

Matthew W. Fields – Biographical Sketch

Professor, Microbiology and Immunology, Montana State University, Bozeman, MT
Director, Center for Biofilm Engineering, Montana State University
Research Fellow, National Center for Genome Resources, Santa Fe, NM

(a) Professional Preparation

Western Kentucky University	Biology/Chemistry	B.S., 1993
Mississippi State University	Biological Sciences	M.S., 1995
Cornell University	Microbiology	Ph.D., 2001
Oak Ridge National Laboratory	Environmental Sci.	Postdoc, 2001

(b) Appointments

Director, Center for Biofilm Engineering	2015-
Interim Department Head, Microbiology & Immunology	2013-2014
Professor, Department of Microbiology and Immunology	2015-
Adjunct Research Fellow, National Center for Genome Resources, Santa Fe, NM	2012-
Associate Professor, Department of Microbiology, CBE, Montana State University	2011-
Assistant Professor, Department of Microbiology, CBE, MSU	2007-2010
Assistant Professor, Department of Microbiology, Miami University	2003-2006
Research Staff Scientist, Environmental Sciences Division, ORNL	2001-2003

(c) Publications – 5 (105 total) most closely related to proposed work (h index=42)

Zelaya, A.J., A.E. Parker, K.L. Bailey, P. Zhang, J. Van Nostrand, N. Daliang, D.A. Elias, J. Zhou, T.C. Hazen, A.P. Arkin, and **M.W. Fields**. 2019. High spatiotemporal variability of bacterial diversity over short time scales with unique hydrochemical associations within a shallow aquifer. *Water Res.* 164:114917

Krantz, G.P., K. Lucas, E.L. -Wunderlich, L.T. Hoang, R. Avci, G. Siuzdak, and **M.W. Fields**. 2019. Bulk phase resource ratio alters carbon steel corrosion rates and endogenously-produced extracellular electron transfer mediators in a sulfate-reducing biofilm. *Biofouling* (in press)

Schweitzer, H., D. Ritter, J. McIntosh, E.P. Barnhart, A.B. Cunningham, D. Vinson, W. Orem, and **M.W. Fields**. 2019. Changes in microbial communities and associated water and gas geochemistry across redox gradients in coal beds: Powder River Basin, USA. *Geochim. Cosmochim. Acta* 245: 495-513

Smith, H.J., A.J. Zelaya, K.B. De León, R. Chakraborty, D.A. Elias, T.C. Hazen, A.P. Arkin, A.B. Cunningham and **M.W. Fields**. 2018. Impact of hydrologic boundaries on microbial planktonic and biofilm communities in shallow terrestrial subsurface environments. *FEMS Microbiol. Ecol.* 94:fify191

He, Z., P. Zhang, L. Wu, A. Rocha, Q. Tu, Z. Shi, B. Wu, Y. Qin, J. Wang, Y.-Q. Yun, D. Curtis, D. Ning, J.D. Van Nostrand, L. Wu, Yunfeng Yang, Dwayne Elias, D. Watson, M. Adams, **M.W. Fields**, E. Alm, T.C. Hazen, P. Adams, A.P. Arkin, and J. Zhou. 2018. Microbial functional gene diversity predicts groundwater contamination and ecosystem functioning. *mBio* 9:e02435-17

Additional Relevant Publications

McKay, L.J., M. Dlakić, **M.W. Fields**, T.O. Delmont, A. Murat Eren, Z.J. Jay, K.B. Klingelsmith, D. Rusch, and W.P. Inskeep. 2019. Co-occurring genomic capacity for anaerobic methane metabolism and dissimilatory sulfite reduction discovered in the Korarchaeota. *Nat. Microbiol.* 4:614-622

McKay, L.J., R. Hatzenpichler, W.P. Inskeep, and **M.W. Fields**. 2017. Occurrence and expression of diverse methane cycling genes by multiple archaeal phyla in hot spring sediments. *Nature Sci. Reports* 7:7252

De Leon, K., G. Zane, V. Trotter, G. Krantz, A.P. Arkin, G. Butland, P. Walian, **M.W. Fields**, and J.D. Wall. 2017. Unintended laboratory-driven evolution reveals genetic requirements for biofilm formation by *Desulfovibrio vulgaris* Hildenborough. *mBio* 8:e01696-17

Brileya, K.A., J.M. Connolly, C. Downey, R. Gerlach, and **M.W. Fields**. 2013. Taxis toward hydrogen gas by *Methanococcus maripaludis*. *Nature Sci. Rep.* 3:3140.

Momeni, B., K.A. Brileya, **M.W. Fields**, and W. Shou. 2013. Strong inter-population cooperation leads to partner intermixing in microbial communities. *eLife*. 2:e00230

(d) Synergistic Activities

Editorial Board, <i>Biofilms</i> (Elsevier)	2019
Organizing Committee, Biofilms8	2018
DOE Panel Reviews	2015, 2016, 2017, 2018
Editorial Board for <i>Applied and Environmental Microbiology</i>	2006-2014
Specialty Editor; <i>Frontiers in Microbiology</i>	2011-2016