

BIOGRAPHICAL SKETCH

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NAME		POSITION TITLE	
Hettich, Robert L.		Distinguished Research Scientist	
EDUCATION/TRAINING (<i>Begin with baccalaureate or other initial professional education, such as nursing, and include postdoctoral training.</i>)			
INSTITUTION AND LOCATION	DEGREE (if applicable)	YEAR(s)	FIELD OF STUDY
South Dakota School of Mines and Technology, Rapid City, South Dakota	B.S.	1982	Chemistry
Purdue University, West Lafayette, Indiana	Ph.D.	1986	Analytical Chemistry

A. Positions and Honors:**Positions and Employment**

1980-1982 Research Assistant, Experiment Station, South Dakota School of Mines and Technology, Rapid City, SD
 1986-2009 Research staff scientist, Oak Ridge National Laboratory
 2009 - Distinguished research scientist, Oak Ridge National Laboratory
 2000- Adjunct faculty member, University of Tennessee, Genome Sciences Graduate School
 2011- Joint faculty member, University of Tennessee, Microbiology Department
 2014- Faculty member, Bredesen Center for Interdisciplinary Research/ Graduate Education.

Other Experience and Professional Memberships

1986 - American Society for Mass Spectrometry
 1990-1996 ORISE Traveling Lecture Program
 1996-1997 East Tennessee Mass Spectrometry Discussion Group, chairman
 2000- Review member for NIH-NCI Study Sections on Quantitative Proteomics
 2001- Dissertation advisor for ORNL-UTK Ph.D. graduate students
 2002 - Review member for NIH General Medicine Study Sections
 2002 Review member for NIH Study Section of High-End Instrumentation Grants
 2004- Review member for NIH-NCI Study Section on IMAT
 2004 Chairman, NIH Study Section for New Technologies for Metabolomics
 2006 Chairman, NIH Study Section for Clinical Proteome Tech. for Cancer Research
 2008 Chairman, NIH Study Section for Clinical Proteomics Applications
 2008- Editorial Advisory Board, *Mass Spectrometry Reviews*
 2010- Editorial Advisory Board, *BMC Genomics*
 2016- Associate editor, *Microbiome*
 2014 Session organizer for International Mass Spectrometry Conference (Geneva, Switzerland)
 2019-2021 ASMS, Vice-president for arrangements

Honors

1993 Martin Marietta Energy Systems Technical Achievement Award
 2018 ORNL Outstanding graduate student mentor award
 2019 ORNL Distinguished Research Award

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

B. Peer-reviewed publications (in chronological order)

1. "Heteronuclear Transition Metal Cluster Ions in the Gas Phase: Photodissociation and Reactivity of VFe^+ ," R.L. Hettich and B.S. Freiser, J. Amer. Chem. Soc., (1985), 107, 6222.
2. "Photodissociation of $FeCH_2^+$ and $CoCH_2^+$: Determination of the Carbene, Carbyne, and Carbide Bond Strengths," R.L. Hettich and B.S. Freiser, J. Amer. Chem. Soc., (1986) 108, 2537.
3. "Gas Phase Photodissociation of Organometallic Ions: Bond Energy and Structural Determinations," R.L. Hettich, T.C. Jackson, E.M. Stanko, and B.S. Freiser, J. Amer. Chem. Soc., (1986) 108, 5086.
4. "Determination of Carbide, Carbyne, and Carbene Bond Energies by Gas Phase Photodissociation of $RhCH_2^+$, $NbCH_2^+$, and $LaCH_2^+$," R.L. Hettich and B.S. Freiser, J. Amer. Chem. Soc., (1987) 109, 3543.
5. "Spectroscopic and Thermodynamic Investigations of Transition Metal Cluster Ions in the Gas Phase: Photodissociation of MFe^+ ," R.L. Hettich and B.S. Freiser, J. Amer. Chem. Soc., (1987) 109, 3537.
6. "Gas Phase Photodissociation of Transition Metal Ion Complexes and Clusters," R.L. Hettich and B.S. Freiser, Fourier Transform Mass Spectrometry: Evolution, Innovation, and Applications, ACS Symposium Series 359, M.V. Buchanan, Editor, 1987.
7. "The Gas Phase Ion Chemistry of Methyl and Ethyl Borate," R.L. Hettich, T. Cole, and B.S. Freiser, Inter. J. Mass Spectrom. Ion Proc., (1987) 81, 203.
8. "Ligand Effects on Transition Metal Ion Reactivity: Primary and Secondary Reactions of Co^+ and Ni^+ with Alkenes," R.L. Hettich and B.S. Freiser, Organomet., (1989), 8, 2447.
9. "The Differentiation of Methyl Guanosine Isomers by Laser Ionization Fourier Transform Mass Spectrometry," R.L. Hettich, Biomed. Environ. Mass Spec., (1989) 18, 265.
10. "Characterization of Photo-induced Pyrimidine Cyclobutane Dimers by Laser Desorption Fourier Transform Mass Spectrometry" R. L. Hettich, M.V. Buchanan, and C.-h. Ho, Biomed. Environ. Mass Spec., (1990) 19, 55.
11. "Structural Investigations of Aluminum Cluster Ions, Al_n^- ($n=3-50$)," R. L. Hettich, J. Amer. Chem. Soc., (1989) 111, 8582.
12. "Investigation of UV Matrix-Assisted Laser Desorption Fourier Transform Mass Spectrometry for Peptides," R. L. Hettich and M. V. Buchanan, J. Amer. Soc. Mass Spec., (1991) 2, 22.
13. "Laser Ablation Studies of Palladium Electrolytically Loaded with Hydrogen and Deuterium," M. J. Shea, R. N. Compton, and R. L. Hettich, Phys. Rev. A, (1990) 42, 3579.
14. "Structural Characterization of Normal and Modified Oligonucleotides by Matrix-Assisted Laser Desorption Fourier Transform Mass Spectrometry," R. L. Hettich and M. V. Buchanan, J. Amer. Soc. Mass Spec. (1991), 2, 402.

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15. "Matrix-Assisted Laser Desorption Fourier Transform Mass Spectrometry for Biological Compounds," Proceedings of the NATO Workshop on Methods and Mechanisms for Producing Ions from Large Molecules, edited by K. Standing and W. Ens, Plenum Press, New York, 1991, 247-255.
16. "Doubly-Charged Negative Ions of Carbon-60," R. L. Hettich, R. N. Compton, and R. H. Ritchie, Phys. Rev. Lett. (1991), 67, 1242.
17. "Matrix-Assisted Laser Desorption Fourier Transform Mass Spectrometry for the Structural Examination of Modified Nucleic Acid Constituents," R. L. Hettich and M. V. Buchanan, Int. J. Mass Spec. Ion Proc. (1991), 111, 365.
18. "Applications of Matrix-Assisted Laser Desorption FTMS for Biomolecules," R. L. Hettich and M. V. Buchanan, Lecture Notes in Physics, 389, Laser Ablation Mechanisms and Applications, J.C. Miller and R. F. Haglund, Jr., editors, Springer-Verlag Publishers, New York, NY 1991, p. 160.
19. "Ion-Molecule Reactions of Carbon Cluster Anions," R. L. Hettich, Lecture Notes in Physics, 389, Laser Ablation Mechanisms and Applications, J.C. Miller and R. F. Haglund, Jr., editors, Springer-Verlag Publishers, New York, NY 1991, p. 280.
20. "Doubly-Charged Negative Ions of Bucky Ball - C₆₀²⁻," R. L. Hettich, R. N. Compton, and R. H. Ritchie, Lecture Notes in Physics, 389, Laser Ablation Mechanisms and Applications, J.C. Miller and R. F. Haglund, Jr., editors, Springer-Verlag Publishers, New York, NY 1991, p. 285.
21. "Applications of Mass Spectrometry to DNA Sequencing," K. B. Jacobson, H. F. Arlinghaus, C. H. Chen, G. L. Glish, R. L. Hettich, M. V. Buchanan, and S. A. McLuckey, Genetic Analysis Tech. Appl., (1991), 8, 223.
22. "Contribution to the Isolation and Characterization of the Buckminsterfullerenes," M. Diack, R.L. Hettich, R.N. Compton, and G. Guiochon, Anal. Chem., (1992) 64, 2143.
23. "Characterization and Stability of Highly Fluorinated Fullerenes," A.A. Tuinman, P. Mukherjee, J.L. Adcock, R.L. Hettich, and R.N. Compton, J. Phys. Chem., (1992) 96, 7584.
24. "Fullerenes from the Geological Environment," P.R. Buseck, S. Tsipursky, and R.L. Hettich, Science, (1992) 257, 215.
25. "Methyl Guanine Isomer Distinction by Hydrogen/Deuterium Exchange using a Fourier Transform Mass Spectrometer," B.D. Nourse and R.L. Hettich, J. Amer. Soc. Mass Spec., (1993) 4, 296-305.
26. "Rapid Extraction and Structural Characterization of Biomolecules in Agarose Gels by Laser Desorption FTMS," J. C. Dunphy, K. L. Busch, R. L. Hettich, and M. V. Buchanan, Anal. Chem., (1993) 65, 1329-1335.
27. "Characterization of Large Biomolecules by FTMS", M.V. Buchanan and R.L. Hettich, Anal. Chem., (1993) 65, 245A-259A.
28. "Characterization of Underivatized Pterins by Laser Desorption Fourier Transform Mass

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.
Spectrometry", K. Bruce Jacobson and Robert L. Hettich, Pteridines, (1993) 4, 72-75.

29. "Laser Ablation and Laser Desorption Techniques with Fourier Transform Mass Spectrometry", Robert Hettich and Changming Jin, invited book chapter for Principles and Applications of Laser Ablation, J.C. Miller, editor, Springer Series in Materials Science 28, Springer-Verlag Publishers, Berlin, (1994), p. 135-154.

30. "Matrix-Assisted Laser Desorption/Ionization Fourier Transform Mass Spectrometry of Oligodeoxyribonucleotides," E.A. Stemmler, R.L. Hettich, G.B. Hurst, and M.V. Buchanan, Rapid Comm. Mass Spec., (1993) 7, 828-826.

31. "Characterization and Identification of Radiation-Induced Products of Thymidine-3'-Monophosphate and Thymidylyl (3'->5') Thymidine by Laser Desorption Fourier Transform Mass Spectrometry," H. Yoshida and R. Hettich, Radiation Research, (1994) 139, 271-279.

32. "Synthesis and Characterization of Molybdenum Carbide Clusters: Mo_nC_{4n} ($n=1-4$)", C. Jin, R.E. Haulfer, R.L. Hettich, R.N. Compton, A.A. Puretzky, and A.V. Dem'yanenko, Science, (1994) 263, 68-71.

33. "Characterization of Naturally-Occurring and Modified Fullerenes by Fourier Transform Mass Spectrometry," R. Hettich, C. Jin, R. Compton (Oak Ridge National Laboratory) and P. Buseck and S. Tsipursky (Arizona State University), American Institute of Physics Conference Proceedings (Laser Ablation: Mechanisms and Applications-II), Vol 288, J.C. Miller and D.B. Goehagan, Editors, AIP Press, New York, (1994), p. 94-99.

34. "Laser Desorption Fourier Transform Mass Spectrometry Studies of Modified Fullerenes," C. Jin, R. Hettich, and R. Compton, American Institute of Physics Conference Proceedings (Laser Ablation: Mechanisms and Applications-II), Vol 288, J.C. Miller and D.B. Goehagan, Editors, AIP Press, New York, (1994), p. 141-147.

35. "Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry for the Structural Characterization of Modified Oligonucleotides," G. Hurst, R. Hettich, and M. Buchanan (Oak Ridge National Laboratory) and E. Stemmler (Bowdoin College, Brunswick, ME), American Institute of Physics Conference Proceedings (Laser Ablation: Mechanisms and Applications-II), Vol 288, J.C. Miller and D.B. Goehagan, Editors, AIP Press, New York, (1994), p. 519-525.

36. "Laser Ablation of Graphite in Different Buffer Gases," A.A. Puretzky, D.B. Goehagan, R.E. Haulfer, R.L. Hettich, X.-Y. Zheng, and R.N. Compton, American Institute of Physics Conference Proceedings (Laser Ablation: Mechanisms and Applications-II), Vol 288, J.C. Miller and D.B. Goehagan, Editors, AIP Press, New York, (1994), p. 365-374.

37. "Characterization of Modified Nucleic Acid Constituents by Matrix-Assisted Laser Desorption Mass Spectrometry," R. Hettich, G. Hurst, M. Buchanan, and E. Stemmler, Polycyclic Aromatic Compounds, M. Zander, editor, (1994) 6, 95-102 (peer-reviewed).

38. "The Structural Characterization of Polycyclic Aromatic Hydrocarbon Dihydrodiol Epoxide DNA

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

Adducts Using Matrix-Assisted Laser Desorption/Ionization Fourier Transform Mass Spectrometry," E.A. Stemmler, M.V. Buchanan, G.B. Hurst, and R.L. Hettich, Anal. Chem., (1994), 66, 1274-1285.

39. "Determination of the Electron Affinities of Fluorinated Fullerenes ($C_{60}F_{44,46}$, $C_{70}F_{52,54}$) by Fourier Transform Mass Spectrometry," R. Hettich, C. Jin, and R. Compton, Int. J. Mass Spec. Ion Proc (Special Issue on Fullerenes), (1994) 138, 263-274.

40. "Direct Solid Phase Hydrogenation of Fullerenes," C. Jin, R. Hettich, R. Compton, D. Joyce, J. Blencoe, and T. Burch, J. Phys. Chem., (1994), 98, 4215-4217.

41. "Attachment of Two Electrons to $C_{60}F_{46,48}$: Shape Resonances in Multiply Charged Anions," C. Jin, R. Hettich, R. Compton, A. Tuinman, Phys. Rev. Lett., (1994) 73, 2821.

42. "Ionic Properties of Hydrogenated and Fluorinated Fullerenes," R.L. Hettich, C. Jin, P.F. Britt, A.A. Tuinman, and R.N. Compton, Mat. Res. Soc. Symp. Proc. (1994) 349, 133.

43. "Thermodynamic Characterization of the Plastic Crystal and Non-Plastic Crystal Phases of C_{70} ," Y. Jin, A. Xenopoulos, J. Cheng, W. Chen, B. Wunderlich, M. Diack, C. Jin, R.L. Hettich, R.N. Compton, and G. Guiochon, Mol. Cryst. Liq. Cryst., (1994) 257, 235-250.

44. "Production and Characterization of Metallofullerene Superatoms," Z.C. Ying, C. Jin, R.L. Hettich, A.A. Puretzky, R.E. Haufler, and R.N. Compton, in *Fullerenes: Recent Advances in the Chemistry and Physics of Fullerenes and Related Materials*, edited by K.M. Kadish and R.S. Ruoff, (The Electrochemical Society, Pennington), 1995, Volume 94-24, p. 1402. (invited paper).

45. "Low Level Detection of Chemical Agent Simulants in Meat and Milk by Ion Trap Mass Spectrometry," M.V. Buchanan, R.L. Hettich, J.H. Xu, L.C. Waters, and A. Watson, J. Hazard. Mat., (1995) 42, 49.

46. "Structural Characterization of Underivatized Pteridines by Laser Desorption Fourier Transform Mass Spectrometry," R. L. Hettich and K. B. Jacobson, J. Mass Spec., (1995) 30, 872.

47. "The Analysis of Modified Oligonucleotides by Matrix-Assisted Laser Desorption/ Ionization Fourier Transform Mass Spectrometry," E.A. Stemmler, M.V. Buchanan, G.B. Hurst, and R.L. Hettich, Anal. Chem., (1995) 67, 2924.

48. "A New Interface for Combining Electrospray with Fourier Transform Ion Cyclotron Resonance Mass Spectrometry," L. Tang, R.L. Hettich, and M.V. Buchanan, Rapid Comm. Mass Spec., (1995) 9, 731.

49. "Mass Spectrometry and Small-Angle X-Ray Scattering and Studies of Gamma-Irradiated C_{60} ," R.L. Hettich, S. Henderson, R.N. Compton, and G. Bakale, J. Phys. Chem., (1996) 100 5426.

50. "Structural Determination and Ionic Properties of Endohedral Lanthanum Fullerenes," R.L. Hettich, Z. C. Ying, and R.N. Compton, *Recent Advances in the Chemistry and Physics of Fullerenes and Related Materials: Volume 3*, K.M. Kadish and R.S. Ruoff, editors, The Electrochemical Society: Pennington, N.J. (1994), p. 1457.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

51. "Instability of Fullerene Molecules at a High Temperature," Z.C. Ying, R.N. Compton, B.A. DiCamillo, R.L. Hettich, and G.A. Guiochon, J. Phys. Chem., in press.
52. "Enrichment and Characterization of a Noble-Gas Fullerene: Ar@C₆₀," B.A. DiCamillo, R.N. Compton, R.L. Hettich, A.A. Tuinman, J. Cross, Jimenez-Vazquez, M. Saunders, J. Phys Chem., (1996) 100, 9197-9201.
53. "Investigation of Oligonucleotide Fragmentation with MALDI-FTMS and Sustained Off-Resonance Irradiation (SORI)," Robert Hettich and Elizabeth Stemmler, Rapid Comm. Mass Spec. (Special ORNL issue), (1996) 10 321.
54. "Concerning Naturally-Occurring Fullerenes in Shungite," R.L. Hettich and P. R. Buseck, Carbon, (1996) 34, 685-687.
55. "Synthesis of Nitrogen-Doped Fullerenes by Laser Ablation," Z.C. Ying, R.L. Hettich, R.N. Compton, and R.E. Haufler, J. Phys. B, (1996) 29, 4935-4942.
56. "Sonochemical Synthesis of C₆₀H₂," D. Mandrus, M. Kele, R.L. Hettich, G. Guiochon, B.C. Sales, and L.A. Boatner, J. Phys. Chem., (1997) 101, 123-128.
57. "Synthesis of Doped Fullerene Clusters and Boron-Nitrogen Tubules Using Laser Ablation," Z.C. Ying, J.G. Zhu, R.N. Compton, L.F. Allard, Jr., R.L. Hettich, R.E. Haufler, invited book chapter for *Nanostructured Materials: Clusters, Composites, and Thin Films*, M. Moskovits and V.M. Shalaev, editors, (1998).
58. "Endohedral Metallofullerenes," A. Lahamer, Z.C. Ying, R.E. Haufler, R.L. Hettich, and R.N. Compton, invited book chapter for *Advances in Metal and Semiconductor Clusters, Vol. IV*, M. Duncan, editor, JAI Press, (1998), pp. 179-203.
59. "Cadiolides A and B, New Metabolites from an Ascidian of the Genus *Botryllus*," C.J. Smith, R.L. Hettich, J. Jompa, A. Tahir, M.V. Buchanan, and C. M. Ireland, J. Org. Chem., (1998), 63, 4147-4150.
60. "Investigation of the Fragmentation and Oxygen Reactivity of Endohedral Metallofullerenes M@C₆₀," R. Hettich, A. Lahamer, and R. Compton, Int. J. Mass Spec. Ion Proc., (Freiser focus issue), (1999) 182/183, 335-348.
61. "Formation and Characterization of Iron-Oligonucleotide Complexes with MALDI-FTICR," R. Hettich, J. Amer Soc. Mass Spec. (Freiser focus issue), (1999), 10, 941-949.
62. "Spacer Length Effect on the PET Fluorescent Probe for Alkali Metal Ions," H.-F. Ji, R. Dabestani, G. M. Brown, and R.L. Hettich, Photochem. Photobiol. J., (1999) 69, 513-516.
63. "Optical Sensing of Cesium Using 1,3-Alternate Calix[4]-mono- and di(anthrylmethyl)-aza-crown-6," Ji, H F, Dabestani R, Hettich R L, Brown G M, Photochemistry and Photobiology (1999) 70 882-886.
64. "Investigating the Effect of Transition Metal Ion Oxidation State on Oligodeoxyribonucleotide Binding by

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Matrix-Assisted Laser Desorption/Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometry," R.L. Hettich, Int. J. Mass Spec., (2001) 204, 55-75.

65. "Characterization of Monomeric and Dimeric Forms of Recombinant *Sml1p-histag* Protein by Electrospray-MS," T. Uchiki, R. Hettich, V. Gupta, and C. Dealwis, Anal. Biochem., (2002) 301, 35-48.
66. "Synthesis and sensing behavior of cyanoanthracene modified 1,3-alternate calix[4]benzocrown-6: a new class of Cs⁺ selective optical sensors," Ji HF, Dabestani R, Brown GM, Hettich RL, Journal of the Chemical Society-Perkin Transactions, 2 (4): 585-591 APR 2001.
67. "N-Phenyl-1,4-Phenylenediamine and Benzidine Oxidation Products Identified Using On-Line Electrochemistry/Electrospray Fourier Transform Mass Spectrometry," V. Kertesz, H. Deng, K.G. Asano, R.L. Hettich, and G.J. Van Berkel, Electroanalysis, (2002), 14, 1027-1030.
68. "Intact Protein Analysis for Site Directed Mutagenesis Overexpression Products: Plasmid Encoded R67 Dihydrofolate Reductase," N.C. VerBerkmoes, M.B. Strader, D. Smiley, E.E. Howell, G.B. Hurst, R. L. Hettich, and J.L. Stephenson, Jr., Anal. Biochem., (2002), 305, 68-81.
69. "Integrating "Top-Down" and "Bottom-Up" Mass Spectrometric Approaches for Proteomic Analysis of *Shewanella oneidensis*," N.C. VerBerkmoes, J.L. Bundy, L. Hauser, K.G. Asano, J. Razumovskaya, F. Larimer, R.L. Hettich, and J.L. Stephenson, Jr., J. Proteome Research, (2002), 1, 239-252.
70. "Protein Surface Mapping by Chemical Oxidation: Structural Analysis by Mass Spectrometry," J. Sharp, J. Becker, and R.L. Hettich, Anal. Biochem. (2003), 313, 216-225.
71. "Mass Spectrometry", Nathan VerBerkmoes, Joshua Sharp, and Robert Hettich, in *Microbial Functional Genomics*, edited by Jizhong Zhou, Dorothea Thompson, Ying Xu, and James Tiedje, John Wiley & Sons, New Jersey, 2004, pp. 241-282.
72. "Comparison of Sustained Off-Resonance Irradiation Collisionally Activated Dissociation and Multipole Storage-Assisted Dissociation for Top-Down Protein Analysis," Karin Keller, Jennifer Brodbelt, Robert Hettich, and Gary Van Berkel, J. Mass Spec. (2004), 39, 402-411.
73. "Identification of Phosphorylation Sites on the Yeast Ribonucleotide Reductase Inhibitor Sml1," T. Uchiki, R. Hettich, and C. Dealwis J. Biol. Chem. (2004), 279, 11293-11303.
74. "New Technologies for Nutrition Research," Sharon Ross, Pothur Srinivas, Andrew Clifford, Stephen Lee, Martin, Philbert, and Robert Hettich, J. Nutrition, (2004), 134, 681-685.
75. "Sml1p is a Dimer in Solution: Characterization of Denaturation and Renaturation of Recombinant Sml1p," Vibha Gupta, Cynthia Peterson, Lezlee Dice, Tomoaki Uchiki, Joseph Racca, Jun-tao Guo, Ying Xu, Robert Hettich, Xiaolan Zhao, Rodney Rothstein, and Chris Dealwis, Biochemistry, (2004), 43, 8568-8578.
76. "A Computational Method for Assessing Peptide-Identification Reliability in Tandem Mass Spectrometry Analysis with Sequest," Jane Razumovskaya, Victor Olman, Dong Xu, Ed Uberbacher, Nathan VerBerkmoes, Robert Hettich, and Ying Xu," Proteomics, (2004), 4, 961-969.
77. "Analysis of Protein Solvent Accessible Surfaces by Photochemical Oxidation and Mass Spectrometry," J. Sharp, J. Becker, and R.L. Hettich, Anal. Chem. (2004), 76, 672-683.
78. "Mass Spectrometric Approaches for Characterizing Bacterial Proteomes," Nathan VerBerkmoes, Heather Connelly, Chongle Pan, and Robert Hettich, Expert Review in Proteomics, (2004), 1, 433-447.
79. "Characterization of the 70S Ribosome from *Rhodospseudomonas palustris* Using an Integrated "Top-Down" and "Bottom-Up" Mass Spectrometric Approach," Strader M, VerBerkmoes N, Tabb D, Connelly H, Barton J, Bruce B, Pelletier D, Davison B, Hettich RL, Larimer F, Hurst G., J Proteome Res., (2004), 3, 965-978.
80. "A Graph-Theoretic Approach to Separation of b and y Ions in Tandem Mass Spectra," Bo Yan, Chongle Pan, Victor Olman, Robert Hettich, and Ying Xu, Bioinformatics, (2005), 21, 563-574.
81. "Transcriptomic and Proteomic Characterization of the Fur Modulon in the Metal-Reducing Bacterium

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- Shewanella oneidensis*,” Xiu-Feng Wan, Nathan C. VerBerkmoes, Lee Ann McCue, Dawn Stanek, Heather Connelly, Loren J. Hauser, Liyou Wu, Xueduan Liu, Tingfen Yan, Adam Leaphart, Robert L. Hettich, Jizhong Zhou, and Dorothea K. Thompson, J. Bacteriology, (2004) 186, 8385-8400.
82. “Photochemical Surface Mapping of C14S-Sml1p for Constrained Computational Modeling of Protein Structure,” Joshua Sharp, Jun-tao Guo, Tomoaki Uchiki, Ying Xu, Chris Dealwis, and Robert Hettich, Anal. Biochem., 2005, 340, 201-212.
 83. “Multipole Storage-Assisted Dissociation (MSAD) for the Characterization of Large Proteins and Protein Mixtures by ESI-FTICR-MS,” Chongle Pan and Robert Hettich, Anal. Chem., 2005, 77, 3072-3082.
 84. “DBDigger: Reorganized Proteome Database Identification Improves Flexibility and Speed,” David Tabb, Chandrasegaran Narasimhan, Michael Strader, and Robert Hettich, Anal. Chem., 2005, 77, 2464-2474.
 85. “Community proteomics identifies key activities in a natural microbial biofilm,” Rachna Ram, Nathan VerBerkmoes, Michael Thelen, Gene Tyson, Brett Baker, Robert Blake II, Manesh Shah, Robert Hettich, and Jillian Banfield,” Science, 2005, 308, 1915-1920.
 86. “MASPIC: An intensity based tandem mass spectrometry scoring scheme that improves peptide identification at high confidence,” Chandrasegaran (Chandra) Narasimhan, David L. Tabb, Nathan C. VerBerkmoes, Melissa R. Thompson, Robert L. Hettich, and Edward C. Uberbacher, Anal. Chem., (2005) 77, 7581-7593.
 87. “Global molecular and morphological effects of 24-hour chromium(VI) exposure on *Shewanella oneidensis* MR-1,” Chourey, K, Thompson MR, Morrell-Falvey J, VerBerkmoes NC, Brown SD, Shah M, Zhou JZ, Doktycz M, Hettich RL, Thompson DK Applied And Environmental Microbiology (2006), 72 6331-6344.
 88. “Determination of peptide and protein ion charge states by Fourier transformation of isotope-resolved mass spectra,” Tabb DL, Shah MB, Strader MB, Connelly HM, Hettich RL, Hurst GB J. Amer. Soc. Mass Spectrometry (2006) 17 903-915.
 89. “Molecular dynamics of the *Shewanella oneidensis* response to chromate stress,” Brown SD, Thompson MR, VerBerkmoes NC, Chourey K, Shah M, Zhou JZ, Hettich RL, Thompson DK, Molecular & Cellular Proteomics (2006) 5 1054-1071.
 90. “Determination and comparison of the baseline proteomes of the versatile microbe *Rhodopseudomonas palustris* under its major metabolic states,” VerBerkmoes NC, Shah MB, Lankford PK, Pelletier DA, Strader MB, Tabb DL, McDonald WH, Barton JW, Hurst GB, Hauser L, Davison BH, Beatty JT, Harwood CS, Tabita FR, Hettich RL, Larimer FW, Journal Of Proteome Research (2006) 5 287-298.
 91. “Proteogenomic approaches for the molecular characterization of natural microbial communities,” Banfield JF, Verberkmoes NC, Hettich RL, Thelen MP, Omics-A Journal Of Integrative Biology, (2005) 9 301-333.
 92. “Characterization of pII family (GlnK1, GlnK2, and GlnB) protein uridylylation in response to nitrogen availability for *Rhodopseudomonas palustris*,” Connelly, H. M., Pelletier, D. A., Lu, Tse-Yuan, Lankford, P. K., Hettich, Robert L. Analytical Biochemistry (2006), 357, 93-104.
 93. “ProRata: A quantitative proteomics program for accurate protein abundance ratio estimation with confidence interval evaluation,” Pan, Chongle, Kora, Guruprasad, McDonald, W. Hayes, Tabb, David L., VerBerkmoes, Nathan C., Hurst, Gregory B., Pelletier, Dale A., Samatova, Nagiza F., Hettich, Robert L. Analytical Chemistry (2006) 78, 7121-7131.
 94. “Robust estimation of peptide abundance ratios and rigorous scoring of their variability and bias in quantitative shotgun proteomics,” Pan, Chongle, Kora, Guruprasad, Tabb, David L., Pelletier, Dale A., McDonald, W. Hayes, Hurst, Gregory B., Hettich, Robert L., Samatova, Nagiza F. Analytical Chemistry (2006) 78, 7110-7120.
 95. “Detecting differential and correlated protein expression in label-free shotgun proteomics,” Zhang, Bing, VerBerkmoes, Nathan C., Langston, Michael A., Uberbacher, Edward, Hettich, Robert L., Samatova, Nagiza F. Journal Proteome Research (2006), 5, 2909-2918.
 96. “High Performance Statistical Computing with Parallel R: Applications to Biology and Climate Modeling,”

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

Samatova, N.F.; Branstetter, M.; Ganguly, A.R.; Hettich, R.L.; Khan, S.; Kora, G.; Li, J.; Ma, X.; Pan, C.; Shoshani, A.; Yoginath, S. Journal of Physics: Conference Series (2006), 46, 505-509.

97. "Strain-resolved community proteomics reveals recombining genomes of acidophilic bacteria," Lo, I; Deneff, VJ; VerBerkmoes, NC; Shah, MB; Goltsman, D; DiBartolo.; Tyson, GW; Allen, EE; Ram, RJ; Detter, Richardson, P; Thelen, MP; Hettich, RL; Banfield, JF, Nature, (2007), 446, 537-541.
98. "Dosage-dependent proteome response of *Shewanella oneidensis* MR-1 to acute chromate challenge," Thompson MR, VerBerkmoes NC, Chourey K, et al., Journal of Proteome Research, (2007), 6, 1745-1757
99. "Implications of strain- and species-level sequence divergence for community and isolate shotgun proteomic analysis," Deneff V J, Shah M B, VerBerkmoes N C, Hettich, R L, Banfield J F, Journal Of Proteome Research, (2007), 6, 3152-3161.
100. "Systematic Assessment of the Benefits and Caveats in Mining Microbial Post-Translational Modifications from Shotgun Proteomic Data: The Response of *Shewanella oneidensis* to Chromate Exposure," Thompson, M.R.; Thompson, D.K.; Hettich, R.L. Journal Of Proteome Research, (2008), 7, 648-658.
101. "Characterization of Anaerobic Catabolism of p-Coumarate in *Rhodospseudomonas palustris* by Integrating Transcriptomics and Quantitative Proteomics," Pan, C.; Oda, Y.; Lankford, P.K.; Zhang, B.; Samatova, N.F.; Pelletier, D.A.; Harwood, C.S.; Hettich, R.L. Molecular and Cellular Proteomics, (2008), 7, 938-948.
102. "Development of an Electrochemical Oxidation Method for Probing Higher Order Protein Structure with Mass Spectrometry," McClintock, C.; Kertesz, V.; Hettich, R.L. Anal. Chem. (2008), 80, 3304-3317.
103. "Cytochrome 572 is a potential iron oxidase directly purified from a natural acidophilic microbial community," Chris Jeans, Steven W. Singer, Clara S. Chan, Nathan C. VerBerkmoes, Manesh Shah, Robert L. Hettich, Jillian F. Banfield and Michael P. Thelen ISME Journal (2008), 2, 542-550.
104. "Characterization of Cytochrome 579, an unusual cytochrome isolated from an iron oxidizing microbial community," Steven W. Singer, Clara S. Chan, Adam Zemla, Nathan C. VerBerkmoes, Mona Hwang, Robert L. Hettich, Jillian F. Banfield, and Michael P. Thelen, AEM (2008), 74, 4454-4462.
105. "Community proteogenomics suggests microbial fine-tuning in enhanced biological phosphorus removal within activated sludge," Paul Wilmes, Anders F Andersson, Mark G Lefsrud, Margaret Wexler, Manesh Shah, Robert L Hettich, Philip L Bond, Nathan C VerBerkmoes, and Jillian F Banfield, ISME Journal (2008), 2, 853-864
106. "Experimental Approach for Deep Proteome Measurements from Small-Scale Microbial Biomass Samples," Thompson, M.R., Chourey, K., Froelich, J.M., Erickson, B.K., VerBerkmoes, N.C., Hettich, R.L., Analytical Chemistry (2008), 8, 9517-9525.
107. "Systems: Biology: Functional Analysis Of Natural Microbial Consortia Using Community Proteomics," VerBerkmoes, N.C., Deneff, V.J., Hettich, R.L., and Banfield, J.F. Nature Reviews Microbiology (2009), 7, 196-205.
108. "Shotgun Metaproteomics of the Human Distal Gut Microbiota," Verberkmoes, N. C., Russell, A. L., Shah, M., Godzik, A., Rosenquist, M., Halfvarson, J., Lefsrud, M.G., Apajalahti, J., Tysk, C., Hettich R.L., Jansson, J., ISME Journal, (2009), 3, 179-189.
109. "Proteomics-inferred genome typing (PIGT) demonstrates inter-population recombination as a strategy for environmental adaptation," Deneff, V. J., VerBerkmoes, N.C., Shah, M.B., Abraham, P., Lefsrud, M., Hettich, R.L., Banfield, J. F., Environmental Microbiology (2009), 11, 313-325.
110. "Comparative Temporal Proteomics of a Response Regulator (SO2426)-Deficient Strain and Wild-Type *Shewanella oneidensis* MR-1 During Chromate Transformation," Chourey, K., Thompson, M.R., Shah, M., Zhang, B., VerBerkmoes, N. C., Thompson, D.K., Hettich, R.L., J. Proteome Research (2009) 8, Special Issue: 59-71.
111. "Environmental Proteomics: a Paradigm Shift in Characterizing Microbial Activities at the Molecular Level," Martin Keller and Robert Hettich, Microbiology and Molecular Biology Reviews, (2009), 62-70.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

112. "Characterizing a model human gut microbiota composed of members of its two dominant bacterial phyla," Mahowald, Michael A.; Rey, Federico E.; Seedorf, Henning; Turnbaugh, Peter J.; Fulton, Robert S.; Wollam, Aye; Shah, Neha;; Wang, Chunyan; Magrini, Vincent; Wilson, Richard K.; Cantarel, Brandi L.; Coutinho, Pedro M.; Henrissat, Bernard; Crock, Lara W.; Russell, Alison; Verberkmoes, Nathan C.; Hettich, Robert L.; Gordon, Jeffrey I. Proc. Natl. Acad. Sci. (2009) 106, 5859-5864.
113. "Differential Accumulation of Form I RubisCO in *Rhodopseudomonas palustris* CGA010 under Photoheterotrophic Growth Conditions with Reduced Carbon Sources," Joshi, Gauri S.; Romagnoli, Simona; VerBerkmoes, Nathan C.; Hettich, Robert L.; Pelletier, Dale; Tabita, F. Robert, Journal of Bacteriology (2009) 191, 4243-4250.
114. "Community Genomic and Proteomic Analyses of Chemoautotrophic Iron-Oxidizing "*Leptospirillum rubarum*" (Group II) and "*Leptospirillum ferrodiazotrophum*" (Group III) Bacteria in Acid Mine Drainage Biofilms," Goltsman, Daniela S. Aliaga; Deneff, Vincent J.; Singer, Steven W.; VerBerkmoes, Nathan C.; Lefsrud, Mark; Mueller, Ryan S.; Dick, Gregory J.; Sun, Christine L.; Wheeler, Korin E.; Zemla, Adam; Baker, Brett J.; Hauser, Loren; Land, Miriam; Shah, Manesh B; Thelen, Michael P.; Hettich, Robert L; Banfield, Jillian F., Applied and Environmental Microbiology (2009) 75, 4599-4615.
115. "Proteogenomic Monitoring of *Geobacter* Physiology during Stimulated Uranium Bioremediation," Wilkins, Michael J.; VerBerkmoes, Nathan C.; Williams, Kenneth H.; Callister, Stephen J.; Mouser, Paula J.; Elifantz, Hila; N'Guessan, A. Lucie; Thomas, Brian C.; Nicora, Carrie D.; Shah, Manesh B; Abraham, Paul; Lipton, Mary S.; Lovley D.R.; Hettich, Robert L.; Long, Philip E.; Banfield, Jillian F., Applied And Environmental Microbiology (2009) 75, 6591-6599.
116. "PTMSearchPlus: Software Tool for Automated Protein Identification and Post-Translational Modification Characterization by Integrating Accurate Intact Protein Mass and Bottom-Up Mass Spectrometric Data Searches," Kertesz V., Connelly H.M., Erickson B.K., Hettich, Robert L., Analytical Chemistry (2009) 81, 8387-8395.
117. "Phenotype Fingerprinting Suggests the Involvement of Single-Genotype Consortia in Degradation of Aromatic Compounds by *Rhodopseudomonas palustris*," Karpinets, Tatiana V.; Pelletier, Dale A.; Pan, Chongle; Uberbacher, Edward C.; Melnichenko, Galina V.; Hettich, Robert L.; Samatova, Nagiza F., PLOS ONE (2009) 4, Article Number: e4615 (Published: FEB 26 2009)
118. "Environmental Metaproteomics: Identifying the Protein Machinery that Impact the Activities and Interactions of Microbes in Isolation or Communities," Richard J. Giannone, Martin Keller, and Robert Hettich, invited book chapter for Environmental Metagenomics, Edited by Robert Li, published Feb. 2010
119. "Influence of geochemistry and biology on community protein expression within a chemoautotrophic biofilm," Mueller, Ryan S.; Deneff, Vincent J.; Kalnejais, Linda H.; Wilmes, Paul; Thomas, Brian C; Shah, Manesh B.; Hettich, Robert L.; VerBerkmoes, Nathan C.; Banfield, Jillian F., Geochimica et Cosmochimica Acta (2009) 73, A912-A912 Supplement: Supplement S.
120. "The biochemistry of unusual cytochromes isolated from acidophilic Fe(II)-oxidizing biofilms," Singer, Steven W.; Erickson, Brian; VerBerkmoes, Nathan C.; Hwang, Mona; Chan, Clara S.; Jeans, Christopher; Banfield, Jillian F.; Hettich, Robert L; Thelen, Michael P., Geochimica et Cosmochimica Acta (2009) 73, A1229-A1229 Supplement: Suppl. S Published: JUN 2009
121. "Proteogenomic basis for ecological divergence of closely related bacteria in natural acidophilic microbial communities" Deneff, Vincent J.; Kalnejais, Linda H.; Mueller, Ryan S.; Wilmes, Paul; Baker, Brett J.; Thomas, Brian C.; VerBerkmoes, Nathan C.; Hettich, Robert L.; Banfield, Jillian F., Proc. Natl. Acad. Sci. (2010) 107 2383-2390.
122. "Computational Prediction and Experimental Validation of Signal Peptide Cleavages in the Extracellular Proteome of a Natural Microbial Community," Brian K. Erickson, Ryan S. Mueller, Nathan C. VerBerkmoes, Manesh Shah, Steven W. Singer, Michael P. Thelen, Jillian F. Banfield, Robert L. Hettich, J. Proteome Res., (2010), 9, 2148-59.
123. "A high-throughput de novo sequencing approach for shotgun proteomics using high-resolution tandem mass spectrometry," Pan C, Park B.H., McDonald W.H., Carey P.A., Banfield J.F., Verberkmoes N.C., Hettich R.L., Samatova N.F. BMC Bioinformatics. (2010), 11, 118.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

124. "Cultivation and quantitative proteomic analyses of acidophilic microbial communities.," Belnap C.P., Pan C, Verberkmoes N.C., Power M.E., Samatova N.F., Carver R.L., Hettich R.L., Banfield J.F. ISME J. (2010) 4, 520-530.
125. "Enigmatic, ultra-small uncultivated Archaea," Brett J. Baker, Luis R. Comolli, Gregory J. Dick, Loren Hauser, Doug Hyatt, Brian Dill, Miriam Land, Nathan C. VerBerkmoes, Robert L. Hettich and Jillian F. Banfield, Proc. Natl. Acad. Sci., (2010), 107, 8806-11.
126. "Proteomics reveals a core molecular response of *Pseudomonas putida* F1 to acute chromate challenge," Thompson, D.K.; Chourey, K.; Wickham, G.S.; Thieman, S.B.; VerBerkmoes, N.C.; Zhang, B.; McCarthy, A.T.; Rudisill, M.A.; Shah, M.; Hettich, R.L. BMC Genomics (2010), 11, 311.
127. "Ecological distribution and population physiology defined by proteomics in a natural microbial community," Mueller, R.S.; Deneff, V.J.; Kalnejais, L.H.; Suttle, K.B.; Thomas, B.C.; Wilmes, P.; Smith, R.L.; Nordstrom, D.K.; McCleskey, R.B.; Shah, M.B.; VerBerkmoes, N.C.; Hettich, R.L.; Banfield, J.F. Molecular Systems Biology (2010), 6, 374.
128. "A Direct Cellular Lysis/Protein Extraction Protocol for Soil Metaproteomics," Chourey, K.; Jansson, J.; VerBerkmoes, N.; Shah, M.; Chavarria, K.; Tom, L.; Brodie, E.; Hettich, R.L. J. Prot. Res., (2010), 9, 6615-6622.
129. "Deletion of the Cel48S cellulase from *Clostridium thermocellum*," Olson, D.G., Tripathi, S.A., Giannone, R.J., Lo, J, Caiazza, N.C., Hogsett, D.A., Hettich, R.L., Guss, A.M., Dubrovsky, G., Lynd, L.R. Proc. Natl Acad. Sci. (2010), 107, 17727-17732.
130. "Metabolome-Proteome Differentiation Coupled to Microbial Divergence," Wilmes, P., Bowen, B.P., Thomas, B.C., Mueller, R.S., Deneff, V.J., VerBerkmoes, N.C., Hettich, R.L., Northen, T.R., Banfield, J.F. mBIO, (2010), 1, Issue: 5 Article Number: e00246 DOI: 10.1128/mBio.00246-10 .
131. "Posttranslational modification and sequence variation of redox-active proteins correlate with biofilm life cycle in natural microbial communities," Singer, S.W., Erickson, B.K., VerBerkmoes, N.C., Hwang, M., Shah, M.B., Hettich, R.L., Banfield, J.F., Thelen, M.P., ISME J., (2010), 4, 1398-1409.
132. "Analysis of Biostimulated Microbial Communities from Two Field Experiments Reveals Temporal and Spatial Differences in Proteome Profiles," Callister, S.J., Wilkins, M.J., Nicora, C.D., Williams, K.H., Banfield, J.F., VerBerkmoes, N.C., Hettich, R.L., N'Guessan, L., Mouser, P.J., Elifantz, H., Smith, R.D., Loyley, D.R., Lipton, M.S., Long, P.E., Environ. Sci. Tech., (2010), 44, 8897-8903.
133. "Quantitative Tracking of Isotope Flows in Proteomes of Microbial Communities," Pan, C.L., Fischer, C.R., Hyatt, D., Bowen, B.P., Hettich, R.L., Banfield, J.F., Mole. Cell. Proteomics, (2011), 10, Issue: 4 Article Number: 006049 DOI: 10.1074/mcp.M110.006049.
134. "Discovery and annotation of small proteins using genomics, proteomics, and computational approaches," Yang, X.H., Tschaplinski, T.J., Hurst, G.B., Jawdy, S., Abraham, P.E., Lankford, P.K., Adams, R.M., Shah, M.B., Hettich, R.L., Lindquist, E., Kalluri, U.C., Gunter, L.E., Pennacchio, C., Tuskan, G.A., Genome Res. (2011), 21, 634-641.
135. "Insights into plant biomass conversion from the genome of the anaerobic thermophilic bacterium *Caldicellulosiruptor bescii* DSM 6725," Dam, P., Kataeva, I., Yang, S.J., Zhou, F.F., Yin, Y.B., Chou, W.C., Poole, F.L., Westpheling, J., Hettich, R., Giannone, R., Lewis, D.L., Kelly, R., Gilbert, H.J., Henrissat, B., Xu, Y., Adams, M.W.W., Nuc. Acid Res. (2011), 39, 3240-3254.
136. "Use of Label-Free Quantitative Proteomics To Distinguish the Secreted Cellulolytic Systems of *Caldicellulosiruptor bescii* and *Caldicellulosiruptor obsidiansis*," Lochner, A., Giannone, R.J., Rodriguez, M., Shah, M.B., Mielenz, J.R., Keller, M., Antranikian, G., Graham, D.E., Hettich, R.L., Appl. Environ. Microbiology, (2011), 77, 4042-4054.
137. "Quantitative proteomic analyses of the response of acidophilic microbial communities to different pH conditions," Belnap, C.P., Pan, C., Deneff, V.J., Samatova, N.F., Hettich, R.L., Banfield, J.F., ISME J. (2011), 5, 1152-1161.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

138. "Identification of Biofilm Matrix-Associated Proteins from an Acid Mine Drainage Microbial Community," Jiao, Y.Q., D'haeseleer, P., Dill, B.D., Shah, M., VerBerkmoes, N.C., Hettich, R.L., Banfield, J.F., Thelen, M.P., Appl. Environ. Microbiology, (2011), 77, 5230-5237.
139. "Proteome changes in the initial bacterial colonist during ecological succession in an acid mine drainage biofilm community," Mueller, R.S., Dill, B.D., Pan, C.L., Belnap, C.P., Thomas, B.C., VerBerkmoes, N.C., Hettich, R.L., Banfield, J.F., Environ. Micro. (2011), 13, 2279-2292.
140. "Proteomic Characterization of Cellular and Molecular Processes that Enable the *Nanoarchaeum equitans-Ignicoccus hospitalis* Relationship," Giannone, R.J., Huber, H., Karpinets, T., Heimerl, T., Kuper, U., Rachel, R., Keller, M., Hettich, R.L., Podar, M., PLoS ONE, (2011), 6, Issue: 8 Article Number: e22942 DOI: 10.1371/journal.pone.0022942.
141. "Comparative shotgun proteomic analysis of *Clostridium acetobutylicum* from butanol fermentation using glucose and xylose," Sivagnanam, K., Raghavan, V.G.S., Shah, M., Hettich, R.L., Verberkmoes, N.C., Lefsrud, M.G., Proteome Sci., (2011), 9, Article Number: 66 DOI: 10.1186/1477-5956-9-66.
142. "Label-free Quantitative Proteomics for the Extremely Thermophilic Bacterium *Caldicellulosiruptor obsidiansis* Reveal Distinct Abundance Patterns upon Growth on Cellobiose, Crystalline Cellulose, and Switchgrass," Lochner, A., Giannone, R.J., Keller, M., Antranikian, G., Graham, D.E., Hettich, R.L., J. Proteome Res. (2011), 10, 5302-5314.
143. "Strategies for Metagenomic-Guided Whole-Community Proteomics of Complex Microbial Environments," Cantarel, B.L.; Erickson, A.R., VerBerkmoes, N., Erickson, B.K.; Carey, P.A.; Pan, C.; Shah, M.; Mongodin, E.F.; Jansson, J.K.; Fraser-Liggett, C.M.; Hettich, R.L.; PLoS One (2011), Volume 6, issue 11, e267173.
144. "Defining the Boundaries and Characterizing the Landscape of Functional Genome Expression in Vascular Tissues of Populus using Shotgun Proteomics," Abraham, P.; Adams, R.; Giannone, R.J.; Kalluri, U.; Ranjan, P; Erickson, B.; Shah, M.; Tuskan, G.A.; Hettich, R.L. J. Proteome Res. (2012), 11, 449-460.
145. "Metal Affinity Enrichment Increases the Range and Depth of Proteome Identification for Extracellular Microbial Proteins," Wheeler, K.E.; Erickson, B.K.; Mueller, R.; Singer, S.W.; VerBerkmoes, N.C.; Hwang, M.; Thelen, M.P.; Hettich, R.L. J. Proteome Res. (2012), 11, 861-870.
146. "Shotgun proteomic monitoring of *Clostridium acetobutylicum* during stationary phase of butanol fermentation using xylose and comparison with the exponential phase," Sivagnanam, K; Raghavan, V.G.S.; Shah, M; Hettich, R.L.; Verberkmoes, N.C.; Lefsrud, Mark G. J. Indust. Microbiol. & Biotech., (2012), 39, 949-955.
147. "Phage-Induced Expression of CRISPR-Associated Proteins Is Revealed by Shotgun Proteomics in *Streptococcus thermophilus*," Young, J.C; Dill, B.D; Pan, C.L; Hettich, R.L.; Banfield, J.F; Shah, M; Fremaux, C; Horvath, P; Barrangou, R; VerBerkmoes, N.C, PLoS ONE, (2012) 7, Article Number: e38077 DOI: 10.1371
148. "Microbes in thawing permafrost: the unknown variable in the climate change equation,"Graham, D. E.; M. D. Wallenstein, T. A. Vishnivetskaya, M. P. Waldrop, T. J. Phelps, S. M. Pfiffner, T. C. Onstott, L. G. Whyte, E. M. Rivkina, D. A. Gilichinsky, D. A. Elias, R. Mackelprang, N. C. VerBerkmoes, R. L. Hettich, D. Wagner, S. D. Wulfschleger, and J. K. Jansson. ISME J (2012) 6:709-712.
149. "The *Clavibacter michiganensis* subsp *michiganensis*-Tomato Interactome Reveals the Perception of Pathogen by the Host and Suggests Mechanisms of Infection," Savidor, Alon; Teper, Doron; Gartemann, Karl-Heinz; Eichenlaub, Rudolf; Chalupowicz, Laura; Manulis-Sasson, Shulamit; Barash, Isaac; Tews, Helena; Mayer, Kerstin; Giannone, Richard J.; Hettich, Robert L.; Sessa, Guido, J. Proteome Res. (2012), 11, 736-750.
150. "Systematic Comparison of Label-Free, Metabolic Labeling, and Isobaric Chemical Labeling for Quantitative Proteomics on LTQ Orbitrap Velos," Li, Zhou; Adams, Rachel M.; Chourey, Karuna; Hurst, Gregory B.; Hettich, Robert L.; Pan, Chongle, J. Proteome Res. (2012), 11, 1582-1590.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

151. "Microbial metaproteomics: identifying the repertoire of proteins that microorganisms use to compete and cooperate in complex environmental communities," Hettich RL, Sharma R, Chourey K, Giannone RJ. Curr Opin Microbiol. (2012) Jun;15 (3):373-80.
152. "Microbial proteomics: the quiet revolution," Seraphin, B.; Hettich R. Curr Opin Microbiol. (2012), 15, 348-350.
153. "Caldicellulosiruptor core and pan genomes reveal determinants for non-cellulosomal thermophilic deconstruction of plant biomass" Sara Blumer-Schuetz, Richard Giannone, Jeffrey Zurawski, Inci Ozdemir, Qin Ma, Yanbin Yin, Ying Xu, Irina Kataeva, Farris Poole II, Michael Adams, Scott Hamilton-Brehm, James Elkins, Frank Larimer, Miriam Land, Loren Hauser, Robert Cottingham, Robert Hettich, and Robert Kelly, J. Bact. (2012), 194, 4015-4028.
154. "Clostridium thermocellum ATCC27405 transcriptomic, metabolomic, and proteomic profiles after ethanol stress," Shihui Yang, Richard J Giannone, Lezlee Dice, Zamin K Yang, Nancy L Engle, Timothy J Tschaplinski, Robert L Hettich and Steven D Brown BMC Genomics (2012), 13, article :336, DOI: 10.1186/1471-2164-13-336
155. "Fermentation, Hydrogen and Sulfur Metabolism in Multiple Uncultivated Bacterial Phyla" Kelly C. Wrighton, Brian C. Thomas, Itai Sharon, Christopher S. Miller, Cindy Castelle, Nathan C. VerBerkmoes, Michael J. Wilkins, Robert L. Hettich, Mary Lipton, Kenneth H. Williams, Philip E. Long, and Jillian F. Banfield, Science, (2012), 337, 1661-1665.
156. "The temporal analysis of yeast exponential phase using shotgun proteomics as a fermentation monitoring technique," Huang, E.L.; Orsat, V.; Shah, M.B.; Hettich, R.L.; VerBerkmoes, N.C.; Lefsrud, M.G. J. Proteomics (2012) 75, 5206-5214.
157. "Heterotrophic archaea contribute to carbon cycling in low-pH suboxic biofilm communities," Justice, N.B.; Pan, C.; Mueller, R.; Spaulding, S.E.; Shah, V.; Sun, C.L.; Yelton, A.P.; Miller, C.S.; Thomas, B.C.; Shah, M.; VerBerkmoes, N.C.; Hettich, R.L.; Banfield, J.F. App. Environ. Microbiol. (2012) 23, 8321-8330.
158. "Comparative c-type cytochrome expression analysis in *Shewanella oneidensis* strain MR-1 and *Anaeromyxobacter dehalogenans* strain 2CP-C grown with soluble and insoluble oxidized metal electron acceptors," Nissen, S.; Liu, X.; Chourey, K.; Hettich, R.L.; Loeffler, F. Biochem. Soc. Transactions (2012), 40, 1204-1210.
159. "Integrated Metagenomics/Metaproteomics Reveals Human Host-Microbiota Signatures of Crohn's Disease," Alison R. Erickson, Brandi L. Cantarel, Regina Lamendella, Youssef Darzi, Emmanuel F. Mongodin, Chongle Pan, Manesh Shah, Jonas Halfvarson, Curt Tysk, Bernard Henrissat, Jeroen Raes, Nathan C. Verberkmoes, Claire M. Fraser, Robert L. Hettich, Janet K. Jansson, PloS ONE, (2012), 10.1371/journal.pone.0049138df.
160. "Coupling a detergent lysis/cleanup methodology with intact protein fractionation for enhanced proteome characterization," Sharma, R.; Dill, B.D.; Chourey, K.; Shah, M.; VerBerkmoes, N.C.; Hettich, R.L., J. Proteome Res. (2012), 11, 6009-6018.
161. "Putting the Pieces Together: High-performance LC-MS/MS Provides Network-, Pathway-, and Protein-level Perspectives in Populus," Abraham, P; Giannone, R.J.; Adams, R.M.; Kalluri, U.; Tuskan, G.A.; Hettich, R.L., Mole. Cell. Proteomics, (2013), 12, 106-119.
162. "Experimental Approach to Controllably Vary Protein Oxidation While Minimizing Electrode Adsorption for Boron-Doped Diamond Electrochemical Surface Mapping Applications," McClintock, C.S.; Hettich, R.L., Anal. Chem. (2013), 85, 213-219.
163. "Role of the CipA Scaffoldin Protein in Cellulose Solubilization, as Determined by Targeted Gene Deletion and Complementation in *Clostridium thermocellum*," Olson, D.G.; Giannone, R.J.; Hettich, R.L.; Lynd, L.R.; J. Bacteriology, (2013) 195, 733-739.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

164. "Characterizing the Range of Extracellular Protein Post-Translational Modifications in a Cellulose-Degrading Bacteria Using a Multiple Proteolytic Digestion/Peptide Fragmentation Approach," Dykstra, A.B.; Rodriguez, M.; Raman, B.; Cook, K.D.; Hettich, R.L. Anal. Chem. (2013) 85, 3144-3151.
165. "Biostimulation induces syntrophic interactions that impact C, S and N cycling in a sediment microbial community," Handley, K.M.; VerBerkmoes, N.C.; Steefel, C.I.; Williams, K.H.; Sharon, I; Miller, C.S.; Frischkorn, K.R.; Chourey, K; Thomas, B.C.; Shah, M.B.; Long, P.E.; Hettich R.L.; Banfield, J.F., ISME J. (2013), 7, 800-816.
166. "Metaproteomics: Harnessing the Power of High Performance Mass Spectrometry to Identify the Suite of Proteins That Control Metabolic Activities in Microbial Communities," Hettich, R.L.; Pan, C.L.; Chourey, K.; Giannone, R.J., Anal. Chem. (2013), 85, 4203-4214.
167. "A new group in the *Leptospirillum* clade: cultivation-independent community genomics, proteomics and transcriptomics of the new species *Leptospirillum* group IV UBA BS.," Aliaga Goltsman DS, Dasari M, Thomas BC, Shah MB, Verberkmoes NC, Hettich RL, Banfield JF. App. Environ. Microbiol., (2013), 79, 5384-5393.
168. "Insights into the structure and metabolic function of microbes that shape pelagic iron-rich aggregates ("iron snow")," Lu S, Chourey K, Reiche M, Nietzsche S, Shah MB, Neu TR, Hettich RL, Küsel K. Appl Environ Microbiol. (2013), 79, 4272-4281.
169. "Moving Away from the Reference Genome: Evaluating a Peptide Sequencing Tagging Approach for Single Amino Acid Polymorphism Identifications in the Genus *Populus*." Abraham P, Adams RM, Tuskan GA, Hettich RL. J Proteome Res. (2013), 12, 3642-3651.
170. "Comparative Informatics Analysis to Evaluate Site-Specific Protein Oxidation in Multidimensional LC-MS/MS Data," McClintock CS, Parks JM, Bern M, Ghattyvenkatakrishna PK, Hettich RL. J. Proteome Res. (2013), 12, 3307-3316.
171. "Fungal community responses to discrete precipitation pulses under altered rainfall intervals," A. Jumpponen, L. Zeglin, M. David, Prestat, E; Brown, S; Dvornik, J; Lothamer, K; Hettich, R.; Jansson, J; Rice, CW; Tringe, S; Myrold, D., Phytopathology, (2013), 103, 182-183.
172. "Effects of Diet on Resource Utilization by a Model Human Gut Microbiota Containing *Bacteroides cellulosilyticus* WH2, a Symbiont with an Extensive Glycobiome," Nathan P. McNulty, Meng Wu, Alison R. Erickson, Chongle Pan, Brian K. Erickson, Eric C. Martens, Nicholas A. Pudlo, Brian Muegge, Bernard Henrissat, Robert L. Hettich, Jeffrey I. Gordon, PLOS Biology, (2013), 11, Issue: 8, Article Number: e1001637.
173. "Isolation and chemical analysis of nanoparticles from English ivy (*Hedera helix* L.)," Lenaghan, SC ; Burris, JN; Chourey, K; Huang, YJ; Xia, LJ; Lady, B; Sharma, R; Pan, CL; LeJeune, Z; Foister, S; Hettich, RL ; Stewart, CN, J. Royal Soc. Interface, (2013), vol 10, issue 87, Article Number: 20130392.
174. "Environmental proteomics reveals early microbial community responses to biostimulation at a uranium- and nitrate-contaminated site," Chourey, K; Nissen, S; Vishnivetskaya, T; Shah, M; Pfiffner, S; Hettich, RL ; Löffler, FE, Proteomics , (2013), 13, 2921-2930.
175. "Identification and Environmental Distribution of *dcpA*, Which Encodes the Reductive Dehalogenase Catalyzing the Dichloroelimination of 1,2-Dichloropropane to Propene in Organohalide-Respiring *Chloroflexi*," Padilla-Crespo, E; Yan, J; Swift, C; Wagner, DD; Chourey, K; Hettich, RL; Ritalahti, KM; Löffler, App. Environ. Microbiol., (2014), 80, 808-818.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

176. "Geochemical, metagenomic and metaproteomic insights into trace metal utilization by methane-oxidizing microbial consortia in sulphidic marine sediments," Glass JB, Yu H, Steele JA, Dawson KS, Sun S, Chourey K, Pan C, Hettich RL, Orphan VJ. Environ Microbiol. (2014) 16, 1592-1611.
177. "Development of a Multipoint Quantitation Method to Simultaneously Measure Enzymatic and Structural Components of the *Clostridium thermocellum* Cellulosome Protein Complex," Dykstra AB, St Brice L, Rodriguez M Jr, Raman B, Izquierdo J, Cook KD, Lynd LR, Hettich RL. J Proteome Res. (2014), 13, 692-701.
178. "Metaproteomics: extracting and mining proteome information to characterize metabolic activities in microbial communities," Paul Abraham, Richard J. Giannone, Weili Xiong, Robert L. Hettich, Current Protocols for Bioinformatics, (2014), 46, 13.26.1-13.16.14.
179. "15 N- and 2 H proteomic stable isotope probing links nitrogen flow to archaeal heterotrophic activity," Justice, N.B.; Li, Z.; Wang, Y; Spaulding, S.E.; Mosier, A.C.; Hettich, R.L.; Pan, C; Banfield, J.F. Environ Microbiol. (2014) 16, 3224-3237.
180. "Diverse and divergent protein post-translational modifications in two growth stages of a natural microbial community," Zhou Li; Yingfeng Wang; Qiuming Yao; Nicholas B. Justice; Tae-Hyuk Ahn; Dong Xu; Robert L. Hettich; Jillian F. Banfield; Chongle Pan, Nature Comm. (2014), 5, 4405.
181. "Recoding of the stop codon UGA to glycine by a BD1-5/SN-2 bacterium and niche partitioning between Alpha- and Gammaproteobacteria in a tidal sediment microbial community naturally selected in a laboratory chemostat," Anna Hanke, Emmo Hamann, Ritin Sharma, Jeanine S. Geelhoed, Theresa Hargesheimer, Beate Kraft, Volker Meyer, Sabine Lenk, Harald Osmers, Rong Wu, Kofi Makinwa, Robert L. Hettich, Jillian F. Banfield, Halina E. Tegetmeyer, and Marc Strous, Frontiers Microbiol., (2014), 5, Article number 231.
182. "Life on the edge: functional genomic response of *Ignicoccus hospitalis* to the presence of *Nanoarchaeum equitans*," Giannone R.J, Wurch L.L, Heimerl T., Martin S., Yang Z., Huber H., Rachel R., Hettich R.L, Podar M., ISME J. (2015), 9, 101-114.
183. "Profile of secreted hydrolases, associated proteins, and SlpA in *Thermoanaerobacterium saccharolyticum* during the degradation of hemicelluloses," Currie D.H, Guss A.M, Herring C.D, Giannone R.J, Johnson C.M, Lankford P.K, Brown S.D, Hettich R.L, Lynd L.R. Appl Environ Microbiol. (2014), 80, 5001-5011
184. "Elevated temperature alters proteomic responses of individual organisms within a biofilm community," Mosier, A.C., Li, Z., Thomas, B.C., Hettich, R.L., Pan, C., Banfield, J.F., ISME J. (2014) 9, 180-194.
185. "The environmental controls that govern the end product of bacterial nitrate respiration" Kraft, B., Tegetmeyer, H.E., Sharma, R., Klotz, M.G., Ferdelman, T.G., Hettich, R.L., Geelhoed, J.S., Strous, M., Science. (2014), 345, 676-679.
186. "Development of an Enhanced Metaproteomic Approach for Deepening the Microbiome Characterization of the Human Infant Gut," Xiong W, Giannone RJ, Morowitz MJ, Banfield JF, Hettich RL. J Proteome Res. (2014) 14, 133-141.
187. "Untargeted metabolomics studies employing NMR and LC-MS reveal metabolic coupling between *Nanoarchaeum equitans* and its archaeal host *Ignicoccus hospitalis*," Timothy Hamerly, Brian P. Tripet, Michelle Tigges, Richard J. Giannone, Louie Wurch, Robert L. Hettich, Mircea Podar, Valerie Copie, Brian Bothner Metabolomics (2015), 11, pp. 895-907.

188. "A comparative multidimensional LC-MS proteomic analysis reveals mechanisms for furan aldehyde detoxification in *Thermoanaerobacter pseudethanolicus* 39E." Sonya M Clarkson, Scott D Hamilton-Brehm, Richard J Giannone, Nancy L Engle, Timothy J Tschaplinski, Robert L Hettich, James G Elkins Biotechnology for Biofuels (2014), 7:165 doi:10.1186/s13068-014-0165-z.
189. "An active atmospheric methane sink in high Arctic mineral cryosols," M.C.Y. Lau, B.T. Stackhouse, A.C. Layton, A. Chauhan, T.A. Vishnivetskaya, K. Chourey, J. Ronholm, N.C.S. Mykytczuk, P.C. Bennett, G. Lamarche-Gagnon, N. Burton, W.H. Pollard, C.R. Omelon, D.M. Medvigy, R.L. Hettich, S.M. Pfiffner, L.G. Whyte, and T.C. Onstott, ISME J. (2015), 9, 1880-1891.
190. "Discrete and Structurally Unique Proteins (Tapirins) Mediate Attachment of Extremely Thermophilic *Caldicellulosiruptor* Species to Cellulose," Sara E. Blumer-Schuetz, Markus Alahuhta, Jonathan M. Conway, Laura L. Lee, Jeffrey V. Zurawski, Richard J. Giannone, Robert L. Hettich, Vladimir V. Lunin, Michael E. Himmel, and Robert M. Kelly, J. Bacteriology, (2015), 290, 10645-10656.
191. "A roadmap for research on crassulacean acid metabolism (CAM) to enhance sustainable food and bioenergy production in a hotter, drier world," Xiaohan Yang¹, John C. Cushman², Anne M. Borland^{1,3}, Erika J. Edwards⁴, Stan D. Wullschlegel⁵, Gerald A. Tuskan¹, Nick A. Owen⁶, Howard Griffiths⁶, J. Andrew C. Smith⁷, Henrike C. De Paoli¹, David J. Weston¹, Robert Cottingham¹, James Hartwell⁸, Sarah C. Davis⁹, Katia Silvera¹⁰, Ray Ming^{11,12}, Karen Schlauch¹³, Paul Abraham¹⁴, J. Ryan Stewart¹⁵, Hao-Bo Guo¹⁶, Rebecca Albion², Jungmin Ha², Sung Don Lim², Bernard W. M. Wone², Won Cheol Yim², Travis Garcia², Jesse A. Mayer², Juli Petereit¹³, Sujithkumar S. Nair⁵, Erin Casey³, Robert L. Hettich¹⁴, Johan Ceusters¹⁷, Priya Ranjan¹, Kaitlin J. Palla¹, Hengfu Yin¹⁸, Casandra Reyes-García¹⁹, José Luis Andrade¹⁹, Luciano Freschi²⁰, Louisa V. Dever⁸, Susanna F. Boxall⁸, Jade Waller⁸, Jack Davies⁸, Phaitun Bupphada⁸, Nirja Kadu⁸, Klaus Winter¹⁰, Rowan F. Sage²¹, Cristobal N. Aguilar²², Jeremy Schmutz^{23,24}, Jerry Jenkins²³, Joseph A.M. Holtum²⁵, New Phytologist, (2015), 207, pp. 491-504.
192. "Metagenome sequencing of a coastal marine microbial community from Monterey Bay, California," Mueller R.S., Bryson S., Kieft B., Li Z., Pett-Ridge J., Chavez F., Hettich R.L., Pan C., Mayali X. Genome Announc. (2015) Apr 30;3(2). pii: e00341-15. doi: 10.1128/genomeA.00341-15.
193. "Critical biogeochemical functions in the subsurface are associated with bacteria from new phyla and little studied lineages," Hug L.A., Thomas B.C., Sharon I., Brown C.T., Sharma R., Hettich R.L., Wilkins M.J., Williams K.H., Singh A., Banfield J.F. Environ Microbiol. (2016) 18, pp. 159-173.
194. "Microbial metaproteomics for characterizing the range of metabolic functions and activities of human gut microbiota," Xiong W., Abraham P.E., Li Z., Pan C., Hettich R.L. Proteomics. (2015), 20, pp. 2423-3438.
195. "Metaproteomics Reveals Functional Shifts in Microbial and Human Proteins During a Preterm Infant Gut Colonization Case," Jacque C. Young, Chongle Pan, Rachel Adams, Brandon Brooks, Jillian F. Banfield, Michael J. Morowitz, Robert L. Hettich. Proteomics, (2015), 15, pp. 3463-3473
196. "Strain-resolved microbial community proteomics reveals simultaneous aerobic and anaerobic function during early stage gastrointestinal tract colonization," Brandon Brooks, Ryan Mueller, Jacque C. Young, Michael J. Morowitz, Robert L. Hettich, Jillian F. Banfield. Frontiers in Microbiology, (2015), 6, article number 654.
197. "Rescuing Those Left Behind: Recovering and Characterizing Underdigested Membrane and Hydrophobic Proteins to Enhance Proteome Sampling," Richard J. Giannone, Louie L. Wurch, Mircea Podar, and Robert L. Hettich. Anal. Chem. (2015), 87, pp. 7720-7728.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

198. "Integrating mRNA and Protein Sequencing for the Identification of Natural Protein Sequence Variants: A Case Study in the Genus *Populus*," Paul E. Abraham, Xiaojing Wang, Priya Ranjan, Intawat Nookaew, Bing Zhang, Gerald A. Tuskan, Robert L. Hettich, J. Proteome Res. (2015), 14, 5318-5326.
199. "Development of a detergent based method for cell lysis and protein extraction from soil samples for metaproteomics measurements/applications," Karuna Chourey and Robert L. Hettich, Book Chapter for Springer, Methods Mol Biol. (2018);1841:293-302. doi: 10.1007/978-1-4939-8695-8_20. PMID: 30259494
200. "Dramatic performance of *Clostridium thermocellum* explained by its wide range of cellulase modalities." by Qi Xu, Michael Resch, Kara Podkaminer, Shihui Yang, John Baker, Bryon Donohoe, Charlotte Wilson, Dawn Klingeman, Daniel Olson, Stephen Decker, Richard Giannone, Robert Hettich, Steven Brown, Lee Lynd, Edward Bayer, Michael Himmel, and Yannick Bomble, Science Advances, (2016), 2, DOI: 10.1126/SciAdv. IE01254.
201. "Genome Sequence of *Streptomyces vitaminophilus* ATCC 31673, a Producer of Pyrrolomycin Antibiotics, Some of Which Contain a Nitro Group," Kristina M. Mahan, Dawn M. Klingeman, Robert L. Hettich, Ronald J. Parry, David E. Graham. Genome Announc. (2016), 4 (1), e01582-15.
202. "Characterization of fatty acids in Crenarchaeota by GC-MS and NMR" Timothy Hamerly, Brian Tripet, Louie Wurch, Robert L. Hettich, Mircea Podar, Brian Bothner and Valerie Copie, Archaea, (2015), Article Number: 472726, DOI: 10.1155/2015/472726.
203. "Enhancing Metaproteomics - The value of models and defined environmental microbial systems," Herbst FA, Lünsmann V, Kjeldal H, Jehmlich N, Tholey A, von Bergen M, Nielsen JL, Hettich RL, Seifert J, Nielsen PH. Proteomics. (2016) 16, 783-798.
204. "Proteomic Stable Isotope Probing Reveals Dynamic Metabolism and Diverse Methanotrophic Activity at Marine Methane Seeps, Jeffrey J. Marlow, Connor T. Skennerton, Zhou Li, Karuna Chourey, Robert L. Hettich, Chongle Pan, Victoria J. Orphan. Frontiers in Microbiology, (2016), 7, 563.
205. "Proteogenomic analyses indicate bacterial methylotrophy and archaeal heterotrophy are prevalent below the grass root zone," Cristina N. Butterfield, Zhou Li, Peter Andeer, Susan Spaulding, Brian C. Thomas, Andrea Singh, Robert Hettich, K. Blake Suttle, Susannah Tringe, Trent Northen, Chongle Pan, Jillian F. Banfield, PeerJ, (2016), Volume 4, Article Number: e2687.
206. "Proteomic-based stable isotope probing reveals taxonomically distinct patterns in amino acid assimilation by coastal marine bacterioplankton" Samuel Bryson, Zhou Li, Jennifer Pett-Ridge, Robert Hettich, Xavier Mayali, Chongle Pan, and Ryan Mueller mSystems, (2016), 1, 27.
207. "Fungi contribute critical but spatially varying roles in nitrogen and carbon cycling in acidic mine drainage," Annika Mosier; Christopher S. Miller; Kyle Frischkorn; Robin Ohm; Zhou Li; Kurt LaButti; Alla Lapidus; Anna Lipzen; Cindy Chen; Jennifer Kaplan; Erika Lindquist; Chongle Pan; Robert Hettich; Igor Grigoriev; Steven Singer; Jillian Banfield Frontiers in Microbiology, (2016), 7, 1-18.
208. "Development of a Plasmid-Based Expression System in *Clostridium thermocellum* and its use to Screen Heterologous Expression of bifunctional alcohol dehydrogenases (adhEs)," Shuen Hon, Anthony A Lanahan; Liang Tian; Richard J Giannone; Robert L Hettich; Daniel G Olson; Lee R Lynd. Metabolic Engineering Communications, (2016), 3, 120-129.
209. "Global Proteome Response to Deletion of Genes Related to Mercury Methylation and Dissimilatory Metal Reduction Reveals Changes in Respiratory Metabolism in *Geobacter sulfurreducens* PCA," C Qian, A Johs, H Chen, BF Mann, X Lu, PE Abraham, RL Hettich, B Gu. Journal of Proteome Research (2016) 15, 3540-3549

210. "Reconstructing a hydrogen-driven microbial metabolic network in Opalinus Clay rock," Alexandre Bagnoud, Karuna Chourey, Robert L. Hettich, Ino de Bruijn, Anders F. Andersson, Olivier X. Leupin, Bernhard Schwyn, Rizlan Bernier-Latmani, Nature Comm. (2016), Volume 7, Article Number 12115, DOI: 10.1038/ncomms12770.
211. "From Genomics-informed isolation and characterization of a symbiotic Nanoarchaeota system from a terrestrial geothermal environment," Wurch L., Gianonne R., Belisle B., Swift C., Utturkar S., Hettich R., Reysenbach A.L. Podar M., Nature Comm. (2016), 7, Article Number: 12115, DOI: 10.1038/ncomms12115
212. "The temporal dynamics of transcripts, proteins, and metabolites that define crassulacean acid metabolism in Agave," Paul E. Abraham, Hengfu Yin, Anne M. Borland, Deborah Weighill, Sung Don Lim, Henrique Cestari De Paoli, Nancy Engle, Ryan Agh, David J. Weston, Stan D. Wullschleger, Timothy Tschaplinski, Daniel Jacobson, John C. Cushman, Robert L. Hettich, Gerald A. Tuskan, Xiaohan Yang, Nature Plant, (2016), 2, Article Number: 16178.
213. "Metabolic Regulation as a consequence of anaerobic 5-methylthioadenosine recycling in *Rhodospirillum rubrum*," Justin A. North, Jaya Srirama, Karuna Chourey, Christopher D. Eckera, Ritin Sharma, John A. Wildenthala, Robert L. Hettich, F. Robert Tabita, mBio, (2016), 7, Article Number: e00855-16.
214. "Transcriptomic and proteomic insights into innate immunity and adaptations to a symbiotic lifestyle in the gutless marine worm *Olavius algarvensis*," Juliane Wippler, Manuel Kleiner, Christian Lott, Alexander Gruhl, Paul Abraham, Rich Giannone, Jacque C. Young, Robert L. Hettich, Nicole Dubilier, BMC Genomics (2016), 17, article 942, DOI: 10.1186/s12864-016-3293-y.
215. "Improved growth rate in *Clostridium thermocellum* hydrogenase mutant via perturbed sulfur metabolism," Ranjita Biswas, Charlotte M. Wilson, Richard J. Giannone, Dawn M. Klingeman, Robert L. Hettich, Steven D. Brown, Adam M. Guss, Biotech. For Biofuels, (2017), 10, Article Number: 6, doi: 10.1186/s13068-016-0684-x.
216. "Simultaneous achievement of high ethanol yield and titer in *Clostridium thermocellum*," Tian, L.; Papanek, B.; Olson, D.G.; Rydzak, T.; Holwerda, E.K.; Zheng, T.Y.; Zhou, J.L.; Maloney, M.; Jiang, N.N.; Giannone, R.J.; Hettich, R.L.; Guss, A.M.; Lynd, L.R. Biotech. For Biofuels, (2016), 9, Article Number: 116.
217. "Integrated 'omics analyses reveal the details of metabolic adaptation of *Clostridium thermocellum* to lignocellulose-derived growth inhibitors released during the deconstruction of switchgrass," Suresh Poudel, Richard J. Giannone; Miguel Rodriguez Jr.; Babu Raman; Madhavi Z. Martin; Nancy L. Engle; Jonathan R. Mielenz; Intawat Nookaew; Steve D. Brown; Timothy J. Tschaplinski; David Ussery; and Robert L. Hettich, Biotech. For Biofuels, (2017), 10, Article Number: 14.
218. "Impacts of chemical gradients on microbial community structure," Marc Strous, Jianwei Chen, Halina Tegetmeyer, Ines Kattmann, Ritin Sharma, Emmo Hamann, Theresa Hargesheimer, Beate Kraft, Sabine Lenk, Robert Hettich, and Jeanine Geelhoed. ISME J. (2017), 920-931.
219. "Genome-resolved meta-omics ties microbial dynamics to process performance in biotechnology for thiocyanate degradation," Rose S. Kantor, Robert J. Huddy, Ramsunder Iyer, Brian C. Thomas, Christopher T. Brown, Karthik Anantharaman, Susannah Tringe, Robert L. Hettich, Susan T. L. Harrison, Jillian F. Banfield, Env. Sci. Tech. (2017), 51, 2944-2953.
220. "Specialized activities and expression differences for *Clostridium thermocellum* biofilm and planktonic cells," Alexandru Dumitrache, Dawn M. Klingeman, Jace Natzke, Miguel Rodriguez, Richard J. Giannone, Robert L. Hettich, Brian H. Davison, Steven D. Brown, Scientific Reports, (2017), 7, Article Number: 43583.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

221. "Pentose sugars inhibit metabolism and invoke a cell-to-cell communication response in *Clostridium thermocellum*," Tobin J. Verbeke, Richard J. Giannone, Dawn M. Klingeman, Nancy L. Engle, Thomas Rydzak, Adam M. Guss, Timothy J. Tschaplinski, Steven D. Brown, Robert L. Hettich, James G. Elkins, Scientific Reports, (2017), 7, Article Number: 43355.
222. "The protein and neutral lipid composition of lipid droplets isolated from the fission yeast, *Schizosaccharomyces pombe*." Meyers, A; Chourey, K; Weiskittel, TM; Pfiffner, S; Dunlap, JR; Hettich, RL; Dalhaimer, P. J. Microbiol, (2017) 55, 112-122.
223. "An optimized extraction method to remove humic acids interferences from a proteome measurement of environmental soil samples," Chen Qian, Robert L. Hettich, J. Proteome Res. (2017), 16, 2537-2546.
224. "The effects of micronutrient deficiencies on the human gut microbiota," Matthew C. Hibberd, Meng Wu, Dmitry A. Rodionov, Xiaoqing Li, Jiye Cheng, Nicholas W. Griffin, Michael J. Barratt, Richard J. Giannone, Robert L. Hettich, Andrei L. Osterman, and Jeffrey I. Gordon, Science Translational Medicine, (2017), 9 Issue: 390 Article Number: eaal4069.
225. "Methane-fuelled syntrophy through extracellular electron transfer: uncovering the genomic traits conserved within diverse bacterial partners of ANME archaea," Connor T. Skennerton, Karuna Chourey, Ramsunder Iyer, Robert L. Hettich, Gene W. Tyson, and Victoria J. Orphan, mBio (2017), 8, Article Number: e01561-17.
226. "Phylogenetically conserved resource partitioning in the coastal microbial loop," Samuel Bryson, Zhou Li, Francisco Chavez, Jennifer Pett-Ridge, Robert L. Hettich, Chongle Pan, Xavier Mayali, Ryan S. Mueller, ISME J. (2017), 11, Pages: 2781-2792
227. "Multi-level Omics Analysis Provides Insight to the *Ignicoccus hospitalis*-*Nanoarchaeum equitans* Association," Rawle RA, Hamerly T, Tripet BP, Giannone RJ, Wurch L, Hettich RL, Podar M, Copié V, Bothner B. Biochim Biophys Acta. (2017) Volume: 1861 Issue: 9 Pages: 2218-2227
228. "Genome-resolved metaproteomic characterization of preterm infant gut microbiota development reveals species-specific metabolic shifts and variabilities during early-life," Weili Xiong, Christopher T. Brown, Michael J. Morowitz, Jillian F. Banfield, Robert L. Hettich, Microbiome (2017), 5, article 72.
229. "Grape Pomace Compost Harbors Organohalide-Respiring *Dehalogenimonas* Species with Novel Reductive Dehalogenase Genes," Yi Yang, Steven A. Higgins, Jun Yan, Burcu Şimşir, Karuna Chourey, Ramsunder Iyer, Robert L. Hettich, Brett Baldwin, Dora M. Ogles, and Frank E. Löffler, ISME J. (2017), Volume: 11 Issue: 12 Pages: 2767-2780.
230. "Eliminating a global regulator of carbon catabolite repression enhances the conversion of aromatic lignin monomers to muconate in *Pseudomonas putida* KT2440," Christopher W. Johnson, Paul E. Abraham, Jeffrey G. Linger, Payal Khanna, Robert L. Hettich, Gregg T. Beckham, Metabolic Engineering Communications, (2017), 5, pages 19-25.
231. "Identification of peptide binding sites within BSA using rapid, laser-induced covalent cross-linking combined with high-performance mass spectrometry," Melinda Hauser, Chen Qian, Steven T. King, Sarah Kauffman, Fred Naider, Robert Hettich, and Jeffrey M. Becker, J. Molecular Recognition, (2018), 31, Article Number: e2680, DOI: 10.1002/jmr.2680
232. "The *Kalanchoë* genome provides insights into convergent evolution 1 and building blocks of crassulacean acid metabolism," Xiaohan Yang, Rongbin Hu, Hengfu Yin, Jerry Jenkins, Shengqiang Shu, Haibao Tang, Degao Liu, Deborah A. Weighill, Won Cheo Yim, Jungmin Ha, Karolina Heyduk, David M. Goodstein, Hao-Bo Guo, Robert C. Moseley, Elisabeth Fitzek, Sara Jawdy, Zhihao Zhang, Meng Xie, James Hartwell, Jane Grimwood, Paul E. Abraham, Ritesh Mewalal, Juan D. Beltrán, Susanna F. Boxall, Louisa V. Dever,

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

Kaitlin J. Palla, Rebecca Albion, Travis Garcia, Jesse Mayer, Sung Don Lim, Ching Man Wai, Paul Peluso, Robert Van Buren, Henrique Cestari De Paoli, Anne M. Borland, Hong Guo, Jin-Gui Chen, Wellington Muchero, Yanbin Yin, Daniel A. Jacobson, Timothy J. Tschaplinski, Robert L. Hettich, Ray Ming, Klaus Winter, James H. Leebens-Mack, J. Andrew C. Smith, John C. Cushman, Jeremy Schmutz, Gerald A. Tuskan, Nature Comm. (2017), 8, Article Number: 1899.

233. “Mechanisms of subzero growth in the cryophile *Planococcus halocryophilus* determined through proteomic analysis,” Isabelle Raymond-Bouchard, Karuna Chourey, Ianina Altshuler, Ramsunder Iyer, Robert L. Hettich, Lyle G. Whyte, Environ. Microbiol. (2017), 19, Pages: 4460-4479.
234. “Functional Analysis of the Glucan Degradation Locus in *Caldicellulosiruptor bescii* Reveals Essential Roles of Component Glycoside Hydrolases in Plant Biomass Deconstruction, Conway, J.M; McKinley, B.S; Seals, N.L; Hernandez, D; Khatibi, P.A; Poudel, S; Giannone, R.J; Hettich, R.L; Williams-Rhaesa, A.M; Lipscomb, G.L; Adams, M.W.W; Kelly, R.M. Appl. Environ. Microbiology (2017), 83, Article Number: UNSP e01828-17.
235. “Deletion of the *hfsB* gene increases ethanol production in *Thermoanaerobacterium saccharolyticum* and several other thermophilic anaerobic bacteria,” Eminoglu, A; Murphy, S.J.L; Maloney, M; Lanahan, A; Giannone, R.J; Hettich, R.L; Tripathi, S.A; Belduz, A.O; Lynd, L.R; Olson, D.G. Biotech for Biofuels, (2017), 10, Article Number: 282, DOI: 10.1186/s13068-017-0968-9.
236. “Maternal consumption of fish oil programs reduced adiposity in broiler chicks,” Beckford, R.C; Howard, S.J; Das, S; Farmer, A.T; Campagna, S.R; Yu, J.L; Hettich, R.L; Wilson, J.L; Voy, B.H. Sci. Reports, (2018), 8, Article Number: 11578.
237. “Quantitative proteome profile of water deficit stress responses in eastern cottonwood (*Populus deltoides*) leaves,” Abraham, PE; Garcia, BJ; Gunter, LE; Jawdy, SS; Engle, N; Yang, XH; Jacobson, DA; Hettich, RL; Tuskan, GA; Tschaplinski, TJ. PlosOne, (2018), 13, Article Number: e0190019, DOI: 10.1371/journal.pone.0190019.
238. “Phosphorylation of Ribosomal Protein RPS6 Integrates Light Signals and Circadian Clock Signals,” Enganti, R; Cho, SK; Toperzer, JD; Urquidi-Camacho, RA; Cakir, OS; Ray, AP; Abraham, PE; Hettich, RL; von Arnim, AG. Front.Plant Sci. (2018), 8, Article Number: 2210, DOI: 10.3389/fpls.2017.02210.
239. “Insights into the Evolution of Host Association through the Isolation and Characterization of a Novel Human Periodontal Pathobiont, *Desulfobulbus oralis*, Karissa L Cross, Payal Chirania, Weili Xiong, Clifford J Beall, James G Elkins, Richard J Giannone, Ann L Griffen, Adam M Guss, Robert L Hettich, Snehal S Joshi, Elaine M Mokrzan, Roman K Martin, Igor B Zhulin, Eugene J Leys, Mircea Podar. mBio, (2018), 9, article number e02061-17
240. “The diversity and specificity of the extracellular proteome in the cellulolytic bacterium *Caldicellulosiruptor bescii* is driven by the nature of the cellulosic growth substrate,” Suresh Poudel, Richard J. Giannone, Mirko Basen, Intawat Nookaew, Farris L. Poole II, Robert M. Kelly, Michael W. W. Adams, Robert L. Hettich. Biotech for Biofuels, (2018), Volume: 11, Article Number: 80
241. “Hospitalized Premature Infants Are Colonized by Related Bacterial Strains with Distinct Proteomic Profiles,” Christopher. T. Brown, Weili Xiong, Matthew R. Olm, Brian C. Thomas, Robyn Baker, Brian Firek Michael J. Morowitz, Robert L. Hettich, Jillian F. Banfield. mBio, (2018), Volume: 9, Article Number: e00441-18
242. “Proteogenomics reveals novel reductive dehalogenases and methyltransferases putatively involved in anaerobic dichloromethane metabolism,” Sara Kleindienst, Karuna Chourey, Gao Chen, Steven A. Higgins, Ramsunder Iyer, Shawn R. Campagna, E. Erin Mack, Edward S. Seger, Robert L. Hettich, Frank E. Löffler. Appl Environ. Microbiol. (2019), 85, article Number: UNSP e02768-18, DOI: 10.1128/AEM.02768-18

243. "Microbial community structure-function relationships in Yaquina Bay estuary reveal spatially distinct carbon and nitrogen cycling capacities," Kieft, B.; Li, Z.; Bryson, S.; Crump, B.C.; Hettich, R.; Pan, C.L.; Mayali, X.; Mueller, R.S. Front. Microbiol., (2018), Volume: 9, Article Number: 1282.
244. "Diel rewiring and positive selection of ancient plant proteins enabled evolution of CAM photosynthesis in Agave." Yin, H.F.; Guo, H.B.; Weston, D.J.; Borland, A.M.; Ranjan, P.; Abraham, P.E.; Jawdy, S.S.; Wachira, J.; Tuskan, G.A.; Tschaplinski, T.J.; Wullschleger, S.D.; Guo, H.; Hettich, R.L.; Gross, S.M.; Wang, Z.; Visel, A.; Yang, X.H. BMC Genomics, (2018), 19, Article Number: 588.
245. "Thermochemical wastewater valorization via enhanced microbial toxicity tolerance," Jayakody, L.N.; Johnson, C.W.; Whitham, J.M.; Giannone, R.J.; Black, B.A.; Cleveland, N.S.; Klingeman, D.M.; Michener, W.E.; Olstad, J.L.; Vardon, D.R.; Brown, R.C.; Brown, S.D.; Hettich, R.L.; Guss, A.M.; Beckham, G.T. Energy & Environ. Sci. (2018), Volume: 11, Issue: 6, Pages: 1625-1638.
246. "Comparative Genomics and Proteomic Analysis of Assimilatory Sulfate Reduction Pathways in Anaerobic Methanotrophic Archaea," Yu H., Susanti D., McGlynn S.E., Skennerton C.T., Chourey K., Iyer R., Scheller S., Tavormina P.L., Hettich R.L., Mukhopadhyay B., Orphan V.J. Front Microbiol. (2018) Dec 3;9:2917. doi: 10.3389/fmicb.2018.02917. eCollection 2018.
247. "Quantitative Proteomic Analysis of Biological Processes and Responses of the Bacterium *Desulfovibrio desulfuricans* ND132 upon Deletion of Its Mercury Methylation Genes." Qian C., Chen H., Johs A., Lu X., An J., Pierce E.M., Parks J.M., Elias D.A., Hettich R.L., Gu B, Proteomics. (2018), Volume: 18 Issue: 17 Article Number: 1700479
248. "Quantifying Codon Usage in Signal Peptides: Gene Expression and Amino Acid Usage Explain Apparent Selection for Inefficient Codons," BBA – Biomembranes, (2018), Cope, A.; Hettich, R.L.; Gilchrist, M.A., 1860, 2479-2485. doi: 10.1016/j.bbamem.2018.09.010.
249. "Transcriptomic and proteomic changes from medium supplementation and strain evolution in high yielding *Clostridium thermocellum* strains," Papanek, B.; O'Dell, K.B.; Manga, P.; Giannone, R.J.; Klingeman, D.M.; Hettich, R.L.; Brown, S.D.; Guss, A.M. Journal of Industrial Microbiology & Biotechnology (2018), Volume: 45 Issue: 11 Pages: 1007-1015.
250. "Biogeochemical cycling by a low-diversity microbial community in deep groundwater," Emma Bell, Tiina Lamminmäki, Johannes Alneberg, Anders F Andersson, Chen Qian, Weili Xiong, Robert L Hettich, Louise Balmer, Manon Frutschi, Guillaume Sommer, Rizlan Bernier-Latmani, (2018), Front. In Microbiol., Volume: 9 Article Number: 2129.
251. "Metaproteomics reveals persistent and phylum-redundant metabolic functional stability in adult human gut microbiomes of Crohn's remission patients despite temporal variations in microbial taxa, genomes, and proteomes," Blakeley-Ruiz, J.A.; Erickson, A.R.; Cantarel, B.L.; Xiong, W.L.; Adams, R.; Jansson, J.K.; Fraser, C.M.; Hettich, R.L. (2019) Microbiome, 7, Article Number: 18.
252. "Scavenging organic nitrogen and remodeling lipid metabolism are key survival strategies adopted by the endophytic fungi, *Serendipita vermifera* and *Serendipita bescii* to alleviate nitrogen and phosphorous starvation in vitro." Ray P., Abraham P.E., Guo Y., Giannone R.J., Engle N.L., Yang Z.K., Jacobson D., Hettich R.L., Tschaplinski T.J., Craven K.D. Environ Microbiol Rep. (2019), 11, 5487-557. doi: 10.1111/1758-2229.12757.
253. "Evaluation of an untargeted nano-liquid chromatography-mass spectrometry approach to expand coverage of low molecular weight dissolved organic matter in Arctic soil," Ladd M.P., Giannone R.J., Abraham P.E., Wullschleger S.D., Hettich R.L. Sci Rep. (2019) 9, article 5810.

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.

254. "Progress and Challenges in Ocean Metaproteomics and Proposed Best Practices for Data Sharing," Saito M.A., Bertrand E.M., Duffy M.E., Gaylord D.A., Held N.A., Hervey W.J. 4th, Hettich R.L., Jagtap P.D., Janech M.G., Kinkade D.B., Leary D.H., McIlvin M.R., Moore E.K., Morris R.M., Neely B.A., Nunn B.L., Saunders J.K., Shepherd A.I., Symmonds N.I., Walsh D.A. J Proteome Res. (2019) 18, pp 11461-1476.
255. "Exploiting the Dynamic Relationship between Peptide Separation Quality and Peptide Co-isolation in a Multiple-Peptide Matches-per-Spectrum Approach Offers a Strategy to Optimize Bottom-Up Proteomics Throughput and Depth," Solis, M.I.V.; Giannone, R.J.; Hettich, R.L.; Abraham, P.E. Anal. Chem. (2019), 91, 7273-7279. DOI: 10.1021/acs.analchem.9b00819.
256. "Targeted detection of *Dehalococcoides mccartyi* microbial protein biomarkers as indicators of reductive dechlorination activity in contaminated groundwater," Solis M.I.V., Abraham P.E., Chourey K., Swift C.M., Löffler F.E., Hettich R.L. Sci Rep. (2019) Jul 22;9(1):10604. doi: 10.1038/s41598-019-46901-6.
257. "Effects of microbiota-directed foods in gnotobiotic animals and undernourished children," Gehrig J.L., Venkatesh S., Chang H.W., Hibberd M.C., Kung V.L., Cheng J., Chen R.Y., Subramanian S., Cowardin C.A., Meier M.F., O'Donnell D., Talcott M., Spears L.D., Semenkovich C.F., Henrissat B., Giannone R.J., Hettich R.L., Ilkayeva O., Muehlbauer M., Newgard C.B., Sawyer C., Head R.D., Rodionov D.A., Arzamasov A.A., Leyn S.A., Osterman A.L., Hossain M.I., Islam M., Choudhury N., Sarker S.A., Huq S., Mahmud I., Mostafa I., Mahfuz M., Barratt M.J., Ahmed T., Gordon J.I. Science. (2019) Jul 12;365(6449). pii: eaau4732. doi: 10.1126/science.aau 4732.
258. "One-time nitrogen fertilization shifts switchgrass soil microbiomes within a context of larger spatial and temporal variation," Chen H., Yang Z.K., Yip D., Morris R.H., Lebreux S.J., Cregger M.A., Klingeman D.M., Hui D., Hettich R.L., Wilhelm S.W., Wang G., Löffler F.E., Schadt C.W. PLoS One. (2019) Jun 18;14(6):e0211310. doi: 10.1371/journal.pone.0211310. eCollection 2019.
259. "Interspecies Competition Impacts Targeted Manipulation of Human Gut Bacteria by Fiber-Derived Glycans," Patnode, ML; Beller, ZW; Han, ND; Cheng, JY; Peters, SL; Terrapon, N; Henrissat, B; Le Gall, S ; Saulnier, L; Hayashi, DK; Meynier, A; Vinoy, S; Giannone, RJ; Hettich, RL; Gordon, JI. Cell (2019), Volume: 179 Issue: 1 Pages: 59- DOI: 10.1016/j.cell.2019.08.011.
260. "Bioremediation of a Common Product of Food Processing by a Human Gut Bacterium." Wolf, AR; Wesener, DA; Cheng, JY; Houtston-Ludlam, AN; Beller, ZW; Hibberd, MC; Giannone, RJ; Peters, SL; Hettich, RL; Leyn, SA; Rodionov, DA; Osterman, AL; Gordon, JI. Cell Host & Microbe, (2019), Volume: 26 Issue: 4 Pages: 463-+ DOI: 10.1016/j.chom.2019.09.001.
261. "Surface Water Microbial Community Response to the Biocide 2,2-Dibromo-3-Nitrilopropionamide, Used in Unconventional Oil and Gas Extraction," Campa, MF; Techtmann, SM; Ladd, MP; Yan, J; Patterson, M; Amaral, AGD; Carter, KE; Ulrich, N; Grant, CJ; Hettich, RL; Lamendella, R; Hazen, TC. Appl. Environ. Microbiol. (2019), Volume: 85 Issue: 21 Article Number: e01336-19.
262. "Impact of Fixed Nitrogen Availability on *Dehalococcoides mccartyi* Reductive Dechlorination Activity," Kaya D, Kjellerup BV, Chourey K, Hettich RL, Taggart DM, Löffler FE. Environ. Sci. Tech., (2019), 53, 14548-14558.
263. "Comparing DNA, RNA and protein levels for measuring microbial dynamics in soil microcosms amended with nitrogen fertilizer," Orellana LH; Hatt JK; Iyer R; Chourey K; Hettich RL; Spain JC; Yang WH; Chee-Sanford JC; Sanford RA; Löffler FE; Konstantinidis KT. Sci Rep. (2019) Volume: 9 Article Number: 17630
264. "Integrated Proteomics and Lipidomics Reveal That the Swarming Motility of *Paenibacillus polymyxa* Is Characterized by Phospholipid Modification, Surfactant Deployment, and Flagellar Specialization Relative to Swimming Motility," Poudel S; Giannone RJ; Farmer AT; Campagna SR; Bible AN; Morrell-Falvey JL;

Principal Investigator/Program Director (Last, First, Middle): Hettich, Robert L.
Elkins JG; Hettich RL. Front Microbiol. (2019) Volume: 10 Article Number: 2594.

265. "Genome-Resolved Proteomic Stable Isotope Probing of Soil Microbial Communities Using ^{13}C and ^{13}C -Methanol," Li Z; Yao Q; Guo X; Crits-Christoph A; Mayes MA; Iv WJH; Lebeis SL; Banfield JF; Hurst GB; Hettich RL; Pan C. Front Microbiol. (2019), Volume: 10, Article Number: 2706.
266. "Outer membrane vesicles catabolize lignin-derived aromatic compounds in *Pseudomonas putida* KT2440," Davinia Salvachúa, Allison Z. Werner, Isabel Pardo, Martyna Michalska, Brenna A. Black, Bryon S. Donohoe, Stephan Haugen, Rui Katahira, Sandra Notonier, Kelsey J. Ramirez, Samuel O. Purvine, Erika M. Zink, Paul E. Abraham, Richard J. Giannone, Suresh Poudel, Phillip Laible, Robert L. Hettich, Gregg T. Beckham, Proc. Natl. Acad. Sci., (2020), 117, 9302-9310.
267. "Active sulfur cycling in the terrestrial deep subsurface" Emma Bell, Tiina Lamminmäki, Johannes Alneberg, Anders Andersson, Chen Qian, Weili Xiong, Robert Hettich, Manon Frutschi, and Rizlan Bernier-Latmani, ISME J. (2020), in press, Paper #ISMEJ-19-01155AR2.
268. "Impact of Fatty-Acid Labeling of *Bacillus subtilis* Membranes on the Cellular Lipidome and Proteome," Jonathan D. Nickels, Suresh Poudel, Sneha Chatterjee, Abigail T. Farmer, Destini Cordner, Shawn R. Campagna, Richard J. Giannone, Robert L. Hettich, Dean A. Myles, Robert F. Standaert, John Katsaras, James G. Elkins. . Front Microbiol. (2020), 11 Article Number: 914, DOI: 10.3389/fmicb.2020.00914
269. "Gene targets for engineering osmotolerance in *Caldicellulosiruptor bescii*," Sander, KB.; Chung, D; Klingeman, DM; Giannone, RJ; Rodriguez, M; Whitham, J; Hettich, RL; Davison, BH; Westpheling, J; Brown, SD, Biotech. For Biofuels, (2019), 13, DOI: 10.1186/s13068-020-01690-3asfd
270. "Peeling back the layers of crassulacean acid metabolism: functional differentiation between *Kalanchoe fedtschenkoi* epidermis and mesophyll proteomes" Abraham, P.E; Castano, N.H.; Cowan-Turner, D.; Barnes, J.; Poudel, S.; Hettich, R.; Flutsch, S.; Santelia, D.; Borland, A.M. Plant J. (2020), DOI: 10.1111/tpj.14757, Early access iconEarly Access: APR 2020
271. "Characterizing Intracellular Proteomes for Microbes: An Experimental Approach Using Label-Free Protein Quantitation," Paul E Abraham and Robert L Hettich Methods Mol Biol. (2020); 2096:81-87. doi: 10.1007/978-1-0716-0195-2_7.
272. "Biogeography of microbial bile acid transformations along the murine gut," Solenne Marion, Lyne Desharnais, Nicolas Studer, Yuan Dong, Matheus D Notter, Suresh Poudel, Laure Menin, Andrew Janowczyk, Robert L Hettich, Siegfried Hapfelmeier, Rizlan Bernier-Latmani, J Lipid Res. (2020) Jul 13;jlr.RA120001021. doi: 10.1194/jlr.RA120001021. Online ahead of print.
273. "Integrating state-of-the-art tandem mass spectrometry with de novo-assisted database peptide sequencing as a viable strategy for the discovery of peptide proteolytic cleavage products (PCPs) in plant-microbe interactions," Manuel Ivan Villalobos Solis, Suresh Poudel, Clemence Bonnot, Him Kumar Shrestha, Robert Lloyd Hettich, Claire Veneault-Fourrey, Francis Martin, Paul Edward Abraham, Mol Plant Microbe Interact. (2020) Jun 29. doi: 10.1094/MPMI-04-20-0082-TA. Online ahead of print.
274. "A carotenoid-deficient mutant of the plant-associated microbe *Pantoea* sp. YR343 displays an altered membrane proteome, Sushmitha Vijaya Kumar, Paul E. Abraham, Gregory B. Hurst, Karuna Chourey, Amber N. Bible, Robert L. Hettich, Mitchel J. Doktycz, Jennifer L. Morrell-Falvey, Sci. Reports, (2020), in press.
275. "A nitrogenase-like enzyme system catalyzes methionine, ethylene, and methane biogenesis," Justin A. North, Adrienne B. Narrowe, Weili Xiong, Kathryn M. Byerly, Guanqi Zhao, Sarah J. Young, Srividya Murali, John A. Wildenthal, William R. Cannon, Kelly C. Wrighton, Robert L. Hettich, and F. Robert Tabita, Science (2020), in press.

276. “Structural and proteomic studies of the *Aureococcus anophagefferens* Virus demonstrate a global distribution of virus-encoded carbohydrate processing, Eric R. Gann, Yuejiao Xian, Paul E. Abraham, Robert L. Hettich, Todd B. Reynolds, Chuan Xiao, and Steven W. Wilhelm, *Front. Microbiol.*, (2020), in press.

Academic Responsibilities as adjunct faculty member at the University of Tennessee - Knoxville

Dissertation Major Advisor - (completed degrees):

- **Dr. Joshua Sharp**, “Development of Hydroxyl Free Radical Chemistry for the Surface Mapping of Proteins,” Ph.D. awarded October 2003. (*academic faculty member at Univ. of Mississippi*)
- **Dr. Tomoaki Uchiki**, “Structure-Function and Regulation of Ribonucleotide Reductase Inhibitor, Sml1,” Ph.D. awarded December 2004. (*research staff position in Japan*)
- **Dr. Nathan VerBerkmoes**, “Mass Spectrometry-Based Proteomics for Studying Microbial Physiology from Isolates to Communities,” Ph.D. awarded May 2005 (*academic faculty member at UT-El Paso*)
- **Dr. Heather Connelly**, “Integrated Computational and Experimental Platform for Characterizing Protein Isoforms and PTMs in Microbial Systems by Top-Down FT-ICR Mass Spectrometry,” Ph.D. awarded August 2006. (*staff member at Hospira, North Carolina*)
- **Dr. Chongle Pan**, “An Integrated Experimental and Computational Approach to Proteomics: Scaling from High Resolution Qualitative Analysis to Quantitative Measurements with Confidence Evaluation,” Ph.D. awarded December 2006 (*staff member at ORNL*)
- **Ms. Demet Ataman**, failed comprehensive exam twice; she chose to leave GST program.
- **Ms. Kanan Vyas**, “The Integration of Oxidative Surface Mapping and Molecular Dynamics Simulation Techniques as a Strategy for Studying Protein Conformational Change,” M.S. awarded August 2006. (*employed at a biotech company in San Francisco area*)
- **Dr. Melissa Thompson**, “Integrating Mass Spectrometry Based Proteomics and Bioinformatics Technologies for the Molecular Level Characterization of *Shewanella Oneidensis* To Chromate Exposure,” Ph.D. awarded November 2007. (*staff member at Pfizer, St. Louis, MO*)
- **Dr. Carlee McClintock**, “Development of an Electrochemical Technique for Oxidative Surface Mapping to Investigate Solution-Phase Protein Dynamics with High Performance Mass Spectrometry and Advanced Informatics,” Ph.D. awarded Dec. 2009. (*staff position at Pain Consultants research clinic, Knoxville, TN*)
- **Dr. Brian Erickson**, “Integrating Mass Spectrometric and Computational Technologies for the Characterization of Extracellular Proteins in a Natural Microbial Community,” Ph.D. awarded Oct. 2010 (*post-doc at Harvard Medical School*)
- **Dr. Andrew Dykstra**, “Advanced Techniques in Mass Spectrometry for Qualitative and Quantitative Protein Characterization,” Ph.D. awarded May 2011 (*research staff member at Amgen, New Hampshire*)
- **Dr. Alison Russell**, “Characterization of the Human Host Gut Microbiome with an Integrated Genomics/Proteomics Approach,” Ph.D. awarded Oct. 2011 (*post-doc at Harvard Medical School*)
- **Dr. Adriane Lochner**, “Proteomic characterization of the cellulolytic enzyme system expressed by the extremely thermophilic bacteria *Caldicellulosiruptor* spp.,” Ph.D. awarded Mar. 2012 (Germany; *currently instructor at Univ. of Hamburg*)
- **Dr. Jacque Young**, “Functional Characterization of Microbial Symbiotic Associations by Metaproteomics,” Ph.D. awarded Nov. 2012 (*post-doctoral position at Univ. of Penn.*)
- **Dr. Paul Abraham**, “Development and application of mass-spectrometry-based proteomics to generate and navigate the proteomes of the genus *Populus*,” Ph.D. awarded April, 2013 (*staff position at ORNL*)
- **Dr. Rachel Adams**, “Development and integration of informatics tools for qualitative and quantitative characterization of proteomic datasets generated by tandem mass spectrometry, Ph.D. awarded May, 2013 (*informatics consultant*)

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- **Mr. Adam Martin**, “Demonstration of a targeted proteome characterization approach for examining specific metabolic pathways in complex bacterial systems,” M.S. awarded Dec. 2013 (*currently instructor at Maryville College*).
- **Dr. Ritin Sharma**, “Development of an experimental and computational platform for enhanced characterization of modified peptides and proteins in environmental proteomics,” Ph.D. awarded April 2014 (*post-doc at Moffitt Cancer Center, Florida*)
- **Dr. Zhou Li**, “Quantitative Characterization of Proteins and Post-Translational Modifications in Complex Proteomes Using High-Resolution Mass Spectrometry-Based Proteomics,” Ph.D. awarded April 2014 (*currently post-doc at ORNL*)
- **Dr. Xiaoxin Liu**, “Comparative Proteomics Reveals Core vs. Unique Molecular Signatures for Dissimilatory Metal Reducing Bacteria Grown with Various Electron Acceptors,” Ph.D. awarded Aug. 2014 (*currently in nursing school*)
- **Dr. Weili Xiong**, “Characterizing Early-life Microbiome Functionality in Premature Infant Gut by a Metaproteomics Approach,” Ph.D. awarded Nov. 2015. (*currently post-doc at FDA*)
- **Dr. Chen Qian**, “Development of MS-based proteomics approaches to examine metabolic pathways and protein:protein interactions in microbial systems,” Ph.D. awarded July 2017. (*currently post-doc in Baltimore*)
- **Dr. Ramsunder (Sarvesh) Iyer**, “Bioinformatic and Experimental Approaches for Deeper Metaproteomic Characterization of Complex Environmental Samples,” Ph.D. awarded August 2017. (*currently post-doc at Northwestern Univ.*)
- **Dr. Suresh Poudel**, “Proteome characterizations of microbial systems using MS-based experimental and informatics approaches to examine key metabolic pathways, proteins of unknown function, and phenotypic adaptation,” awarded July 2018 (*currently post-doc at ORNL*)
- **Dr. Mallory Ladd**, “Development and application of an untargeted exometabolomics approach to identify biogeochemical hotspots of dissolved organic matter vulnerability in Arctic soils,” awarded November 2018. (*intern position Naval Research lab, Wash. D.C.*).

Dissertation Major Advisor for following graduate students (current):

- Mr. Manuel Ivan Villalobos-Solis, graduate student in GST-Ph.D. program
- Mr. Alfredo Blakeley-Ruiz, graduate student in GST-Ph.D. program
- Mr. David Reeves, graduate student in the ORNL-UTK Bredesen program.
- Mr. Alex Cope, graduate student in GST-Ph.D. program.
- Ms. Payal Chirania, graduate student in GST-Ph.D. program.
- Ms. Samantha Peters, graduate student in GST-Ph.D. program.
- Mr. Him Shrestha, graduate student in GST-Ph.D. program.
- Mr. Matthew Keller, graduate student in GST-Ph.D. program.
- Ms. Manasa Appidi, graduate student in GST-Ph.D. program.

Academic teaching:

- I am the lead instructor for the GST-II (LS 521) class on Analytical Technologies, and teach 10 MS lectures each spring.
- Every other fall, I am the lead instructor for an advanced graduate level class (LS 695) on Biological Mass Spectrometry. (*this class has been held every other year for the past 12 years*)
- Every semester, I teach a mass spectrometry-based journal discussion class at ORNL.

Expertise Summary

Dr. Robert Hettich is a distinguished research scientist and leader of the Bioanalytical Mass Spectrometry Group of the Biosciences Division at Oak Ridge National Laboratory and a joint faculty member in the Microbiology Department at the University of Tennessee. He has over 34 years of experience in biological mass spectrometry, with a particular focus on high performance mass spectrometry. His research interests involve the development and application of advanced mass spectrometry technology for characterizing complex biological mixtures, such as

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microbial and plant proteomes. His group has pioneered the technology of metaproteomics, which is the characterization of the extensive protein inventory in natural microbial communities. His work in this arena spans from environmental microbiology (i.e. microbial communities in soil and groundwater ecosystems) to bioenergy (engineering of microbial solubilization of cellulosic biomass to generate biofuels and bioproducts) and finally to the human microbiome (characterizing the microbial connections between human health and disease). His group is active in development of both experimental and computational informatic methodologies for proteome research. He has authored more than 275 publications (h-index = 67 Google Scholar), and is active in mentoring graduate students, as well as teaching analytical technology and advanced biological mass spectrometry classes. He is an associate editor for Microbiome, a member of the editorial advisory boards of Mass Spectrometry Reviews, BMC Genomics, and Journal of Integrated Omics, and is an active review member on several NIH review study sections, in particular for NIH-GM and NIH-NCI-IMAT.