

Curriculum vitae Martin Ackermann

May 2023

Career

2023 - present	Director, Eawag, Switzerland
2022 - present	Full Professor for Microbial Systems Ecology, EPFL, Switzerland
2015 - present	Full Professor for Microbial Systems Ecology, ETH Zurich, Switzerland
2012 - 2022	Head of the Department of Environmental Microbiology, Eawag, Switzerland
2008 - 2015	Associate Professor for Molecular Microbial Ecology, ETH Zurich, Switzerland
2006 - 2008	SNF Professor for Microbial Evolution, ETH Zurich, Switzerland
2004 - 2006	Senior Scientist, ETH Zurich, Switzerland, in the group of Prof. S. Bonhoeffer
2002 - 2004	Postdoctoral Researcher, UC San Diego, USA, in the group of Prof. L. Chao
2002	PhD in Biology , University of Basel, Switzerland, with Profs. U. Jenal and S. Stearns

Research Group

Since 2023: co-lead of the research group with Dr. Olga Schubert

Microbial Systems Ecology group, Eawag, ETH and EPFL: Located at the Eawag Department of Environmental Microbiology, and the ETH Department of Environmental Systems Science.

Supervision of students and postdocs since 2006

- 14 PhD theses (6 ongoing) as direct scientific supervisor
- 26 Postdocs and scientists (8 ongoing)

All former group members have successfully completed their PhD or postdoctoral studies.
22 out of 26 PhD students and postdocs under my direct supervision stayed in academic research.
14 former group members are now principal investigators in academia or have been offered PI positions.

Research Interests

Microbial communities: assembly, dynamics and emergent properties of microbial communities.

Bacterial interactions: mutualistic and antagonistic interactions between bacterial cells; the impact of interactions on processes in natural systems.

Bacterial individuality: molecular mechanisms and adaptive significance of phenotypic heterogeneity in clonal populations; relevance of phenotypic heterogeneity in natural environments; bacterial behaviour in dynamic environments.

Bacteria under adverse conditions: genetic and phenotypic responses to antibiotics and other stressors at the single-cell level.

Invited Seminars and Talks

More than 120 invited talks since 2009.

Selected examples 2014-2022:

2022 Weizmann Institute, Rehovot, Israel

Kavli Institute of Physics, UC Santa Barbara, Santa Barbara CA, USA

Biozentrum of the University of Basel, Basel, CH

Belgian Society for Microbiology annual symposium, Brussels, BE

Anniversary Symposium of the Max Planck Institute for Marine Microbiology, Bremen, DE

Department of Biology, Humboldt University, Berlin, DE

Microbial Community Meeting, Ohio State University, USA (online)

2021 (all seminars before August 2021 cancelled because of work in the Swiss National COVID-19

Science Task Force see below)

PBEE Seminar, Emory University, USA (online)

Center for Biofilm Engineering, Montana State University, USA (online)

2020 (all seminars cancelled because of work in the Swiss National COVID-19 Science Task Force,
see below)

2019 Institute of Physics, London, UK

Department of Biology, Leiden Univ., NL

FEMS Conference, Glasgow, UK

Max Planck Institute for Marine Microbiology, Bremen, DE

SPP1617 International Meeting, Schloss Hohenkammer, DE

University of Luxembourg, Luxembourg, LU

2018 Weizmann Institute, Rehovot, IL

Gordon Conference on Marine Microbiology, Lucca, IT

Lake Arrowhead Microbial Genomics Meeting, Lake Arrowhead, CA, USA

Meeting of the Royal Society on Single-cell Ecology, London, UK

Department of Biology, KU Leuven, BE

2017 Northeastern University, Department of Biology, Boston, MA, USA

Gordon Conference on Molecular Mechanisms of Evolution, Easton, MA, USA

FEMS Conference, Valencia, ES

2016 Yale University, Dep. of Molecular, Cellular and Developmental Biology, New Haven, CO, USA

MIT, Environmental Sciences Seminar, Boston, MA, USA

Caltech, CEMI symposium, Pasadena, CA, USA

UC San Diego, qBio Seminar, San Diego, CA, USA

Gordon conference on microbial stress response, South Hadley, MA

March Meeting of the American Physical Society, Baltimore, MD, USA

2015 California Institute of Technology, Pasadena, CA, USA

University of British Columbia, Biodiversity Seminar, Vancouver, CA

University of Groningen, Groningen Lectures in Theoretical Biology, Groningen, NL

IOP Institute of Physics, The Physics of Microorganisms, London, UK

ESF conference on Bacterial Networks, San Feliu, ES

2014 Meeting of the Joint Genome Institute, Walnut Creek, Ca, USA

International Conference on Microbial Communication, Jena, DE

SPP1617 International conference on Phenotypic Heterogeneity, Kloster Irsee, DE

International Society of Microbial Ecology Conference, Seoul, South Korea

Isaac Newton Institute for Mathematical Sciences, University of Cambridge, UK

Main Recent Grants

- 2021 Simons Foundation, 10 PI project on 'Principles of Microbial Ecosystems' (lead:
Roman Stocker and Otto Cordero), 1.2 million USD for 5 years.

2020	SNF NCCR, 20 PI project on 'Microbiomes' (lead: Jan-Roelof van der Meer and Julia Vorholt), 677'000 SFR. for MA for 4 years.
2019	SNF, research grant, single PI: SFR. 1'073'000 for 4 years.
2017	Simons Foundation, 10 PI project on 'Theory of Microbial Ecosystems' (lead: Roman Stocker and Otto Cordero), 1.4 million USD for MA for 5 years.
2016	SNF, research grant, single PI: SFR. 783'000 for 3 years.
2009-21	Host for 7 Marie-Curie Postdoctoral Fellows, 5 ETH Fellows, 2 SNF Ambizione Fellows, 1 SystemsX Fellow, 2 EMBO Fellows

Recognition and Service to the Academic Community

Member of the European Academy of Microbiology, from 2017 on.

Member of the Advisory Board for the MPI in Terristrial Microbiology, Marburg, 2016-2020.

Editorial Board member of 'Environmental Microbiology' since 2014.

Reviewer for over 50 different scientific journals (including Nature, Science, PNAS, PLOS Biology, Elife), and for 12 national and international funding agencies (including the Swiss National Science Foundation, NSF, ERC, MacArthur Foundation).

Member of more than 60 doctoral committees in Switzerland and abroad.

Public Service

August 2020 - August 2021: President of the Swiss National COVID-19 Science Task Force.

April 2020 - July 2020: Vice-president of the Swiss National COVID-19 Science Task Force.

Publications

Google Scholar: H-index 52, i10 index 93, total 10640 citations (May 2023).

<https://scholar.google.com/citations?user=kk5lwecAAAAJ&hl=en>

In addition to the publications listed below, members of my research group have published more than 40 papers since 2006 while being in the group without me as a co-author. I am only a co-author on papers from my group if I make a concrete scientific contribution.

D'Souza, G., Ebrahimi, A., Stubbensch, A., Daniels, M., Keegstra, J., Stocker, R., Cordero, O. & Ackermann, M. Cell aggregation is associated with enzyme secretion strategies in marine polysaccharide-degrading bacteria. *The ISME Journal*, 2023, 1-9.

Dal Co, A., Ackermann, M., & van Vliet, S. (2023). Spatial self-organization of metabolism in microbial systems: A matter of enzymes and chemicals. *Cell Systems*, 2023, 14(2), 98-108.

Schemm, S., Grund, D., Knutti, R., Wernli, H., Ackermann, M., & Evensen, G. Learning from weather and climate science to prepare for a future pandemic. *Proceedings of the National Academy of Sciences*, 2023, 120(4), e2209091120.

- Daniels, M., van Vliet, S., & Ackermann, M. Changes in interactions over ecological time scales influence single-cell growth dynamics in a metabolically coupled marine microbial community. *The ISME Journal*, 2023, 1-11.
- Huelsmann M, Ackermann M. Community instability in the microbial world. *Science*. 2022 Oct 7;378(6615):29-30.
- Wang M, Chen X, Liu X, Fang Y, Zheng X, Huang T, Tang YQ, Ackermann M, Nie Y, Wu XL. Even allocation of benefits stabilizes microbial community engaged in metabolic division of labor. *Cell Reports*, 2022, Sep 27;40(13):111410.
- Hemmerle, L., Maier, B. A., Bortfeld-Miller, M., Ryback, B., Gäbelein, C. G., Ackermann, M., & Vorholt, J. A. Dynamic character displacement among a pair of bacterial phyllosphere commensals in situ. *Nature communications*, 2022, 13(1), 1-14.
- van Vliet, S., Hauert, C., Fridberg, K., Ackermann, M., & Dal Co, A. Global dynamics of microbial communities emerge from local interaction rules. *PLoS computational biology*, 2022, 18(3), e1009877
- Hockenberry, A. M., Micali, G., Takacs, G., Weng, J., Hardt, W. D., and M. Ackermann. Microbiota-derived metabolites inhibit *Salmonella* virulent subpopulation development by acting on single-cell behaviors. *Proceedings of the National Academy of Sciences*, 2021. 8(3).
- van Gestel, J., Bareia, T., Tenenbaum, B., Dal Co, A., Guler, P., Aframian, N., Puyesky, S., Grinberg, I., D'Souza, G.G., Erez, Z., Ackermann, M. and A. Eldar. Short-range quorum sensing controls horizontal gene transfer at micron scale in bacterial communities. *Nature communications*, 2021. 12(1), pp.1-11.
- Nguyen, J., Fernandez, V., Pontrelli, S., Sauer, U., Ackermann, M., and R. Stocker, R. A distinct growth physiology enhances bacterial growth under rapid nutrient fluctuations. *Nature Communications*, 2021. 12(1), 1-12.
- D'Souza, G. G., Povolo, V. R., Keegstra, J. M., Stocker, R., and M. Ackermann. Nutrient complexity triggers transitions between solitary and colonial growth in bacterial populations. *The ISME Journal*, 2021. 1-13.
- Rodríguez-Verdugo, A., and M. Ackermann. Rapid evolution destabilizes species interactions in a fluctuating environment. *The ISME Journal*, 2021. 15(2), 450-460.
- Benz, F., J.S. Huisman, E. Bakkeren, J.A. Herter, T. Stadler, M. Ackermann, M. Diard, A. Egli, A.R. Hall, and W.-D. Hardt, Plasmid-and strain-specific factors drive variation in ESBL-plasmid spread in vitro and in vivo. *The ISME journal*, 2021. 15(3): p. 862-878.
- Sun, L., P. Ashcroft, M. Ackermann, and S. Bonhoeffer, Stochastic gene expression influences the selection of antibiotic resistance mutations. *Molecular biology and evolution*, 2020. 37(1): p. 58-70.
- Schreiber, F. and M. Ackermann, Environmental drivers of metabolic heterogeneity in clonal microbial populations. *Current Opinion in Biotechnology*, 2020. 62: p. 202-211.
- Moreno-Gámez, S., D.J. Kiviet, C. Vulin, S. Schlegel, K. Schlegel, G.S. van Doorn, and M. Ackermann, Wide lag time distributions break a trade-off between reproduction and survival in bacteria. *Proceedings of the National Academy of Sciences*, 2020. 117(31): p. 18729-18736.

- Gorter, F.A., M. Manhart, and M. Ackermann, Understanding the evolution of interspecies interactions in microbial communities. *Philosophical Transactions of the Royal Society B*, 2020. 375(1798): p. 20190256.
- Dal Co, A., S. van Vliet, D.J. Kiviet, S. Schlegel, and M. Ackermann, Short-range interactions govern the dynamics and functions of microbial communities. *Nature Ecology & Evolution*, 2020. 4(3): p. 366-375.
- Bosshard, L., S. Peischl, M. Ackermann, and L. Excoffier, Dissection of the mutation accumulation process during bacterial range expansions. *BMC Genomics*, 2020. 21: p. 1-11.
- van Gestel, J., M. Ackermann, and A. Wagner, Microbial life cycles link global modularity in regulation to mosaic evolution. *Nature Ecology & Evolution*, 2019.
- Schiessl, K.T., A. Ross-Gillespie, D.M. Cornforth, M. Weigert, C. Bigosch, S.P. Brown, M. Ackermann, and R. Kummerli, Individual- versus group-optimality in the production of secreted bacterial compounds. *Evolution*, 2019.
- Rodriguez-Verdugo, A., C. Vulin, and M. Ackermann, The rate of environmental fluctuations shapes ecological dynamics in a two-species microbial system. *Ecology letters*, 2019.
- Povolo, V.R. and M. Ackermann, Disseminating antibiotic resistance during treatment. *Science*, 2019. 364(6442): p. 737-738.
- Leventhal, G.E., M. Ackermann, and K.T. Schiessl, Why microbes secrete molecules to modify their environment: the case of iron-chelating siderophores. *Journal of the Royal Society Interface*, 2019. 16(150).
- Dal Co, A., S. van Vliet, and M. Ackermann, Emergent microscale gradients give rise to metabolic cross-feeding and antibiotic tolerance in clonal bacterial populations. *Philosophical Transactions of the Royal Society B*, 2019.
- Dal Co, A., M. Ackermann, and S. van Vliet, Metabolic activity affects response of single cells to a nutrient switch in structured populations. *Journal of the Royal Society Interface*, 2019.
- Bosshard, L., S. Peischl, M. Ackermann, and L. Excoffier, Mutational and selective processes involved in evolution during bacterial range expansions. *Molecular biology and evolution*, 2019.
- Balaban, N.Q., S. Helaine, K. Lewis, M. Ackermann, B. Aldridge, D.I. Andersson, M.P. Brynildsen, D. Bumann, A. Camilli, and J.J. Collins, Definitions and guidelines for research on antibiotic persistence. *Nature Reviews Microbiology*, 2019. 17(7): p. 441-448.
- Zimmermann, M., S. Escrig, G. Lavik, M.M.M. Kuypers, A. Meibom, M. Ackermann, and F. Schreiber, Substrate and electron donor limitation induce phenotypic heterogeneity in different metabolic activities in a green sulphur bacterium. *Environmental Microbiology Reports*, 2018. 10(2): p. 179-183.
- Vulin, C., N. Leimer, M. Huemer, M. Ackermann, and A.S. Zinkernagel, Prolonged bacterial lag time results in small colony variants that represent a sub-population of persisters. *Nature Communications*, 2018. 9.
- van Vliet, S., A. Dal Co, A.R. Winkler, S. Spriewald, B. Stecher, and M. Ackermann, Spatially Correlated Gene Expression in Bacterial Groups: The Role of Lineage History, Spatial Gradients, and Cell-Cell Interactions. *Cell Systems*, 2018. 6(4): p. 496-+.

- Sun, L., H.K. Alexander, B. Bogos, D.J. Kiviet, M. Ackermann, and S. Bonhoeffer, Effective polyploidy causes phenotypic delay and influences bacterial evolvability. *Plos Biology*, 2018. 16(2).
- Proctor, C.R., M.D. Besmer, T. Langenegger, K. Beck, J.-C. Walser, M. Ackermann, H. Burgmann, and F. Hammes, Phylogenetic clustering of small low nucleic acid-content bacteria across diverse freshwater ecosystems. *Isme Journal*, 2018. 12(5): p. 1344-1359.
- Louca, S., M.F. Polz, F. Mazel, M.B.N. Albright, J.A. Huber, M.I. O'Connor, M. Ackermann, A.S. Hahn, D.S. Srivastava, S.A. Crowe, M. Doebeli, and L.W. Parfrey, Function and functional redundancy in microbial systems. *Nature Ecology & Evolution*, 2018. 2(6): p. 936-943.
- Dal Co, A., C. Brannon, and M. Ackermann, Division of labor in bacteria. *eLife*, 2018. 7.
- van Vliet, S. and M. Ackermann, Stochastic gene expression: bacterial elites in chemotaxis. *Molecular Systems Biology*, 2017. 13(1).
- Schiessl, K.T., E.M.L. Janssen, S.M. Kraemer, K. McNeill, and M. Ackermann, Magnitude and Mechanism of Siderophore-Mediated Competition at Low Iron Solubility in the *Pseudomonas aeruginosa* Pyochelin System. *Frontiers in Microbiology*, 2017. 8.
- Probst-Rued, S., K. McNeill, and M. Ackermann, Thiouridine residues in tRNAs are responsible for a synergistic effect of UVA and UVB light in photoinactivation of *Escherichia coli*. *Environmental Microbiology*, 2017. 19(2): p. 434-442.
- Nikolic, N., F. Schreiber, A. Dal Co, D.J. Kiviet, T. Bergmiller, S. Littmann, M.M.M. Kuypers, and M. Ackermann, Cell-to-cell variation and specialization in sugar metabolism in clonal bacterial populations. *Plos Genetics*, 2017. 13(12).
- Mathis, R. and M. Ackermann, Asymmetric cellular memory in bacteria exposed to antibiotics. *Bmc Evolutionary Biology*, 2017. 17.
- Marchal, M., F. Goldschmidt, S.N. Derksen-Muller, S. Panke, M. Ackermann, and D.R. Johnson, A passive mutualistic interaction promotes the evolution of spatial structure within microbial populations. *Bmc Evolutionary Biology*, 2017. 17.
- Diard, M., E. Bakkeren, J.K. Cornuault, K. Moor, A. Hausmann, M.E. Sellin, C. Loverdo, A. Aertsen, M. Ackermann, M. De Paepe, E. Slack, and W.-D. Hardt, Inflammation boosts bacteriophage transfer between *Salmonella* spp. *Science*, 2017. 355(6330): p. 1211-1215.
- Bosshard, L., I. Dupanloup, O. Tenaillon, R. Bruggmann, M. Ackermann, S. Peischl, and L. Excoffier, Accumulation of deleterious mutations during bacterial range expansions. *Genetics*, 2017. 207(2): p. 669-684.
- Vizcarra, I.A., V. Hosseini, P. Kollmannsberger, S. Meier, S.S. Weber, M. Arnoldini, M. Ackermann, and V. Vogel, How type 1 fimbriae help *Escherichia coli* to evade extracellular antibiotics. *Scientific Reports*, 2016. 6.
- Stamm, C., K. Rasanen, F.J. Burdon, F. Altermatt, J. Jokela, A. Joss, M. Ackermann, and R.I.L. Eggen, Unravelling the Impacts of Micropollutants in Aquatic Ecosystems: Interdisciplinary Studies at the Interface of Large-Scale Ecology, in *Advances in Ecological Research*, Vol 55: Large-Scale Ecology: Model Systems to Global Perspectives, A.J. Dumbrell, R.L. Kordas, and G. Woodward, Editors. 2016. p. 183-223.

*Schreiber, F., S. Littmann, G. Lavik, S. Escrig, A. Meibom, M.M.M. Kuypers, and M. Ackermann, Phenotypic heterogeneity driven by nutrient limitation promotes growth in fluctuating environments. *Nature Microbiology*, 2016. 1(6).

Nelson, M.B., A.B. Chase, J.B.H. Martiny, R. Stocker, N. Jen, K. Lloyd, R.T. Oshiro, D.B. Kearns, J.P. Schneider, P.D. Ringel, M. Basler, C.A. Olson, H.E. Vuong, E.Y. Hsiao, B.R.K. Roller, M. Ackermann, C. Smillie, D. Chien, E. Alm, and A.J. Jermy, The Microbial Olympics 2016. *Nature Microbiology*, 2016. 1(8).

Mathis, R. and M. Ackermann, Response of single bacterial cells to stress gives rise to complex history dependence at the population level. *Proceedings of the National Academy of Sciences of the United States of America*, 2016. 113(15): p. 4224-4229.

Boehm, A., M. Arnoldini, T. Bergmiller, T. Roeoesli, C. Bigosch, and M. Ackermann, Genetic Manipulation of Glycogen Allocation Affects Replicative Lifespan in E-coli. *Plos Genetics*, 2016. 12(4).

Zimmermann, M., S. Escrig, T. Huebschmann, M.K. Kirf, A. Brand, R.F. Inglis, N. Musat, S. Mueller, A. Meibom, M. Ackermann, and F. Schreiber, Phenotypic heterogeneity in metabolic traits among single cells of a rare bacterial species in its natural environment quantified with a combination of flow cell sorting and NanoSIMS. *Frontiers in Microbiology*, 2015. 6.

van Vliet, S. and M. Ackermann, Bacterial Ventures into Multicellularity: Collectivism through Individuality. *Plos Biology*, 2015. 13(6).

Johnson, D.R., D.E. Helbling, T.K. Lee, J. Park, K. Fenner, H.-P.E. Kohler, and M. Ackermann, Association of Biodiversity with the Rates of Micropollutant Biotransformations among Full-Scale Wastewater Treatment Plant Communities. *Applied and Environmental Microbiology*, 2015. 81(2): p. 666-675.

Hall, A.R., D.C. Angst, K.T. Schiessl, and M. Ackermann, Costs of antibiotic resistance - separating trait effects and selective effects. *Evolutionary Applications*, 2015. 8(3): p. 261-272.

Ackermann, M. and F. Schreiber, A growing focus on bacterial individuality. *Environmental Microbiology*, 2015. 17(7): p. 2193-2195.

Ackermann, M., The usefulness of evolutionary principles: predicting the unexpected. *Environmental Microbiology Reports*, 2015. 7(1): p. 4-5.

Ackermann, M., A functional perspective on phenotypic heterogeneity in microorganisms. *Nature Reviews Microbiology*, 2015. 13(8): p. 497-508.

Ocampo, P.S., V. Lazar, B. Papp, M. Arnoldini, P.A. zur Wiesch, R. Busa-Fekete, G. Fekete, C. Pal, M. Ackermann, and S. Bonhoeffer, Antagonism between Bacteriostatic and Bactericidal Antibiotics Is Prevalent. *Antimicrobial Agents and Chemotherapy*, 2014. 58(8): p. 4573-4582.

Kuemmerli, R., K.T. Schiessl, T. Waldvogel, K. McNeill, and M. Ackermann, Habitat structure and the evolution of diffusible siderophores in bacteria. *Ecology Letters*, 2014. 17(12): p. 1536-1544.

Kaiser, P., R.R. Regoës, T. Dolowschiak, S.Y. Wotzka, J. Lengfeld, E. Slack, A.J. Grant, M. Ackermann, and W.-D. Hardt, Cecum Lymph Node Dendritic Cells Harbor Slow-Growing Bacteria Phenotypically Tolerant to Antibiotic Treatment. *Plos Biology*, 2014. 12(2).

- Fischer, B.B., M. Kwiatkowski, M. Ackermann, J. Krismer, S. Roffler, M.J.F. Suter, R.I.L. Eggen, and B. Matthews, Phenotypic plasticity influences the eco-evolutionary dynamics of a predator-prey system. *Ecology*, 2014. 95(11): p. 3080-3092.
- *Diard, M., M.E. Sellin, T. Dolowschiak, M. Arnoldini, M. Ackermann, and W.-D. Hardt, Antibiotic Treatment Selects for Cooperative Virulence of *Salmonella Typhimurium*. *Current Biology*, 2014. 24(17): p. 2000-2005.
- Bodenhausen, N., M. Bortfeld-Miller, M. Ackermann, and J.A. Vorholt, A Synthetic Community Approach Reveals Plant Genotypes Affecting the Phyllosphere Microbiota. *Plos Genetics*, 2014. 10(4).
- Blank, D., L. Wolf, M. Ackermann, and O.K. Silander, The predictability of molecular evolution during functional innovation. *Proceedings of the National Academy of Sciences of the United States of America*, 2014. 111(8): p. 3044-3049.
- Arnoldini, M., I.A. Vizcarra, R. Pena-Miller, N. Stocker, M. Diard, V. Vogel, R.E. Beardmore, W.-D. Hardt, and M. Ackermann, Bistable Expression of Virulence Genes in *Salmonella* Leads to the Formation of an Antibiotic-Tolerant Subpopulation. *Plos Biology*, 2014. 12(8).
- Nikolic, N., T. Barner, and M. Ackermann, Analysis of fluorescent reporters indicates heterogeneity in glucose uptake and utilization in clonal bacterial populations. *Bmc Microbiology*, 2013. 13.
- Mohr, W., T. Vagner, M.M.M. Kuypers, M. Ackermann, and J. LaRoche, Resolution of Conflicting Signals at the Single-Cell Level in the Regulation of Cyanobacterial Photosynthesis and Nitrogen Fixation. *Plos One*, 2013. 8(6).
- Inglis, R.F., B. Bayramoglu, O. Gillor, and M. Ackermann, The role of bacteriocins as selfish genetic elements. *Biology Letters*, 2013. 9(3).
- Diard, M., V. Garcia, L. Maier, M.N.P. Remus-Emsermann, R.R. Regoes, M. Ackermann, and W.-D. Hardt, Stabilization of cooperative virulence by the expression of an avirulent phenotype. *Nature*, 2013. 494(7437): p. 353-356.
- Ackermann, M., Microbial individuality in the natural environment. *Isme Journal*, 2013. 7(3): p. 465-467.
- Stecher, B., R. Denzler, L. Maier, F. Bernet, M.J. Sanders, D.J. Pickard, M. Barthel, A.M. Westendorf, K.A. Krogfelt, A.W. Walker, M. Ackermann, U. Dobrindt, N.R. Thomson, and W.-D. Hardt, Gut inflammation can boost horizontal gene transfer between pathogenic and commensal Enterobacteriaceae. *Proceedings of the National Academy of Sciences of the United States of America*, 2012. 109(4): p. 1269-1274.
- Silander, O.K., N. Nikolic, A. Zaslaver, A. Bren, I. Kikoin, U. Alon, and M. Ackermann, A Genome-Wide Analysis of Promoter-Mediated Phenotypic Noise in *Escherichia coli*. *Plos Genetics*, 2012. 8(1).
- Pena-Miller, R., D. Laehnemann, H. Schulenburg, M. Ackermann, and R. Beardmore, The optimal deployment of synergistic antibiotics: a control-theoretic approach. *Journal of the Royal Society Interface*, 2012. 9(75): p. 2488-2502.
- Pena-Miller, R., D. Laehnemann, H. Schulenburg, M. Ackermann, and R. Beardmore, Selecting Against Antibiotic-Resistant Pathogens: Optimal Treatments in the Presence of Commensal Bacteria. *Bulletin of Mathematical Biology*, 2012. 74(4): p. 908-934.

- Johnson, D.R., F. Goldschmidt, E.E. Lilja, and M. Ackermann, Metabolic specialization and the assembly of microbial communities. *Isme Journal*, 2012. 6(11): p. 1985-1991.
- Ispolatov, I., M. Ackermann, and M. Doebeli, Division of labour and the evolution of multicellularity. *Proceedings of the Royal Society B-Biological Sciences*, 2012. 279(1734): p. 1768-1776.
- Helbling, D.E., M. Ackermann, K. Fenner, H.-P.E. Kohler, and D.R. Johnson, The activity level of a microbial community function can be predicted from its metatranscriptome. *Isme Journal*, 2012. 6(4): p. 902-904.
- Bergmiller, T., M. Ackermann, and O.K. Silander, Patterns of Evolutionary Conservation of Essential Genes Correlate with Their Compensability. *Plos Genetics*, 2012. 8(6).
- Arnoldini, M., R. Mostowy, S. Bonhoeffer, and M. Ackermann, Evolution of Stress Response in the Face of Unreliable Environmental Signals. *Plos Computational Biology*, 2012. 8(8).
- Sturm, A., M. Heinemann, M. Arnoldini, A. Benecke, M. Ackermann, M. Benz, J. Dormann, and W.-D. Hardt, The Cost of Virulence: Retarded Growth of *Salmonella Typhimurium* Cells Expressing Type III Secretion System 1. *Plos Pathogens*, 2011. 7(7).
- Pecson, B.M., M. Ackermann, and T. Kohn, Framework for Using Quantitative PCR as a Nonculture Based Method To Estimate Virus Infectivity. *Environmental Science & Technology*, 2011. 45(6): p. 2257-2263.
- Bergmiller, T., R. Pena-Miller, A. Boehm, and M. Ackermann, Single-cell time-lapse analysis of depletion of the universally conserved essential protein YgjD. *Bmc Microbiology*, 2011. 11.
- Bergmiller, T. and M. Ackermann, Pole Age Affects Cell Size and the Timing of Cell Division in *Methylobacterium extorquens* AM1. *Journal of Bacteriology*, 2011. 193(19): p. 5216-5221.
- Boehm, A., M. Kaiser, H. Li, C. Spangler, C.A. Kasper, M. Ackermann, V. Kaever, V. Sourjik, V. Roth, and U. Jenal, Second Messenger-Mediated Adjustment of Bacterial Swimming Velocity. *Cell*, 2010. 141(1): p. 107-116.
- Silander, O.K. and M. Ackermann, The constancy of gene conservation across divergent bacterial orders. *BMC research notes*, 2009. 2: p. 2.
- Novak, M., T. Pfeiffer, M. Ackermann, and S. Bonhoeffer, Bacterial growth properties at low optical densities. *Antonie Van Leeuwenhoek International Journal of General and Molecular Microbiology*, 2009. 96(3): p. 267-274.
- Boehm, A., S. Steiner, F. Zaehringer, A. Casanova, F. Hamburger, D. Ritz, W. Keck, M. Ackermann, T. Schirmer, and U. Jenal, Second messenger signalling governs *Escherichia coli* biofilm induction upon ribosomal stress. *Molecular Microbiology*, 2009. 72(6): p. 1500-1516.
- Wanner, R.M., C. Guethlein, B. Springer, E.C. Boettger, and M. Ackermann, Stabilization of the genome of the mismatch repair deficient *Mycobacterium tuberculosis* by context-dependent codon choice. *Bmc Genomics*, 2008. 9.
- Freed, N.E., O.K. Silander, B. Stecher, A. Boehm, W.-D. Hardt, and M. Ackermann, A Simple Screen to Identify Promoters Conferring High Levels of Phenotypic Noise. *Plos Genetics*, 2008. 4(12).
- Ackermann, M., B. Stecher, N.E. Freed, P. Songhet, W.-D. Hardt, and M. Doebeli, Self-destructive cooperation mediated by phenotypic noise. *Nature*, 2008. 454(7207): p. 987-990.

CV Martin Ackermann

- Ackermann, M., A. Schauerte, S.C. Stearns, and U. Jenal, Experimental evolution of aging in a bacterium. *Bmc Evolutionary Biology*, 2007. 7.
- Ackermann, M., L. Chao, C.T. Bergstrom, and M. Doebeli, On the evolutionary origin of aging. *Aging Cell*, 2007. 6(2): p. 235-244.
- Salathe, M., M. Ackermann, and S. Bonhoeffer, The effect of multifunctionality on the rate of evolution in yeast. *Molecular Biology and Evolution*, 2006. 23(4): p. 721-722.
- Ackermann, M. and L. Chao, DNA sequences shaped by selection for stability. *Plos Genetics*, 2006. 2(2): p. 224-230.
- Ackermann, M. and M. Doebeli, Evolution of niche width and adaptive diversification. *Evolution*, 2004. 58(12): p. 2599-2612.
- Ackermann, M. and L. Chao, Evolution of cooperation: Two for one? *Current Biology*, 2004. 14(2): p. R73-R74.
- Ackermann, M., S. Stearns, and U. Jenal, Senescence in a bacterium with asymmetric division. *Science*, 2003. 300(5627): p. 1920.
- Denecker, G., S. Totemeyer, L. Mota, P. Troisfontaines, I. Lamberton, C. Youta, I. Stainier, M. Ackermann, and G. Cornelis, Effect of low- and high-virulence *Yersinia enterocolitica* strains on the inflammatory response of human umbilical vein endothelial cells. *Infection and Immunity*, 2002. 70(7): p. 3510-3520.
- Kern, S., M. Ackermann, S. Stearns, and T. Kawecki, Decline in offspring viability as a manifestation of aging in *Drosophila melanogaster*. *Evolution*, 2001. 55(9): p. 1822-1831.
- Ackermann, M., R. Bijlsma, A. James, L. Partridge, B. Zwaan, and S. Stearns, Effects of assay conditions in life history experiments with *Drosophila melanogaster*. *Journal of Evolutionary Biology*, 2001. 14(2): p. 199-209.
- Stearns, S., M. Ackermann, M. Doebeli, and M. Kaiser, Experimental evolution of aging, growth, and reproduction in fruitflies. *Proceedings of the National Academy of Sciences of the United States of America*, 2000. 97(7): p. 3309-3313.
- Stearns, S., M. Ackermann, and M. Doebeli, The experimental evolution of aging in fruitflies. *Experimental Gerontology*, 1998. 33(7-8): p. 785-792.
- Doebeli, M., A. Blarer, and M. Ackermann, Population dynamics, demographic stochasticity, and the evolution of cooperation. *Proceedings of the National Academy of Sciences of the United States of America*, 1997. 94(10): p. 5167-5171.