# Matthew W. Fields

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<b>Education</b>	
1996-2000	Ph.D. Department of Microbiology, Cornell University
	Major: Microbiology
	Minors: Biochemistry, Agricultural/Biological Engineering
1993-95	M.S. Biological Sciences, Mississippi State University
	Major: Microbiology Minor: Biochemistry
1989-93	B.S. Biology, Western Kentucky University
	Major: Biology/Chemistry (Honors Program)
Professional Experi	ence

Professional	Experience
	Research Staff Scientist, Oak Ridge National Laboratory
-	Assess microbial ecology of contaminated sites using molecular- and
	culture-based techniques, isolation of metal-reducing, denitrifying, and
	heavy-metal resistant microbes, development of functional, community,
	and metagenomic DNA microarrays, physiological characterization of
	bacterial isolates
2000-2001	Postdoctoral Research Associate, Oak Ridge Associated Universities
	Coordinate microbial characterization of DOE field research site at
	ORNL, assess microbial ecology of contaminated sites, assist in
	development of functional and community DNA microarrays
1996-2000	Graduate Research Assistant, Department of Microbiology, Cornell
	University, conducted and initiated independent research combining
	microbial physiology and molecular biology, advised and coordinated
	undergraduate research, assisted in manuscript reviews and grant writing.
	Researched the role and regulation of polysaccharide hydrolases from
1000	anaerobic rumen bacteria  Tanahira Aggistant Camall University Bacterial Physiology
1999	<b>Teaching Assistant</b> , Cornell University, <b>Bacterial Physiology Laboratory</b>
1997-99	Instructor, Cornell University, co-developed microbiology course
1777-77	Instructor, Cornell University, Undergraduate Microbiology
	Instructor, Cornell University, Microbiology Ambassador Program
1993-95	Graduate Research Assistant, Department of Biological Sciences,
1775-75	Mississippi State University, conducted scientific research dealing with
	gastrointestinal bacteria, assisted undergraduate researchers
1993-95	Instructor, Mississippi State University, Undergraduate Microbiology
	Laboratory and Undergraduate Botany Laboratory
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- Microbiological Consultant, Grand Gulf Nuclear Power Station, Port Gibson, Mississippi

  1992 02 Undowgrodusta Honors Program research Western Kentucky
- 1992-93 **Undergraduate Honors Program research**, Western Kentucky University
- 1991-93 **Undergraduate Teaching Assistant and Laboratory Technician**, Biology Department, Western Kentucky University

### **Manuscripts**

- **Fields, M.W.** and J.B. Russell. 2001. Purification and characterization of a novel glucomannokinase of *Prevotella bryantii*  $B_14$ , and its possible role in  $\beta$ -glucanase expression. Microbiol. 147:1035-1043
- Jarvis, G.N., **M.W. Fields**, D.A. Adamovich, C.E. Arthurs, and J.B. Russell. 2001. The mechanism of carbonate killing of *Escherichia coli*. Letts. Appl. Microbiol. 33:196-200
- **Fields, M.W.**, S. Mallik, and J.B. Russell. 2000. *Fibrobacter succinogenes* S85 ferments cellulose just as fast as cellobiose until limited by cellulose surface area. Appl. Microbiol. and Biotech. 54:570-574
- **Fields, M.W.** and J.B. Russell. 2000. Alternative pathways of glucose transport in *Prevotella bryantii* B<sub>1</sub>4. FEMS Microbiol. Letts. 183:137-142
- **Fields, M.W.** and J.B. Russell. 1998. The role of ruminal carboxymethylcellulases in the degradation of β-glucans from cereal grains. FEMS Microbiol. Ecol. 27:261-268
- **Fields, M.W.**, D.B. Wilson, and J.B. Russell. 1997. A mutant of *Prevotella ruminicola*  $B_14$  deficient in  $\beta$ -1,4-endoglucanase and mannanase activities. FEMS Microbiol. Letts. 154:9-15
- Gardner, R.G., J.E. Wells, **M.W. Fields**, D.B. Wilson, and J.B. Russell. 1997. A *Prevotella ruminicola* B<sub>1</sub>4 operon encoding extracellular polysaccharide hydrolases. Curr. Microbiol. 35:274-277
- **Fields, M.W.**, P.E. Ryals, and K.L. Anderson. 1997. Polysaccharide-inducible outer membrane proteins of *Bacteroides xylanolyticus* X5-1. Anaerobe 3:43-48
- **Fields, M.W**. and J.B. Russell. 2001. The glucomannokinase of *Prevotella bryantii* B<sub>1</sub>4: sequence homology with the regulatory glucokinases of gram-positive microorganisms. J. Mol. Microbiol. Biotech. (submitted)
- Beliaev, A.S., D.K. Thompson, **M.W. Fields**, L. Wu, D.P. Lies, K.H. Nealson, and J. Zhou. 2001. Microarray expression profiling in *Shewanella oneidensis* MR-1 indicates involvement of *etrA* in global gene regulation. J. Bac. (submitted)

- Yan, T., **M.W. Fields**, L. Wu, S. Carroll, J.M. Teidje, and J. Zhou. 2001. The diversity of nitrite reductase genes (*nir*S and *nir*K) from nitrate- and uranium-contaminated groundwater. Appl. Environ. Microbiol. (submitted)
- **Fields, M.W.**, S.L. Carroll, and J. Zhou 2001. Denitrifying bacteria isolated from nitrate- and uranium-contaminated waste sites and the ability to denitrify processed groundwater. Environ. Microbiol. (in preparation)
- **Fields, M.W.**, T. Yan, S.L. Carroll, and J. Zhou. 2001. Bacterial community composition from groundwater contaminated with different levels of nitrate, uranium, and heavy metals. Appl. Environ. Microbiol. (in preparation)

#### **Technical Reports**

**Fields, M.W.** 1999. Accurate quantitation of RNA for hybridization blots using the VersaFluor fluorometer. Tech Note 2436, US/EG Rev A, Bio-Rad, Inc.

### **Grants/Support**:

- PI Community-Wide Analysis of Unique Sequences and Functions from Uncultured Microorganisms using DNA Microarrays, LDRD-ORNL (funded)
- Co-PI Integrated analysis of protein complexes and regulatory networks involved in anaerobic energy metabolism of *Shewanella oneidensis* MR-1, LAB 01-20, DOE Microbial Cell Project (funded)

#### **Abstracts**:

- 2001 9th International Conference on Microbial Genomes, Gatlinburg, TN The Diversity of Nitrite Reductase Genes (nirS and nirK) from Bacterial Communities in Groundwater Contaminated with Nitrate, Uranium and Heavy Metals
- 2001 Department of Energy, 4<sup>th</sup> NABIR Workshop, Warrenton, VA Microbial characterization of mixed waste sites and the development of DNA microarray methodology
- 2001 Department of Energy, 4<sup>th</sup> NABIR Workshop, Warrenton, VA Field-scale evaluation of uranium-contaminated groundwater at the NABIR field research site at Oak Ridge, TN
- 2001 Department of Energy, 4<sup>th</sup> NABIR Workshop, Warrenton, VA Understanding the roles of spatial isolation and carbon in microbial community structure, dynamics and activity for bioremediation
- 2000 American Society for Microbiology, 100 <sup>th</sup> General Meeting, Los Angeles, CA The regulation of β-glucanase expression by the ruminal bacterium, *Prevotella bryantii* B<sub>1</sub>4

- 1998 International Symposium on Microbial Ecology, 8 th Symposium, Halifax, Canada The role of ruminal carboxymethylcellulases in the degradation of glucans from cereal grains
- 1998 American Society for Microbiology, 98<sup>th</sup> General Meeting, Atlanta, GA Microbiology Ambassadors: An undergraduate teaching experience in high school classrooms
- 1997 American Society for Microbiology, 97<sup>th</sup> General Meeting, Miami Beach, FL Properties of a *Prevotella ruminicola* B<sub>1</sub>4 mutant deficient in CMCase and mannanase activities
- 1995 The Rumen Conference, 23<sup>rd</sup> Meeting, Chicago, IL Conjugal transfer of transposon Tn1545 into *Eubacterium cellulosolvens* 5494
- 1995 American Society for Microbiology, South Central Branch, Little Rock, AR
  Presence of polysaccharide inducible proteins in the outer membrane of
  Bacteroides xylanolyticus X5-1
- 1995 American Society for Microbiology, 95<sup>th</sup> General Meeting, Washington, D.C. SDS-PAGE shows polysaccharide inducible membrane proteins from *Bacteroides xylanolyticus* X5-1 and *Prevotella ruminicola* 23
- 1994 American Society for Microbiology, South Central Branch, Shreveport, LA Detection of inducible proteins in *Bacteroides thetaiotaomicron*, *Bacteroides xylanolyticus*, *Prevotella ruminicola* with SDS-PAGE