Sren J Srensen

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

66 18,550 350 123 h-index g-index citations papers 6.83 6.7 386 23,348 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
350	IncHI1A plasmids potentially facilitate a horizontal flow of antibiotic resistance genes to pathogens in microbial communities of urban residential sewage <i>Molecular Ecology</i> , 2022 ,	5.7	2
349	Artificial selection of stable rhizosphere microbiota leads to heritable plant phenotype changes. <i>Ecology Letters</i> , 2022 , 25, 189-201	10	0
348	Deciphering bacteria associated with a pre-parasitic stage of the root-knot nematode Meloidogyne hapla in nemato-suppressive and nemato-conducive soils. <i>Applied Soil Ecology</i> , 2022 , 172, 104344	5	1
347	Genome binning of viral entities from bulk metagenomics data <i>Nature Communications</i> , 2022 , 13, 965	17.4	3
346	Metabolic Profiling of Interspecies Interactions During Sessile Bacterial Cultivation Reveals Growth and Sporulation Induction in in Response to Frontiers in Cellular and Infection Microbiology, 2022 , 12, 805473	5.9	
345	Importance of substrate quality and clay content on microbial extracellular polymeric substances production and aggregate stability in soils. <i>Biology and Fertility of Soils</i> , 2022 , 58, 435-457	6.1	O
344	Critical Assessment of Metagenome Interpretation: the second round of challenges <i>Nature Methods</i> , 2022 ,	21.6	14
343	Impact of intensive lifestyle intervention on gut microbiota composition in type 2 diabetes: a analysis of a randomized clinical trial <i>Gut Microbes</i> , 2022 , 14, 2005407	8.8	O
342	A novel and affordable bioaugmentation strategy with microbial extracts to accelerate the biodegradation of emerging contaminants in different media <i>Science of the Total Environment</i> , 2022 , 155234	10.2	1
341	Long range PCR reveals the genetic cargo of IncP-1 plasmids in the complex microbial community of an on-farm biopurification system treating pesticide contaminated wastewater. <i>Applied and Environmental Microbiology</i> , 2021 , AEM0164821	4.8	
340	Neonatal metabolome of cesarean section and risk of childhood asthma. <i>European Respiratory Journal</i> , 2021 ,	13.6	4
339	Emergent bacterial community properties induce enhanced drought tolerance in Arabidopsis. <i>Npj Biofilms and Microbiomes</i> , 2021 , 7, 82	8.2	4
338	Broad Dissemination of Plasmids across Groundwater-Fed Rapid Sand Filter Microbiomes. <i>MBio</i> , 2021 , e0306821	7.8	1
337	Unexpected diversity among small-scale sample replicates of defined plant root compartments. <i>ISME Journal</i> , 2021 ,	11.9	1
336	CRISPR-Cas systems are widespread accessory elements across bacterial and archaeal plasmids. <i>Nucleic Acids Research</i> , 2021 ,	20.1	6
335	EMBRACE-WATERS statement: Recommendations for reporting of studies on antimicrobial resistance in wastewater and related aquatic environments. <i>One Health</i> , 2021 , 13, 100339	7.6	2
334	The treasure inside barley seeds: microbial diversity and plant beneficial bacteria. <i>Environmental Microbiomes</i> , 2021 , 16, 20	5.6	6

(2021-2021)

333	Urbanized microbiota in infants, immune constitution, and later risk of atopic diseases. <i>Journal of Allergy and Clinical Immunology</i> , 2021 , 148, 234-243	11.5	11
332	Temporal oral microbiome changes with brushing in children with cleft lip and palate. <i>Heliyon</i> , 2021 , 7, e06513	3.6	О
331	The Airway Microbiota Modulates Effect of Azithromycin Treatment for Episodes of Recurrent Asthma-like Symptoms in Preschool Children: A Randomized Clinical Trial. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021 , 204, 149-158	10.2	9
330	Extended-Spectrum Lactamase and Carbapenemase Genes are Substantially and Sequentially Reduced during Conveyance and Treatment of Urban Sewage. <i>Environmental Science & Environmental Science & Technology</i> , 2021 , 55, 5939-5949	10.3	3
329	Lead Drives Complex Dynamics of a Conjugative Plasmid in a Bacterial Community. <i>Frontiers in Microbiology</i> , 2021 , 12, 655903	5.7	0
328	The infant gut resistome associates with E. coli, environmental exposures, gut microbiome maturity, and asthma-associated bacterial composition. <i>Cell Host and Microbe</i> , 2021 , 29, 975-987.e4	23.4	14
327	Kin discrimination promotes horizontal gene transfer between unrelated strains in Bacillus subtilis. <i>Nature Communications</i> , 2021 , 12, 3457	17.4	7
326	Metagenomic analysis of a keratin-degrading bacterial consortium provides insight into the keratinolytic mechanisms. <i>Science of the Total Environment</i> , 2021 , 761, 143281	10.2	9
325	The initial inoculation ratio regulates bacterial coculture interactions and metabolic capacity. <i>ISME Journal</i> , 2021 , 15, 29-40	11.9	12
324	Modeling transfer of vaginal microbiota from mother to infant in early life. ELife, 2021, 10,	8.9	10
323	The effect of early probiotic exposure on the preterm infant gut microbiome development. <i>Gut Microbes</i> , 2021 , 13, 1951113	8.8	3
322	Microbial community analysis of soils under different soybean cropping regimes in the Argentinean south-eastern Humid Pampas. <i>FEMS Microbiology Ecology</i> , 2021 , 97,	4.3	2
321	Large-scale association analyses identify host factors influencing human gut microbiome composition. <i>Nature Genetics</i> , 2021 , 53, 156-165	36.3	80
320	Reduced tillage, cover crops and organic amendments affect soil microbiota and improve soil health in Uruguayan vegetable farming systems. <i>FEMS Microbiology Ecology</i> , 2021 , 97,	4.3	3
319	Tillage shapes the soil and rhizosphere microbiome of barley-but not its susceptibility towards Blumeria graminis f. sp. hordei. <i>FEMS Microbiology Ecology</i> , 2021 , 97,	4.3	5
318	Identification of Beneficial Microbial Consortia and Bioactive Compounds with Potential as Plant Biostimulants for a Sustainable Agriculture. <i>Microorganisms</i> , 2021 , 9,	4.9	9
317	A community resource for paired genomic and metabolomic data mining. <i>Nature Chemical Biology</i> , 2021 , 17, 363-368	11.7	32
316	Root exposure to apple replant disease soil triggers local defense response and rhizoplane microbiome dysbiosis. <i>FEMS Microbiology Ecology</i> , 2021 , 97,	4.3	7

315	Distinct rhizomicrobiota assemblages and plant performance in lettuce grown in soils with different agricultural management histories. <i>FEMS Microbiology Ecology</i> , 2021 , 97,	4.3	2
314	Comparative analysis of the alveolar microbiome in COPD, ECOPD, Sarcoidosis, and ILD patients to identify respiratory illnesses specific microbial signatures. <i>Scientific Reports</i> , 2021 , 11, 3963	4.9	12
313	Importance of microbial communities at the root-soil interface for extracellular polymeric substances and soil aggregation in semiarid grasslands. <i>Soil Biology and Biochemistry</i> , 2021 , 159, 10830	1 ^{7.5}	2
312	Spatial analysis of the root system coupled to microbial community inoculation shed light on rhizosphere bacterial community assembly. <i>Biology and Fertility of Soils</i> , 2021 , 57, 973-989	6.1	Ο
311	Metagenomic evidence for co-occurrence of antibiotic, biocide and metal resistance genes in pigs. <i>Environment International</i> , 2021 , 158, 106899	12.9	4
310	CRISPRCasTyper: Automated Identification, Annotation, and Classification of CRISPR-Cas Loci. <i>CRISPR Journal</i> , 2020 , 3, 462-469	2.5	26
309	Discovery of multiple anti-CRISPRs highlights anti-defense gene clustering in mobile genetic elements. <i>Nature Communications</i> , 2020 , 11, 5652	17.4	24
308	Metal-induced bacterial interactions promote diversity in river-sediment microbiomes. <i>FEMS Microbiology Ecology</i> , 2020 , 96,	4.3	6
307	Ecological succession in the vaginal microbiota during pregnancy and birth. ISME Journal, 2020, 14, 232	5-12:3335	18
306	Compatibility of X-ray computed tomography with plant gene expression, rhizosphere bacterial communities and enzyme activities. <i>Journal of Experimental Botany</i> , 2020 , 71, 5603-5614	7	12
305	Prokaryotic Community Composition and Extracellular Polymeric Substances Affect Soil Microaggregation in Carbonate Containing Semiarid Grasslands. <i>Frontiers in Environmental Science</i> , 2020 , 8,	4.8	9
304	Salmonella persistence in soil depends on reciprocal interactions with indigenous microorganisms. <i>Environmental Microbiology</i> , 2020 , 22, 2639-2652	5.2	12
303	Azo dying of Ekeratin material improves microbial keratinase screening and standardization. <i>Microbial Biotechnology</i> , 2020 , 13, 984-996	6.3	5
302	Comparison of antibiotic-resistant bacteria and antibiotic resistance genes abundance in hospital and community wastewater: A systematic review. <i>Science of the Total Environment</i> , 2020 , 743, 140804	10.2	36
301	Prenatal dietary supplements influence the infant airway microbiota in a randomized factorial clinical trial. <i>Nature Communications</i> , 2020 , 11, 426	17.4	13
300	Community-intrinsic properties enhance keratin degradation from bacterial consortia. <i>PLoS ONE</i> , 2020 , 15, e0228108	3.7	11
299	Plasmids persist in a microbial community by providing fitness benefit to multiple phylotypes. <i>ISME Journal</i> , 2020 , 14, 1170-1181	11.9	23
298	Type IV CRISPR-Cas systems are highly diverse and involved in competition between plasmids. <i>Nucleic Acids Research</i> , 2020 , 48, 2000-2012	20.1	57

(2019-2020)

297	Interspecific interactions in dual-species biofilms of soil bacteria: effects of fertilization practices. Journal of Soils and Sediments, 2020 , 20, 1494-1501	3.4	4
296	Exploring microbial determinants of apple replant disease (ARD): a microhabitat approach under split-root design. <i>FEMS Microbiology Ecology</i> , 2020 , 96,	4.3	7
295	Epigenetic landscape links upper airway microbiota in infancy with allergic rhinitis at 6 years of age. Journal of Allergy and Clinical Immunology, 2020 , 146, 1358-1366	11.5	14
294	Composted Sewage Sludge Influences the Microbiome and Persistence of Human Pathogens in Soil. <i>Microorganisms</i> , 2020 , 8,	4.9	8
293	Delivery mode and gut microbial changes correlate with an increased risk of childhood asthma. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	33
292	Environmental shaping of the bacterial and fungal community in infant bed dust and correlations with the airway microbiota. <i>Microbiome</i> , 2020 , 8, 115	16.6	10
291	Rhizosphere microbial communities associated to rose replant disease: links to plant growth and root metabolites. <i>Horticulture Research</i> , 2020 , 7, 144	7.7	9
2 90	The microbiome of captive hamadryas baboons. <i>Animal Microbiome</i> , 2020 , 2, 25	4.1	2
289	Persistence and progression of staphylococcal infection in the presence of public goods. <i>Npj Biofilms and Microbiomes</i> , 2020 , 6, 55	8.2	1
288	Unravelling plasmidome distribution and interaction with its hosting microbiome. <i>Environmental Microbiology</i> , 2020 , 22, 32-44	5.2	7
287	Selection and propagation of IncP conjugative plasmids following long-term anthropogenic metal pollution in river sediments. <i>Journal of Hazardous Materials</i> , 2020 , 382, 121173	12.8	6
286	Impact of Long-Term Organic and Mineral Fertilization on Rhizosphere Metabolites, Root-Microbial Interactions and Plant Health of Lettuce. <i>Frontiers in Microbiology</i> , 2020 , 11, 597745	5.7	6
285	Community-intrinsic properties enhance keratin degradation from bacterial consortia 2020 , 15, e02281	108	
284	Community-intrinsic properties enhance keratin degradation from bacterial consortia 2020 , 15, e02281	108	
283	Community-intrinsic properties enhance keratin degradation from bacterial consortia 2020 , 15, e02281	108	
282	Community-intrinsic properties enhance keratin degradation from bacterial consortia 2020 , 15, e02281	108	
281	Interspecies interactions reduce selection for a biofilm optimized variant in a four-species biofilm model. <i>Environmental Microbiology</i> , 2019 ,	5.2	1
280	Deciphering links between bacterial interactions and spatial organization in multispecies biofilms. <i>ISME Journal</i> , 2019 , 13, 3054-3066	11.9	28

279	Priority of Early Colonizers but No Effect on Cohabitants in a Synergistic Biofilm Community. <i>Frontiers in Microbiology</i> , 2019 , 10, 1949	5.7	6
278	Manure and Doxycycline Affect the Bacterial Community and Its Resistome in Lettuce Rhizosphere and Bulk Soil. <i>Frontiers in Microbiology</i> , 2019 , 10, 725	5.7	20
277	Effect of long-term organic and mineral fertilization strategies on rhizosphere microbiota assemblage and performance of lettuce. <i>Environmental Microbiology</i> , 2019 , 21, 2426-2439	5.2	22
276	Different Degrees of Niche Differentiation for Bacteria, Fungi, and Myxomycetes Within an Elevational Transect in the German Alps. <i>Microbial Ecology</i> , 2019 , 78, 764-780	4.4	8
275	The scent of symbiosis: gut bacteria may affect social interactions in leaf-cutting ants. <i>Animal Behaviour</i> , 2019 , 150, 239-254	2.8	19
274	Fate of CMY-2-Encoding Plasmids Introduced into the Human Fecal Microbiota by Exogenous. <i>Antimicrobial Agents and Chemotherapy</i> , 2019 , 63,	5.9	10
273	Heavy metal exposure causes changes in the metabolic health-associated gut microbiome and metabolites. <i>Environment International</i> , 2019 , 126, 454-467	12.9	69
272	Bacterial community analysis for investigating bacterial transfer from tonsils to the pig carcass. <i>International Journal of Food Microbiology</i> , 2019 , 295, 8-18	5.8	8
271	Construction of Simplified Microbial Consortia to Degrade Recalcitrant Materials Based on Enrichment and Dilution-to-Extinction Cultures. <i>Frontiers in Microbiology</i> , 2019 , 10, 3010	5.7	24
270	Amplicon sequencing provides more accurate microbiome information in healthy children compared to culturing. <i>Communications Biology</i> , 2019 , 2, 291	6.7	29
269	The needle mycobiome of Picea glauca 🖪 dynamic system reflecting surrounding environment and tree phenological traits. <i>Fungal Ecology</i> , 2019 , 41, 177-186	4.1	9
268	Improvement of pesticide removal in contaminated media using aqueous extracts from contaminated biopurification systems. <i>Science of the Total Environment</i> , 2019 , 691, 749-759	10.2	10
267	Quantitative Flow Cytometry to Understand Population Heterogeneity in Response to Changes in Substrate Availability in and Chemostats. <i>Frontiers in Bioengineering and Biotechnology</i> , 2019 , 7, 187	5.8	16
266	Interspecies interactions reduce selection for a biofilm-optimized variant in a four-species biofilm model. <i>Environmental Microbiology Reports</i> , 2019 , 11, 835-839	3.7	4
265	Infant airway microbiota and topical immune perturbations in the origins of childhood asthma. <i>Nature Communications</i> , 2019 , 10, 5001	17.4	52
264	Biocontrol of Bacterial Wilt Disease Through Complex Interaction Between Tomato Plant, Antagonists, the Indigenous Rhizosphere Microbiota, and. <i>Frontiers in Microbiology</i> , 2019 , 10, 2835	5.7	32
263	The effect of acetate on population heterogeneity in different cellular characteristics of Escherichia coli in aerobic batch cultures. <i>Biotechnology Progress</i> , 2019 , 35, e2796	2.8	4
262	DNA- and RNA-SIP Reveal spp. as Key Drivers of Nitrification in Groundwater-Fed Biofilters. <i>MBio</i> , 2019 , 10,	7.8	18

(2018-2019)

261	Tailings microbial community profile and prediction of its functionality in basins of tungsten mine. <i>Scientific Reports</i> , 2019 , 9, 19596	4.9	14
260	Intra- and inter-field diversity of 2,4-dichlorophenoxyacetic acid-degradative plasmids and their tfd catabolic genes in rice fields of the Mekong delta in Vietnam. <i>FEMS Microbiology Ecology</i> , 2019 , 95,	4.3	1
259	Impact of long-term agricultural management practices on soil prokaryotic communities. <i>Soil Biology and Biochemistry</i> , 2019 , 129, 17-28	7.5	59
258	Soil amendment with sewage sludge affects soil prokaryotic community composition, mobilome and resistome. <i>FEMS Microbiology Ecology</i> , 2019 , 95,	4.3	6
257	Bulk soil and maize rhizosphere resistance genes, mobile genetic elements and microbial communities are differently impacted by organic and inorganic fertilization. <i>FEMS Microbiology Ecology</i> , 2018 , 94,	4.3	19
256	Nutrient Fortification of Human Donor Milk Affects Intestinal Function and Protein Metabolism in Preterm Pigs. <i>Journal of Nutrition</i> , 2018 , 148, 336-347	4.1	20
255	Micro-scale intermixing: a requisite for stable and synergistic co-establishment in a four-species biofilm. <i>ISME Journal</i> , 2018 , 12, 1940-1951	11.9	24
254	Estimating the Transfer Range of Plasmids Encoding Antimicrobial Resistance in a Wastewater Treatment Plant Microbial Community. <i>Environmental Science and Technology Letters</i> , 2018 , 5, 260-265	11	59
253	Fluorescence Recovery Allows the Implementation of a Fluorescence Reporter Gene Platform Applicable for the Detection and Quantification of Horizontal Gene Transfer in Anoxic Environments. <i>Applied and Environmental Microbiology</i> , 2018 , 84,	4.8	10
252	Soil bacteria show different tolerance ranges to an unprecedented disturbance. <i>Biology and Fertility of Soils</i> , 2018 , 54, 189-202	6.1	19
251	Microbial Diversity and Putative Opportunistic Pathogens in Dishwasher Biofilm Communities. <i>Applied and Environmental Microbiology</i> , 2018 , 84,	4.8	31
250	Maturation of the gut microbiome and risk of asthma in childhood. <i>Nature Communications</i> , 2018 , 9, 14	117.4	216
249	Investigation of the bacteriophage community in induced lysates of undefined mesophilic mixed-strain DL-cultures using classical and metagenomic approaches. <i>International Journal of Food Microbiology</i> , 2018 , 272, 61-72	5.8	2
248	Rifaximin has minor effects on bacterial composition, inflammation, and bacterial translocation in cirrhosis: A randomized trial. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2018 , 33, 307-314	4	34
247	Long-term industrial metal contamination unexpectedly shaped diversity and activity response of sediment microbiome. <i>Journal of Hazardous Materials</i> , 2018 , 344, 299-307	12.8	52
246	Genetic barcoding of dark-spored myxomycetes (Amoebozoa)-Identification, evaluation and application of a sequence similarity threshold for species differentiation in NGS studies. <i>Molecular Ecology Resources</i> , 2018 , 18, 306-318	8.4	19
245	Microbial indicators for soil quality. Biology and Fertility of Soils, 2018, 54, 1-10	6.1	189
244	A culture-independent method for studying transfer of IncI1 plasmids from wild-type Escherichia coli in complex microbial communities. <i>Journal of Microbiological Methods</i> , 2018 , 152, 18-26	2.8	2

243	The bacterial microbiota in first-void urine from men with and without idiopathic urethritis. <i>PLoS ONE</i> , 2018 , 13, e0201380	3.7	18
242	Synergistic Interactions in Microbial Biofilms Facilitate the Establishment of Opportunistic Pathogenic Fungi in Household Dishwashers. <i>Frontiers in Microbiology</i> , 2018 , 9, 21	5.7	25
241	How Microbial Aggregates Protect against Nanoparticle Toxicity. <i>Trends in Biotechnology</i> , 2018 , 36, 117	′1£ ჭ.1 87	2 81
240	Monitoring plasmid-mediated horizontal gene transfer in microbiomes: recent advances and future perspectives. <i>Plasmid</i> , 2018 , 99, 56-67	3.3	33
239	An intriguing relationship between the cyclic diguanylate signaling system and horizontal gene transfer. <i>ISME Journal</i> , 2018 , 12, 2330-2334	11.9	16
238	Bacterial social interactions and the emergence of community-intrinsic properties. <i>Current Opinion in Microbiology</i> , 2018 , 42, 104-109	7.9	44
237	coagulases are exploitable yet stable public goods in clinically relevant conditions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, E11771-E11779	11.5	5
236	Stability and resilience of the intestinal microbiota in children in daycare - a 12 month cohort study. <i>BMC Microbiology</i> , 2018 , 18, 223	4.5	10
235	Long-term soil metal exposure impaired temporal variation in microbial metatranscriptomes and enriched active phages. <i>Microbiome</i> , 2018 , 6, 223	16.6	20
234	Free-Living Nematodes Together With Associated Microbes Play an Essential Role in Apple Replant Disease. <i>Frontiers in Plant Science</i> , 2018 , 9, 1666	6.2	16
233	Short- and long-term impacts of azithromycin treatment on the gut microbiota in children: A double-blind, randomized, placebo-controlled trial. <i>EBioMedicine</i> , 2018 , 38, 265-272	8.8	37
232	Changes in rumen bacterial and archaeal communities over the transition period in primiparous Holstein dairy cows. <i>Journal of Dairy Science</i> , 2018 , 101, 9847-9862	4	19
231	Enrichment and characterization of an environmental microbial consortium displaying efficient keratinolytic activity. <i>Bioresource Technology</i> , 2018 , 270, 303-310	11	29
230	Enhanced bacterial mutualism through an evolved biofilm phenotype. ISME Journal, 2018, 12, 2608-261	8 11.9	19
229	Low-abundant species facilitates specific spatial organization that promotes multispecies biofilm formation. <i>Environmental Microbiology</i> , 2017 , 19, 2893-2905	5.2	38
228	Alterations of the murine gut microbiome in allergic airway disease are independent of surfactant protein D. <i>Heliyon</i> , 2017 , 3, e00262	3.6	5
227	Whole-Genome Sequence of EK007-RG4, a Promising Biocontrol Agent against a Broad Range of Bacteria, Including the Fire Blight Bacterium. <i>Genome Announcements</i> , 2017 , 5,		1
226	Autogenic succession and deterministic recovery following disturbance in soil bacterial communities. <i>Scientific Reports</i> , 2017 , 7, 45691	4.9	50

(2017-2017)

225	Deciphering conjugative plasmid permissiveness in wastewater microbiomes. <i>Molecular Ecology</i> , 2017 , 26, 3556-3571	5.7	41	
224	It is elemental: soil nutrient stoichiometry drives bacterial diversity. <i>Environmental Microbiology</i> , 2017 , 19, 1176-1188	5.2	154	
223	Critical Assessment of Metagenome Interpretation-a benchmark of metagenomics software. <i>Nature Methods</i> , 2017 , 14, 1063-1071	21.6	412	
222	Community structure of the metabolically active rumen bacterial and archaeal communities of dairy cows over the transition period. <i>PLoS ONE</i> , 2017 , 12, e0187858	3.7	20	
221	Microbiomes associated with infective stages of root-knot and lesion nematodes in soil. <i>PLoS ONE</i> , 2017 , 12, e0177145	3.7	49	
220	In situ metabolomic- and transcriptomic-profiling of the host-associated cyanobacteria Prochloron and Acaryochloris marina. <i>ISME Journal</i> , 2017 ,	11.9	6	
219	Synergistic Interactions within a Multispecies Biofilm Enhance Individual Species Protection against Grazing by a Pelagic Protozoan. <i>Frontiers in Microbiology</i> , 2017 , 8, 2649	5.7	28	
218	Antagonism correlates with metabolic similarity in diverse bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, 10684-10688	11.5	74	
217	Response of the bacterial community in an on-farm biopurification system, to which diverse pesticides are introduced over an agricultural season. <i>Environmental Pollution</i> , 2017 , 229, 854-862	9.3	25	
216	Metagenomic Analysis of Dairy Bacteriophages: Extraction Method and Pilot Study on Whey Samples Derived from Using Undefined and Defined Mesophilic Starter Cultures. <i>Applied and Environmental Microbiology</i> , 2017 , 83,	4.8	13	
215	A meta-proteomics approach to study the interspecies interactions affecting microbial biofilm development in a model community. <i>Scientific Reports</i> , 2017 , 7, 16483	4.9	35	
214	Metal stressors consistently modulate bacterial conjugal plasmid uptake potential in a phylogenetically conserved manner. <i>ISME Journal</i> , 2017 , 11, 152-165	11.9	68	
213	Distinct gene expression profile of Xanthomonas retroflexus engaged in synergistic multispecies biofilm formation. <i>ISME Journal</i> , 2017 , 11, 300-303	11.9	35	
212	A post-planktonic era of in vitro infectious models: issues and changes addressed by a clinically relevant wound like media. <i>Critical Reviews in Microbiology</i> , 2017 , 43, 453-465	7.8	13	
211	Comparative Genomic Analysis Reveals Ecological Differentiation in the Genus. <i>Frontiers in Microbiology</i> , 2017 , 8, 357	5.7	17	
210	Effects of Soil Pre-Treatment with Basamid Granules, , and on Bacterial and Fungal Communities at Two Apple Replant Disease Sites. <i>Frontiers in Microbiology</i> , 2017 , 8, 1604	5.7	38	
209	Legacy Effects on the Recovery of Soil Bacterial Communities from Extreme Temperature Perturbation. <i>Frontiers in Microbiology</i> , 2017 , 8, 1832	5.7	54	
208	A novel baiting microcosm approach used to identify the bacterial community associated with Penicillium bilaii hyphae in soil. <i>PLoS ONE</i> , 2017 , 12, e0187116	3.7	12	

207	Diverse gene functions in a soil mobilome. Soil Biology and Biochemistry, 2016, 101, 175-183	7.5	16
206	Coping with copper: legacy effect of copper on potential activity of soil bacteria following a century of exposure. <i>FEMS Microbiology Ecology</i> , 2016 , 92,	4.3	91
205	Enhanced plasmid loss in bacterial populations exposed to the antimicrobial compound irgasan delivered from interpenetrating polymer network silicone hydrogels. <i>Plasmid</i> , 2016 , 87-88, 72-78	3.3	13
204	Bacterial communities in termite fungus combs are comprised of consistent gut deposits and contributions from the environment. <i>Microbial Ecology</i> , 2016 , 71, 207-20	4.4	29
203	Plasmid metagenomics reveals multiple antibiotic resistance gene classes among the gut microbiomes of hospitalised patients. <i>Journal of Global Antimicrobial Resistance</i> , 2016 , 6, 57-66	3.4	10
202	Draft Genome Sequences of Two Kocuria Isolates, K. salsicia G1 and K. rhizophila G2, Isolated from a Slaughterhouse in Denmark. <i>Genome Announcements</i> , 2016 , 4,		1
201	Coexistence facilitates interspecific biofilm formation in complex microbial communities. <i>Environmental Microbiology</i> , 2016 , 18, 2565-74	5.2	53
200	Studying Bacterial Multispecies Biofilms: Where to Start?. <i>Trends in Microbiology</i> , 2016 , 24, 503-513	12.4	109
199	Ecological patterns, diversity and core taxa of microbial communities in groundwater-fed rapid gravity filters. <i>ISME Journal</i> , 2016 , 10, 2209-22	11.9	81
198	Exploring the complex response to linuron of bacterial communities from biopurification systems by means of cultivation-independent methods. <i>FEMS Microbiology Ecology</i> , 2016 , 92,	4.3	16
197	Back to the Future of Soil Metagenomics. Frontiers in Microbiology, 2016, 7, 73	5.7	82
196	Interspecific Bacterial Interactions are Reflected in Multispecies Biofilm Spatial Organization. <i>Frontiers in Microbiology</i> , 2016 , 7, 1366	5.7	92
195	Type 3 Fimbriae Encoded on Plasmids Are Expressed from a Unique Promoter without Affecting Host Motility, Facilitating an Exceptional Phenotype That Enhances Conjugal Plasmid Transfer. <i>PLoS ONE</i> , 2016 , 11, e0162390	3.7	12
194	Genome Sequence of Kocuria varians G6 Isolated from a Slaughterhouse in Denmark. <i>Genome Announcements</i> , 2016 , 4,		2
193	Large-scale benchmarking reveals false discoveries and count transformation sensitivity in 16S rRNA gene amplicon data analysis methods used in microbiome studies. <i>Microbiome</i> , 2016 , 4, 62	16.6	93
192	The developing hypopharyngeal microbiota in early life. <i>Microbiome</i> , 2016 , 4, 70	16.6	34
191	Metagenomes provide valuable comparative information on soil microeukaryotes. <i>Research in Microbiology</i> , 2016 , 167, 436-50	4	28
190	Genomic and exoproteomic analyses of cold- and alkaline-adapted bacteria reveal an abundance of secreted subtilisin-like proteases. <i>Microbial Biotechnology</i> , 2016 , 9, 245-56	6.3	6

(2015-2016)

189	Natural decay process affects the abundance and community structure of Bacteria and Archaea in Picea abies logs. <i>FEMS Microbiology Ecology</i> , 2016 , 92,	4.3	37
188	Vitamin D and allergic airway disease shape the murine lung microbiome in a sex-specific manner. <i>Respiratory Research</i> , 2016 , 17, 116	7.3	22
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(2013-2014)

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Microbiology Letters, 2007, 266, 250-6 The multiple antibiotic resistance IncP-1 plasmid pKJKS isolated from a soil environment is phylogenetically divergent from members of the previously established alpha, beta and delta sub-groups. Plasmid, 2007, 58, 31-43 Establishment and early succession of a multispecies biofilm composed of soil bacteria. Microbial Ecology, 2007, 54, 352-62 444 Substrate specificity of the OqxAB multidrug resistance pump in Escherichia coli and selected enteric bacteria. Journal of Antimicrobial Chemotherapy, 2007, 60, 145-7 Conjugative transfer facilitates stable maintenance of IncP-1 plasmid pKJKS in Escherichia coli cells colonizing the gastrointestinal tract of the germfree rat. Applied and Environmental Microbiology, 2007, 73, 341-3 The cda GenoTox assay: a new and sensitive method for detection of environmental genotoxins, including nitroarenes and aromatic amines. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2007, 631, 77-84 Cultivation-independent examination of horizontal transfer and host range of an IncP-1 plasmid among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and Environmental Microbiology, 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & Comp. Technology, 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Muta	78		11.9	47
phylogenetically divergent from members of the previously established alpha, beta and delta sub-groups. <i>Plasmid</i> , 2007, 58, 31-43 Establishment and early succession of a multispecies biofilm composed of soil bacteria. <i>Microbiolal Ecology</i> , 2007, 54, 352-62 44 Substrate specificity of the OqxAB multidrug resistance pump in Escherichia coli and selected enteric bacteria. <i>Journal of Antimicrobial Chemotherapy</i> , 2007, 60, 145-7 Conjugative transfer facilitates stable maintenance of IncP-1 plasmid pKJK5 in Escherichia coli cells colonizing the gastrointestinal tract of the germfree rat. <i>Applied and Environmental Microbiology</i> , 2007, 73, 341-3 The cda GenoTox assay: a new and sensitive method for detection of environmental genotoxins, including nitroarenes and aromatic amines. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2007, 631, 77-84 Cultivation-independent examination of horizontal transfer and host range of an IncP-1 plasmid among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. <i>Applied and Environmental Microbiology</i> , 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. <i>Applied and Environmental Microbiology</i> , 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. <i>Environmental Science & Dispersional Processor</i> , 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. <i>Environmental Pollution</i> , 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in micr	77		2.9	43
Substrate specificity of the OqxAB multidrug resistance pump in Escherichia coli and selected enteric bacteria. Journal of Antimicrobial Chemotherapy, 2007, 60, 145-7 Conjugative transfer facilitates stable maintenance of IncP-1 plasmid pKJK5 in Escherichia coli cells colonizing the gastrointestinal tract of the germfree rat. Applied and Environmental Microbiology, 2007, 73, 341-3 The cda GenoTox assay: a new and sensitive method for detection of environmental genotoxins, including nitroarenes and aromatic amines. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2007, 631, 77-84 Cultivation-independent examination of horizontal transfer and host range of an IncP-1 plasmid among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and Environmental Microbiology, 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & Description (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	76	phylogenetically divergent from members of the previously established alpha, beta and delta	3.3	62
conjugative transfer facilitates stable maintenance of IncP-1 plasmid pKJK5 in Escherichia coli cells colonizing the gastrointestinal tract of the germfree rat. Applied and Environmental Microbiology, 2007, 73, 341-3 The cda GenoTox assay: a new and sensitive method for detection of environmental genotoxins, including nitroarenes and aromatic amines. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2007, 631, 77-84 Cultivation-independent examination of horizontal transfer and host range of an IncP-1 plasmid among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and Environmental Microbiology, 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & amp; Technology, 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	75		4.4	47
colonizing the gastrointestinal tract of the germfree rat. Applied and Environmental Microbiology, 2007, 73, 341-3 The cda GenoTox assay: a new and sensitive method for detection of environmental genotoxins, including nitroarenes and aromatic amines. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2007, 631, 77-84 Cultivation-independent examination of horizontal transfer and host range of an IncP-1 plasmid among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and Environmental Microbiology, 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & amp; Technology, 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 4 flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	74		5.1	176
including nitroarenes and aromatic amines. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2007, 631, 77-84 Cultivation-independent examination of horizontal transfer and host range of an IncP-1 plasmid among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and Environmental Microbiology, 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & Description and Environmental Pollution, 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	73	colonizing the gastrointestinal tract of the germfree rat. Applied and Environmental Microbiology,	4.8	19
among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and Environmental Microbiology, 2006, 72, 6687-92 Enhanced biofilm formation and increased resistance to antimicrobial agents and bacterial invasion are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & amp; Technology, 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	72	including nitroarenes and aromatic amines. Mutation Research - Genetic Toxicology and	3	14
are caused by synergistic interactions in multispecies biofilms. Applied and Environmental Microbiology, 2006, 72, 3916-23 Biodegradation, bioaccessibility, and genotoxicity of diffuse polycyclic aromatic hydrocarbon (PAH) pollution at a motorway site. Environmental Science & Delay 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	71	among gram-positive and gram-negative bacteria indigenous to the barley rhizosphere. Applied and	4.8	81
pollution at a motorway site. Environmental Science & Samp; Technology, 2006, 40, 3293-8 Mineralisation of 2,6-dichlorobenzamide (BAM) in dichlobenil-exposed soils and isolation of a BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	70	are caused by synergistic interactions in multispecies biofilms. Applied and Environmental	4.8	470
BAM-mineralising Aminobacter sp. Environmental Pollution, 2006, 144, 289-95 A flow cytometry-optimized assay using an SOS-green fluorescent protein (SOS-GFP) whole-cell biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	69		10.3	42
biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2006, 603, 164-72 Microbial degradation of street dust polycyclic aromatic hydrocarbons in microcosms simulating diffuse pollution of urban soil. Environmental Microbiology, 2006, 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. Current Opinion in Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	68		9.3	36
diffuse pollution of urban soil. <i>Environmental Microbiology</i> , 2006 , 8, 535-45 Making bio-sense of toxicity: new developments in whole-cell biosensors. <i>Current Opinion in Biotechnology</i> , 2006 , 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	67	biosensor for the detection of genotoxins in complex environments. Mutation Research - Genetic	3	35
Biotechnology, 2006, 17, 11-6 Reporter Gene Technology in Soil Ecology; Detection of Bioavailability and Microbial Interactions	66		5.2	49
	65		11.4	103
	64			5

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62	The prevalence of the OqxAB multidrug efflux pump amongst olaquindox-resistant Escherichia coli in pigs. <i>Microbial Drug Resistance</i> , 2005 , 11, 378-82	2.9	61
61	Studying plasmid horizontal transfer in situ: a critical review. <i>Nature Reviews Microbiology</i> , 2005 , 3, 700-	·1 <u>10</u> 2.2	485
60	Use of a whole-cell biosensor and flow cytometry to detect AHL production by an indigenous soil community during decomposition of litter. <i>Microbial Ecology</i> , 2005 , 50, 221-9	4.4	36
59	Construction of an extended range whole-cell tetracycline biosensor by use of the tet(M) resistance gene. <i>FEMS Microbiology Letters</i> , 2005 , 253, 201-5	2.9	33
58	In vivo detection and quantification of tetracycline by use of a whole-cell biosensor in the rat intestine. <i>Antimicrobial Agents and Chemotherapy</i> , 2004 , 48, 1112-7	5.9	36
57	Effect of tetracycline on transfer and establishment of the tetracycline-inducible conjugative transposon Tn916 in the guts of gnotobiotic rats. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 758	3- 4 :8	50
56	Plasmid-encoded multidrug efflux pump conferring resistance to olaquindox in Escherichia coli. <i>Antimicrobial Agents and Chemotherapy</i> , 2004 , 48, 3332-7	5.9	161
55	Impact of DNA extraction method on bacterial community composition measured by denaturing gradient gel electrophoresis. <i>Soil Biology and Biochemistry</i> , 2004 , 36, 1607-1614	7.5	84
54	Recovery of GFP-labeled bacteria for culturing and molecular analysis after cell sorting using a benchtop flow cytometer. <i>Microbial Ecology</i> , 2004 , 48, 239-45	4.4	29
53	Quantification of plasmid loss in Escherichia coli cells by use of flow cytometry. <i>FEMS Microbiology Letters</i> , 2004 , 232, 45-9	2.9	22
52	Direct detection and quantification of horizontal gene transfer by using flow cytometry and gfp as a reporter gene. <i>Current Microbiology</i> , 2003 , 47, 129-33	2.4	41
51	Presence of N-acyl homoserine lactones in soil detected by a whole-cell biosensor and flow cytometry. <i>Microbial Ecology</i> , 2003 , 45, 226-36	4.4	67
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49	Conjugative plasmid conferring resistance to olaquindox. <i>Antimicrobial Agents and Chemotherapy</i> , 2003 , 47, 798-9	5.9	69
48	Quantification of bioavailable chlortetracycline in pig feces using a bacterial whole-cell biosensor. <i>Veterinary Microbiology</i> , 2002 , 87, 51-7	3.3	27
47	The diversity and function of soil microbial communities exposed to different disturbances. <i>Microbial Ecology</i> , 2002 , 44, 49-58	4.4	146
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(2000-2001)

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43	The effect of long-term mercury pollution on the soil microbial community 2001 , 36, 11-19		139
42	Adaptation of the bacterial community to mercury contamination. <i>FEMS Microbiology Letters</i> , 2001 , 204, 49-53	2.9	33
41	Enhanced degradation of phenoxyacetic acid in soil by horizontal transfer of the tfdA gene encoding a 2,4-dichlorophenoxyacetic acid dioxygenase. <i>FEMS Microbiology Ecology</i> , 2001 , 35, 75-84	4.3	42
40	Effects of mercury contamination on the culturable heterotrophic, functional and genetic diversity of the bacterial community in soil. <i>FEMS Microbiology Ecology</i> , 2001 , 36, 1-9	4.3	106
39	The effect of long-term mercury pollution on the soil microbial community. <i>FEMS Microbiology Ecology</i> , 2001 , 36, 11-19	4.3	51
38	Effect of genomic location on horizontal transfer of a recombinant gene cassette between Pseudomonas strains in the rhizosphere and spermosphere of barley seedlings. <i>Current Microbiology</i> , 2001 , 42, 160-7	2.4	44
37	Adaptation of the bacterial community to mercury contamination. <i>FEMS Microbiology Letters</i> , 2001 , 204, 49-53	2.9	38
36	An examination of the biodiversity acosystem function relationship in arable soil microbial communities. <i>Soil Biology and Biochemistry</i> , 2001 , 33, 1713-1722	7.5	210
35	Effects of tylosin as a disturbance on the soil microbial community. <i>Soil Biology and Biochemistry</i> , 2001 , 33, 2061-2071	7.5	136
34	Detection of oxytetracycline production by Streptomyces rimosus in soil microcosms by combining whole-cell biosensors and flow cytometry. <i>Applied and Environmental Microbiology</i> , 2001 , 67, 239-44	4.8	66
33	Ecosystem response of pasture soil communities to fumigation-induced microbial diversity reductions: an examination of the biodiversity cosystem function relationship. <i>Oikos</i> , 2000 , 90, 279-29	94 ⁴	436
32	Influence of fungal-bacterial interactions on bacterial conjugation in the residuesphere. <i>FEMS Microbiology Ecology</i> , 2000 , 31, 39-45	4.3	4
31	Detection and quantification of tetracyclines by whole cell biosensors. <i>FEMS Microbiology Letters</i> , 2000 , 190, 273-8	2.9	64
30	Versatile biosensor vectors for detection and quantification of mercury. <i>FEMS Microbiology Letters</i> , 2000 , 193, 123-7	2.9	84
29	A new method for determining the metabolic activity of specific bacterial populations in soil using tritiated leucine and immunomagnetic separation. <i>Biology and Fertility of Soils</i> , 2000 , 32, 202-208	6.1	4
28	Application of a mer-lux biosensor for estimating bioavailable mercury in soil. <i>Soil Biology and Biochemistry</i> , 2000 , 32, 639-646	7.5	136

27	Utilization of phenoxyacetic acid, by strains using either the ortho or meta cleavage of catechol during phenol degradation, after conjugal transfer of tfdA, the gene encoding a 2,4-dichlorophenoxyacetic acid/2-oxoglutarate dioxygenase. <i>Applied Microbiology and</i>	5.7	15
26	Biotechnology, 1999, 51, 207-14 Predation by protozoa on Escherichia coli K12 in soil and transfer of resistance plasmid RP4 to indigenous bacteria in soil. <i>Applied Soil Ecology</i> , 1999, 11, 79-90	5	30
25	Transfer of plasmid RP4 in the spermosphere and rhizosphere of barley seedling. <i>Antonie Van Leeuwenhoek</i> , 1998 , 73, 69-77	2.1	32
24	The effect of longterm exposure to mercury on the bacterial community in marine sediment. <i>Current Microbiology</i> , 1998 , 36, 291-7	2.4	70
23	The effect of the lacY gene on the induction of IPTG inducible promoters, studied in Escherichia coli and Pseudomonas fluorescens. <i>Current Microbiology</i> , 1998 , 36, 341-7	2.4	52
22	Methods for detection of conjugative plasmid transfer in aquatic environments. <i>Current Microbiology</i> , 1998 , 37, 274-80	2.4	20
21	Chromosomal insertion of the entire Escherichia coli lactose operon, into two strains of Pseudomonas, using a modified mini-Tn5 delivery system. <i>Gene</i> , 1997 , 186, 167-73	3.8	47
20	Conjugation in aquatic environments 1996 , 95-123		2
19	Conjugal transfer at natural population densities in a microcosm simulating an estuarine environment. <i>FEMS Microbiology Ecology</i> , 1995 , 16, 43-54	4.3	22
18	Transfer of plasmid RP4 from Escherichia coli K-12 to indigenous bacteria of seawater. <i>Microbial Releases: Viruses, Bacteria, Fungi</i> , 1993 , 2, 135-41		5
17	Mobilization of nonconjugative pBR322-derivative plasmids from laboratory strains of Escherichia coli to bacteria isolated from seawater. <i>Microbial Releases: Viruses, Bacteria, Fungi,</i> 1992 , 1, 17-22		1
16	Survival of Escherichia coli K12 in seawater. <i>FEMS Microbiology Ecology</i> , 1991 , 8, 161-167	4.3	
15	Survival of Escherichia coli K12 in seawater. FEMS Microbiology Letters, 1991, 85, 161-167	2.9	9
14	Deep mutational scanning by FACS-sorting of encapsulated E. coli micro-colonies		1
13	Effects of mercury contamination on the culturable heterotrophic, functional and genetic diversity of the bacterial community in soil		2
12	Detection and quantification of tetracyclines by whole cell biosensors		2
11	Conjugative dissemination of plasmids in rapid sand filters: a trojan horse strategy to enhance pesticide degradation in groundwater treatment		3
10	CRISPRCasTyper: An automated tool for the identification, annotation and classification of CRISPR-Cas loci		9

LIST OF PUBLICATIONS

9	Discovery of multiple anti-CRISPRs uncovers anti-defense gene clustering in mobile genetic elements	2
8	Large-scale association analyses identify host factors influencing human gut microbiome composition	9
7	DAtest: a framework for choosing differential abundance or expression method	17
6	Type IV CRISPR-Cas systems are highly diverse and involved in competition between plasmids	5
5	CRISPR-Cas systems are widespread accessory elements across bacterial and archaeal plasmids	1
4	Hundreds of viral families in the healthy infant gut	3
3	Critical Assessment of Metagenome Interpretation - the second round of challenges	3
2	Metagenomic evidence for co-occurrence of antibiotic, biocide and metal resistance genes in pigs	1
1	Comparative genomics analysis of Chryseobacterium sp. KMC2 reveals metabolic pathways involved in keratinous utilization and natural product biosynthesis	1