/18 – 03/31/19 Contractual
NIH/NIGMS \$150,200 (direct)

Metal-oxide Photo-Oxidation of Proteins (MoPOP) for Biopharmaceutical Higher Order Structure Analysis

The major goals of this project are to create a new and preferred means of performing flash photo-oxidation of proteins without the use of hydrogen peroxide to enable hydroxyl radical protein foot-printing as used for biopharmaceutical higher order structural analysis.

Role: Other Key Personnel (Analytical Support, CTO)

R01 GM096049A (Sharp) 07/01/2012 – 06/30/2018 1 calendar
NIH/NIGMS \$190,000/year (direct)

Improved Hydroxyl Radical Footprinting for Modeling Protein Structure

The major goals of this project are to develop methods for generating amino acid solvent accessible surface area values from hydroxyl radical protein footprinting, and to use these technologies to develop models of the interaction between gp120 of HIV and a set of broadly neutralizing antibodies

Role: Principal Investigator

R43GM125420-01 (Weinberger) 09/01/17 – 08/31/18 Contractual
NIH/NIGMS \$224,995 (total)

Tailoring the Hydroxyl Radical Footprinting Approach to Provide a Solution for the Higher Order Structural Analytical Needs of the Biopharmaceutical Industry

The major goals of this SBIR project are to develop a stand-alone prototype instrument to perform Fast Photooxidation of Proteins chemistry, creating a single vendor solution for covalent labeling photochemistry to support the biopharmaceutical industry

Role: Other Key Personnel (Analytical Support, CTO)

9P20GM104932 (McCurdy) 09/01/2015 – 08/31/2017 0 calendar
NIH/NIGMS \$25,000/year (direct, Sharp only)

Center for Research Excellence in Natural Products Neuroscience (CORE-NPN)

The major goal of this pilot project under the CORE-NPN COBRE center is to investigate the potential of hydroxyl radical small molecule footprinting as a rapid *in vitro* screening tool for identifying ligands from a pool of natural products.

Role: Pilot Project Principal Investigator

1 R43 GM109602-01 (Leach) 04/01/2014 – 03/31/2017 0 calendar
NIH/NIGMS

Automated Conformational Analysis of Pharmaceutical Proteins

The major goals of this project are to develop and commercialize an automated platform for performing high-throughput hydroxyl radical protein footprinting analyses of protein pharmaceuticals for rapid conformational equivalency testing.

Role: Co-Investigator

5 R01 AI 077569-03 (Maier) 12/01/2008 – 11/31/2013
2.0 calendar
NIH/NAIAD \$20,000/year (direct, Sharp only)

Amino Acid Repair in *Helicobacter pylori*

The major goals of this project are to determine the amino acids recognized and repaired by the methionine sulfoxide reductase protein in *H. pylori* for various proteins in the oxidative stress response pathway.

Role: Subcontractor

R01GM061268 (Prestegard) 01/01/11 – 12/31/14
0 calendar (to be funded up to 1 calendar as project demands)
NIH/NIGMS \$8,151/year (direct, Sharp only)

NMR Approaches to Membrane Associated ARF Structures

The major goals of this project are to determine the structure of ARF associated with cell membranes, using primarily state-of-the-art NMR techniques.

Role: MS support

1R43 GM100634-01A1 (Becker) 09/11/2012-04/10/2014
0.24 calendar
NIH/NIGMS \$40,000/year (direct, Sharp only)

Byologic: Software for Mass Spectrometric Protein Drug Assays

The major goals of this project are to develop commercial software that will improve our ability to characterize the detailed composition of therapeutic proteins.

Role: Co-Investigator

R01GM033225 (Prestegard) 04/01/09 – 03/31/13
0 calendar (to be funded up to 2 calendar as project demands)
NIH/NIGMS \$16,200/year (direct, Sharp only)

NMR of Cell Surface Oligosaccharides

The major goals of this project are the characterization of oligosaccharide conformation in solution, bound to protein and bound to membrane surfaces using newly developed NMR methods. Target systems include glycans on glycol-proteins that mediate protein-protein interactions.

Role: MS Support

Office of the Vice President of Research, University of Georgia Faculty Research Grant in the amount of \$12,105 awarded for January 2009-January 2011. Supports preliminary research into oxidation-induced conformational changes in Spo0F and improved tandem mass spectrometry for determining sites of oxidation.

Professional Societies

Member, American Society for Mass Spectrometry
Member, American Chemical Society (Analytical and Carbohydrate Chemistry divisions)
Member, Society for Glycobiology

Patents and Patent Applications

1. US Patent No. 10816468; PCT US 18/34682: Flash Photo-Oxidation Device and Higher Order Structural Analysis
2. US Patent No. 11181529; Intrinsic Radical Dosimetry System and Methods for Improved Hydroxyl Radical Protein Foot-Printing Analysis of Biopharmaceuticals and Biological Molecules
3. US Patent No. 20230296615A1; Radical Dosimetry Methods for In Vivo Hydroxyl Radical Protein Foot-Printing
4. US 63407959 (provisional): Identification of Residue(s) Specific Oxidation Events in Peptides Separated by Liquid Chromatography/Mass Spectrometry
5. Application Serial No. 63/502,133; US 63502133 (provisional); PCT US2024/029220: Photochemical Footprinting of Targeted Proteins from Mammalian Bodily Fluids

Clinical Trials

NCT04490239 “Intranasal Heparin Tolerability Study”. Co-PI, Responsible Party

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Venkatesh, S., Tomer, K.B., **Sharp, J.S.** (2007) Rapid identification of oxidation-induced conformational changes by kinetic analysis. *Rapid Commun Mass Spectrom* **21**, 3927-3936

Sharp, J.S. and Tomer, K.B. (2007) Analysis of the oxidative damage-induced conformational changes of apo- and holo-calmodulin by dose dependant protein oxidative surface mapping. *Biophys J* **92**, 1682-1692.

Sharp, J.S. and Tomer, K.B. (2006) Effects of Anion Proximity in Peptide Primary Sequence on the Rate and Mechanism of Leucine Oxidation. *Analytical Chemistry* **78**, 4885-4893.

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Sharp, J.S., Nelson, S., Brown, D.T., and Tomer, K.B. (2006) Structural characterization of E2 glycoprotein of Sindbis by lysine biotinylation and LC-MS/MS. *Virology* **348**, 216-223.

Sharp, J.S. and Tomer, K.B. (2005) Formation of $[b_{(n-1)+}OH + H]^+$ ions by solution-phase chemistry. *Journal of the American Society for Mass Spectrometry* **16**, 607-621.

Sharp, J.S., Guo, J., Uchiki, T., Xu, Y., Dealwis, C., and Hettich, R.L (2005). Photochemical surface mapping of C14S-Sml1p for constrained computational modeling of protein structure. *Analytical Biochemistry* **340**, 201-212.

Sharp, J.S., Becker, J.M., and Hettich, R.L. (2004) Analysis of protein solvent accessible surfaces by photochemical oxidation and mass spectrometry. *Analytical Chemistry* **76**, 672-683.

VerBerkmoes, N., **Sharp, J.S.**, and Hettich, R.L. (2004) Mass Spectrometry. In *Microbial Functional Genomics* (eds. J. Zhou, D.K. Thompson, Y. Xu, and J.M. Tiedje), Ch. 9, pg 241-283. John Wiley & Sons, Hoboken, NJ.

Sharp, J.S., Becker, J.M., and Hettich, R.L. (2002) Protein surface mapping by chemical oxidation: structural analysis by mass spectrometry. *Analytical Biochemistry* **313**, 216-225.

Koh, S., Wiles, A.M., **Sharp, J.S.**, Naider, F.R., Becker, J.M., Stacey, G. (2002) An oligopeptide transporter gene family in Arabidopsis. *Plant Physiology* **128**, 21-29.

Invited Lectures

“Radical Protein Footprinting in Stabilized Whole Blood.” **2024** CASSS Symposium on the Practical Applications of Mass Spectrometry in the Biotechnology Industry. September 11th, Bethesda, Maryland.
***Plenary Speaker**

“Radical Footprinting Applications in Structural Glycobiology.” **2023** Translational Glycomics Symposium Welcome Event. September 28th, Wauwatosa, Wisconsin. ***Invited “Rising Star” presentation.**

“Hydroxyl radical protein footprinting: a versatile tool for challenging structural problems.” **2023** Medical College of Wisconsin Department of Biochemistry. September 27, Milwaukee, Wisconsin.

“Broadening the Scope of Hydroxyl Radical Protein Footprinting Applications.” **2023** Genentech, Inc. March 15, South San Francisco, California

“Putting Together Protein Structure from Only a Handful of the Pieces: Hydroxyl Radical Protein Footprinting and Challenging Structural Problems.” **2023** Mississippi State University Department of Chemistry. February 9, Starkville, Mississippi

“LC-FOX: A mass spectrometry-based approach for measuring the higher order structure of dynamic systems.” **2022** Southeastern Regional Meeting of the American Chemical Society, Mass Spectrometry: Transcending Boundaries with Innovations in Methods & Technology symposium. Chair: Christopher Chouinard. October 19-22, San Juan, Puerto Rico

“Addressing Challenges in Structural Glycobiology Through Covalent Labeling Mass Spectrometry.” **2022** University at Albany – State University of New York Department of Chemistry. Scheduled September 27, Albany, New York

“Blinded determination of the structure of human NRG1-Ig by HR-HRPF-informed computational modeling, validated by NMR.” **2022** 2022 CASSS International Symposium on Higher Order Structure. April 5, Gaithersburg, Maryland (meeting moved to Virtual due to COVID-19)

“Determining the Structure of Neuregulin by Multi-Dose FPOP Coupled with Computational Modeling.” **2021** Southeastern Regional Meeting of the American Chemical Society, Measuring More than Mass: Innovations in Mass Spectrometry Experiments and Applications symposium. Chair: Amanda Patrick, Ph.D. November 11, Birmingham, Alabama

“Introduction to Hydroxyl Radical Protein Footprinting.” **2021** St. Jude’s Children’s Research Hospital Department of Chemical Biology and Therapeutics, October 11, Virtual

“Revolutionizing Protein Higher Order Structural Analysis with Flash Oxidation (FOX) Protein Footprinting.” **2021** AAPS eChalk Talks Series, sponsored by GenNext Technologies. June 17, Virtual.

“Probing Protein-Carbohydrate Interactions Using Mass Spectrometry.” **2021** Boston Glycobiology Discussion Group. May 12, Virtual.

“Responding to COVID-19.” **2021** Oxford Newcomers and Friends Lecture Series. Co-hosted with Prof. Lauren Bloodworth. February 10, Virtual.

“COVID-19 Vaccines.” **2021** University of Mississippi Alumni Association UPFRONT Series. February 9, Virtual.

“HRPF Data Processing and Higher Order Structural Analysis.” **2020** Protein Footprinting Seminar Series, hosted by NeoProteomics, Inc. and GenNext Technologies, Inc. December 9, Virtual.

“COVID-19 and heparan sulfate: the carbohydrate handle that SARS-CoV-2 uses to grab your cells.” **2020** Oxford Science Café, November 17, Virtual.

“Fast Photochemical Oxidation of Proteins (FPOP) HRPF.” **2020** Protein Footprinting Seminar Series, hosted by NeoProteomics, Inc. and GenNext Technologies, Inc. November 11, Virtual.

“Introduction to Protein Footprinting.” **2020** Protein Footprinting Seminar Series, hosted by NeoProteomics, Inc. and GenNext Technologies, Inc. Co-hosted with Prof. Mark Chance, Case Western Reserve University Medical Center. September 23, Virtual.

“Clarity from Complexity: Probing Protein-Glycosaminoglycan Interactions Using Mass Spectrometry.” **2019** University of Massachusetts Department of Chemistry, October 17th, Amherst, Massachusetts

“FOX: Laser-Free Sub-Millisecond Broadband Flash Oxidation for HRP.” **2019** Advancing Mass Spectrometry Meeting, July 21-25, Amherst, Massachusetts

“Compensated Hydroxyl Radical Protein Footprinting: A Flexible, Quantitative Probe of Protein Topography.” **2019** 8th International Symposium on Higher Order Structure of Protein Therapeutics (HOS 2019), April 8-10, San Mateo, California

“Qualitative and Quantitative Measurements of Protein Topography by Hydroxyl Radical Protein Footprinting.” **2019** 10th Annual Biotherapeutics Analytical Summit, March 4-8, Alexandria, Virginia

“Reproducibility and Robustness in FPOP: the Need for Community Standards for Radical Dosimetry.” **2019** ASMS Sanibel Meeting on Chemical Cross-Linking and Covalent Labeling: From Proteins to Cellular Networks, January 24-27, St. Petersburg, Florida

“Probing Protein-Glycosaminoglycan Interactions Using Mass Spectrometry: Extracting Clarity from Complexity.” **2018** Virginia Commonwealth University Institute for Structural Biology, Drug Discovery and Development, May 2nd, Richmond, Virginia

“High Resolution Hydroxyl Radical Protein Footprinting: Moving Towards Structural Determination.” **2018** University of Kansas Department of Chemistry, March 16, Lawrence, Kansas

“Quantitative Protein Topography Measurements by High Resolution Hydroxyl Radical Protein Footprinting.” **2017** Genentech, Inc. December 14th, San Francisco, California

“High Resolution Hydroxyl Radical Footprinting: Integrating Molecular Modeling with Covalent Labeling LC-MS.” **2017** 37th International Symposium and Exhibit on the Separation of Proteins, Peptides and Polynucleotides, July 19-21, Philadelphia, Pennsylvania

“Hydroxyl Radicals as Quantitative Probes for Protein Topography.” **2017** North Carolina State University Department of Chemistry, January 13, Raleigh, North Carolina

“Hydroxyl Radical Protein Footprinting: Structural Biology by the Masses.” **2015** University of Mississippi Department of Chemistry, October 8, Oxford, Mississippi

“Progress Towards Automated Sequencing of Heparin/Heparan Sulfate.” **2013** 33rd Symposium on the Separation and Characterization of Biologically Important Molecules, July 17-July 19, Boston, Massachusetts

“Hydroxyl Radical Protein Footprinting: A Mass Spectrometry-Based Measurement of Protein Conformation and Ligand Binding.” **2010** The University of Georgia College of Pharmacy, September 30, Athens, Georgia

“Structural Analysis of Mixtures of Chondroitin Sulfate.” **2009** Waters Corporation, November 3 and November 5, Beverly, Massachusetts

“Hydroxyl Radical Footprinting: A Versatile Probe of Protein Structure” **2008** University of Notre Dame Radiation Laboratory, June 23, South Bend, Indiana

“Hydroxyl Radical Footprinting: A Versatile Probe of Protein Structure and Folding.” **2007** Atlanta Area Mass Spectrometry Discussion Group, October 22, Athens, Georgia.

“Hydroxyl Radical Footprinting and Oxidation-Induced Conformational Change: Cautionary Tales for Covalent Labelers.” **2007** Hydrogen Exchange and Covalent Labeling Interest Group, 55th Annual ASMS Conference on Mass Spectrometry, June 3 - June 7, Indianapolis, Indiana

“New methods for characterizing the structural consequences of radical-mediated protein oxidation.” **2007** University of North Carolina Department of Biochemistry and Biophysics, January 16, Chapel Hill, North Carolina.

“Probing structural changes due to oxidative damage in proteins.” **2006** Vanderbilt Institute of Chemical Biology, December 20, Nashville, Tennessee

Accepted Oral Presentations at Regional, National, and International Meetings

Sharp J.S., Tobin L., Misra S.K., Luo H. and Jones L.M. **2024** Radical Protein Footprinting in Stabilized Whole Blood. 2024 International Mass Spectrometry Conference, August 17-23, Melbourne, Australia

Sharp J.S., Tobin L., Misra S.K., Luo H. and Jones, L.M. **2024** Radical Protein Footprinting in Mammalian Whole Blood. 2024 American Society for Mass Spectrometry Annual Conference, June 2-6, Anaheim, California

Sharp J.S., Misra S.K., Liu H., Xu L., and Townsend S.D. **2022** Fast Photochemical Oxidation of Carbohydrates (FPOC) for the Characterization of Carbohydrate-Protein Interactions. 2022 Mid-South Glycoscience Meeting, July 16th, Oxford, Mississippi

Cheng Z. and **Sharp J.S.** **2022** Measuring the Structural Effects of Dynamic Post-Translational Modifications by Gradient Ion Exchange LC-FOX. ASMS 2022, June 5-9, Minneapolis, Minnesota

Abolhasani Khaje N., Mobley C.K., Eletsy A., Biehn S.E., Mishra S.K. Doerksen R.J., Lindert S., Prestegard J. and **Sharp J.S.** **2021** Validated Determination of a Protein Structure by HR-HRPF Combined with Computational Modeling. ASMS 2021, October 31-Nov 5th, Philadelphia, Pennsylvania

Sharp J.S., Chea E.E., Misra S.K., Orlando R., Popov M., Egan R.W., Holman D. and Weinberger S.W. **2021** The Flash Oxidation (FOX) System: A Novel Laser-Free Fast Photochemical Oxidation Footprinting Platform. 2021 ABRF Annual Meeting, March 7-March 11, Virtual.

Misra S.K., Sood A., Soares P.A., Pomin V.H., Woods R.J. and **Sharp J.S.** **2018** Identification of the Fondaparinux Binding Site of JR-FL gp120 by High Resolution Hydroxyl Radical Protein Footprinting and Blind Computational Docking. 66th Annual ASMS Conference on Mass Spectrometry and Allied Topics, June 3-June 7, San Diego, California

Li Z., Moniz H., Wang S., Ramiah A., Zhang F., Moremen, K.W., Linhardt R.J. and **Sharp J.S.** **2015** ETD-Based High Spatial Resolution Hydroxyl Radical Protein Footprinting Reveals an Extended Robo1-Heparin Binding Interface. 63rd Annual ASMS Conference on Mass Spectrometry, May 31-June 4, St. Louis, Missouri

Huang R., Condac E., Chiu Y., Heiss C., Ishihara, M. Kailemia M.J., Amster J., Azadi P., Wang L. and **Sharp J.S.** **2013** Semi-automated sequencing of affinity-purified heparan sulfate oligosaccharides. 2013 61st Annual ASMS Conference on Mass Spectrometry, June 9-June 13, Minneapolis, Minnesota

Huang R. and **Sharp J.S.** **2012** Sequencing of Mixtures of Heparin/Heparan Sulfate Oligomers by Chemical Derivatization and LC-MS/MS. 4th Charles Warren Workshop, August 8-11, Athens, Georgia

Sharp J.S. 2012 Towards Quantitative Interpretations of Hydroxyl Radical Footprinting Data. Hydrogen Exchange and Covalent Labeling Interest Group, 60th ASMS Conference on Mass Spectrometry and Allied Topics, May 20-24, Vancouver, British Columbia, Canada

Watson C. and **Sharp J.S. 2012** Rapid Conformational Analysis of Protein Pharmaceuticals by an Abbreviated Hydroxyl Radical Footprinting Method. 60th ASMS Conference on Mass Spectrometry and Allied Topics, May 20-24, Vancouver, British Columbia, Canada

Huang R. and **Sharp J.S. 2011** Separation and Structural Characterization of Glycosaminoglycan Oligomers by LC-MSⁿ Using a Chemical Derivatization Strategy. 2011 International Symposium on the Separation of Proteins, Peptides & Polynucleotides, October 26, Alexandria, Virginia.

Huang R., Pomin V.H. and **Sharp J.S. 2011** A Chemical Derivatization Strategy for Structural Analysis of Isomeric Glycosaminoglycan Oligosaccharides Using Reverse Phase LC-MSⁿ. 59th ASMS Conference on Mass Spectrometry and Allied Topics, June 5-9, Denver, Colorado

Watson C., Wang X., **Sharp J.S.** and Prestegard J.H. **2011** Novel structural model for CCL5 (RANTES) oligomerization using hydroxyl radical footprinting. 59th ASMS Conference on Mass Spectrometry and Allied Topics, June 5-9, Denver, Colorado

Watson C., Wang X., Pomin V.H., Prestegard J.H. and **Sharp J.S. 2011** Structural Investigation of CCL5 Protein and CCL5-Glycosaminoglycan Complex Using Mass Spectrometry. 61st Pittsburgh Conference on Analytical Chemistry and Applied Spectroscopy, March 13-18, Atlanta, Georgia

Saladino J. and **Sharp J.S. 2011** Identifying and Quantifying Hydroxyl Radical Oxidation Products in Peptides and Proteins Using Improved Mass Spectrometric Techniques. 61st Pittsburgh Conference on Analytical Chemistry and Applied Spectroscopy, March 13-18, Atlanta, Georgia

Huang R. and **Sharp J.S. 2010** Structure Analysis of Derivatized Chondroitin Sulfate Oligomers with CID MS/MS. Glycomics and Glycoproteins Interest Group, 58th Annual ASMS Conference on Mass Spectrometry, May 23-May 27, Salt Lake City, Utah (given by Ms. Rongrong Huang in my absence)

Watson C. and **Sharp J.S. 2010** Application of Isotope Labeling Techniques in Protein Footprinting Studies. **2010** Hydrogen Exchange and Covalent Labeling Interest Group, 58th Annual ASMS Conference on Mass Spectrometry, May 23-May 27, Salt Lake City, Utah (given by Ms. Caroline Watson in my absence)

Saladino J. and **Sharp J.S. 2009** Improving Techniques for the Detection and Quantification of Oxidized Amino Acids in Peptides and Proteins. 10th Athens-Atlanta Mass Spectrometry Discussion Group Summer Symposium, October 6-October 7, Atlanta, Georgia

C. Yu F., Prestegard J. and **Sharp J.S. 2009** A Hydroxyl Radical Protein Footprinting Approach to the Structure of CCL5-Chondroitin Sulfate (CS) Complex. 10th Athens-Atlanta Mass Spectrometry Discussion Group Summer Symposium, October 6-October 7, Atlanta, Georgia

Watson C., Yu F., Prestegard J. and **Sharp, J.S. 2009** Hydroxyl Radical Footprinting of CCL5-Chondroitin Sulfate Complex Reveals both the Binding Interface and a Ligand-Induced Conformational Change. 57th ASMS Conference on Mass Spectrometry and Allied Topics, May 31-June 4, Philadelphia, Pennsylvania.

Watson C., O'Donnell D., Janik I., Orlando R. and **Sharp J.S. 2008** Demonstration of a Novel Methodology for Sub-Microsecond Hydroxyl Radical Protein Footprinting by Electron Beam Pulse Water Radiolysis. 59th Pittsburgh Conference on Analytical Chemistry and Applied Spectroscopy, March 1-7, New Orleans, Louisiana.

Sharp J.S. and Tomer K.B. **2004** Structural characterization of $[b_{(n-1)}+OH + H]^+$ ions by solution phase chemistry and mass spectrometry. 2004 Southeastern Regional Meeting of the American Chemical Society, November 10-13, Research Triangle Park, North Carolina.

Sharp J.S. and Tomer K.B. **2004** Solution-phase formation of $b_{(n-1)}+H_2O$ peptide ions for position-specific fragmentation at the C-terminus. 52nd Annual Conference on Mass Spectrometry and Allied Topics, May 23-27, Nashville, Tennessee.

Sharp J.S., Guo J., Gupta V., Uchiki T., Ellrott K., Becker J.M., Xu D., Xu Y., Dealwis C., and Hettich R.L. **2003** Photochemical protein surface mapping as an indicator of computational structure model accuracy. 51st Annual Conference on Mass Spectrometry and Allied Topics, June 8-12, Montreal, Canada.

Sharp J.S., Becker J.M., and Hettich R.L. **2002** Characterization of protein structure by chemical oxidation and mass spectrometry. Student Research Award Travel Grant presentation for the East Tennessee Mass Spectrometry Discussion Group, Knoxville, Tennessee

Sharp J.S., VerBerkmoes N., Asano K., Becker J., and Hettich R. **2002** Integrated Proteome Characterization: Beyond Protein Identification. Southeastern Regional Yeast Meeting, March 8-10, Gatlinburg, Tennessee

Poster Presentations at Regional, National, and International Meetings

Over 130 poster presentations at various regional, national, and international meetings.