

# Trent R Northen

## 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.

172  
papers

8,395  
citations

42  
h-index

89  
g-index

193  
ext. papers

11,671  
ext. citations

8.9  
avg, IF

6.21  
L-index

#	Paper	IF	Citations
172	Vertical movement of soluble carbon and nutrients from biocrusts to subsurface mineral soils. <i>Geoderma</i> , <b>2022</b> , 405, 115495	6.7	3
171	SIMILE enables alignment of tandem mass spectra with statistical significance.. <i>Nature Communications</i> , <b>2022</b> , 13, 2510	17.4	0
170	Development of platforms for functional characterization and production of phenazines using a multi-chassis approach via CRAGE. <i>Metabolic Engineering</i> , <b>2021</b> , 69, 188-188	9.7	1
169	Identification of Effector Metabolites Using Exometabolite Profiling of Diverse Microalgae. <i>MSystems</i> , <b>2021</b> , e0083521	7.6	0
168	Long-read metagenomics of soil communities reveals phylum-specific secondary metabolite dynamics. <i>Communications Biology</i> , <b>2021</b> , 4, 1302	6.7	2
167	Ecological generalism drives hyperdiversity of secondary metabolite gene clusters in xylarialean endophytes. <i>New Phytologist</i> , <b>2021</b> ,	9.8	3
166	GNPS Dashboard: collaborative exploration of mass spectrometry data in the web browser. <i>Nature Methods</i> , <b>2021</b> ,	21.6	5
165	Rhizosphere Carbon Turnover from Cradle to Grave: The Role of MicrobePlant Interactions. <i>Rhizosphere Biology</i> , <b>2021</b> , 51-73	0.8	12
164	A multi-omic characterization of temperature stress in a halotolerant <i>Scenedesmus</i> strain for algal biotechnology. <i>Communications Biology</i> , <b>2021</b> , 4, 333	6.7	6
163	Specialized Plant Growth Chamber Designs to Study Complex Rhizosphere Interactions. <i>Frontiers in Microbiology</i> , <b>2021</b> , 12, 625752	5.7	3
162	Metabolomic Patterns of Septoria Canker Resistant and Susceptible Genotypes 24 Hours Postinoculation. <i>Phytopathology</i> , <b>2021</b> , PHYTO02210053R	3.8	0
161	Anaerobic gut fungi are an untapped reservoir of natural products. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2021</b> , 118,	11.5	11
160	Decrypting bacterial polyphenol metabolism in an anoxic wetland soil. <i>Nature Communications</i> , <b>2021</b> , 12, 2466	17.4	8
159	Genomics, Exometabolomics, and Metabolic Probing Reveal Conserved Proteolytic Metabolism of and Three Candidate Species From China and Japan. <i>Frontiers in Microbiology</i> , <b>2021</b> , 12, 632731	5.7	2
158	A multiplexed nanostructure-initiator mass spectrometry (NIMS) assay for simultaneously detecting glycosyl hydrolase and lignin modifying enzyme activities. <i>Scientific Reports</i> , <b>2021</b> , 11, 11803	4.9	1
157	A genomic catalog of Earth's microbiomes. <i>Nature Biotechnology</i> , <b>2021</b> , 39, 499-509	44.5	120
156	Faster, better, and cheaper: harnessing microfluidics and mass spectrometry for biotechnology. <i>RSC Chemical Biology</i> , <b>2021</b> , 2, 1331-1351	3	3

155	Reply to: Examining microbe-metabolite correlations by linear methods. <i>Nature Methods</i> , <b>2021</b> , 18, 40-41	11.6	2
154	Bioactive diterpenoids impact the composition of the root-associated microbiome in maize ( <i>Zea mays</i> ). <i>Scientific Reports</i> , <b>2021</b> , 11, 333	4.9	11
153	Cocultivation of Anaerobic Fungi with Rumen Bacteria Establishes an Antagonistic Relationship. <i>MBio</i> , <b>2021</b> , 12, e0144221	7.8	0
152	Getting back to the grass roots: harnessing specialized metabolites for improved crop stress resilience. <i>Current Opinion in Biotechnology</i> , <b>2021</b> , 70, 174-186	11.4	2
151	CRAGE-CRISPR facilitates rapid activation of secondary metabolite biosynthetic gene clusters in bacteria. <i>Cell Chemical Biology</i> , <b>2021</b> ,	8.2	2
150	Novel and Emerging Capabilities that Can Provide a Holistic Understanding of the Plant Root Microbiome. <i>Phytobiomes Journal</i> , <b>2021</b> , 5, 122-132	4.8	3
149	Biofilm Interaction Mapping and Analysis (BIMA) of Interspecific Interactions in Co-culture Biofilms.. <i>Frontiers in Microbiology</i> , <b>2021</b> , 12, 757856	5.7	1
148	Drought and plant litter chemistry alter microbial gene expression and metabolite production. <i>ISME Journal</i> , <b>2020</b> , 14, 2236-2247	11.9	26
147	Untangling the sequence of events during the S <sub>2</sub> -R transition in photosystem II and implications for the water oxidation mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2020</b> , 117, 12624-12635	11.5	65
146	Function-driven single-cell genomics uncovers cellulose-degrading bacteria from the rare biosphere. <i>ISME Journal</i> , <b>2020</b> , 14, 659-675	11.9	29
145	A structural and kinetic survey of GH5_4 endoglucanases reveals determinants of broad substrate specificity and opportunities for biomass hydrolysis. <i>Journal of Biological Chemistry</i> , <b>2020</b> , 295, 17752-17769	5.4	2
144	Mass Spectrometry for Natural Product Discovery <b>2020</b> , 263-306		6
143	CRAGE-Duet Facilitates Modular Assembly of Biological Systems for Studying Plant-Microbe Interactions. <i>ACS Synthetic Biology</i> , <b>2020</b> , 9, 2610-2615	5.7	5
142	Cervicovaginal Microbiome Composition Is Associated with Metabolic Profiles in Healthy Pregnancy. <i>MBio</i> , <b>2020</b> , 11,	7.8	12
141	Root morphology and exudate availability are shaped by particle size and chemistry in. <i>Plant Direct</i> , <b>2020</b> , 4, e00207	3.3	8
140	Shed Light in the DaRk LineagES of the Fungal Tree of Life-STRES. <i>Life</i> , <b>2020</b> , 10,	3	4
139	Taxonomic and Metabolic Incongruence in the Ancient Genus. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 2170	5.7	23
138	EcoFABs: advancing microbiome science through standardized fabricated ecosystems. <i>Nature Methods</i> , <b>2019</b> , 16, 567-571	21.6	39

137	Probing the active fraction of soil microbiomes using BONCAT-FACS. <i>Nature Communications</i> , <b>2019</b> , 10, 2770	17.4	53
136	A High-Throughput Mass Spectrometric Enzyme Activity Assay Enabling the Discovery of Cytochrome P450 Biocatalysts. <i>Angewandte Chemie - International Edition</i> , <b>2019</b> , 58, 10114-10119	16.4	18
135	A High-Throughput Mass Spectrometric Enzyme Activity Assay Enabling the Discovery of Cytochrome P450 Biocatalysts. <i>Angewandte Chemie</i> , <b>2019</b> , 131, 10220-10225	3.6	3
134	Mediterranean grassland soil C-N compound turnover is dependent on rainfall and depth, and is mediated by genomically divergent microorganisms. <i>Nature Microbiology</i> , <b>2019</b> , 4, 1356-1367	26.6	70
133	Lessons from Two Design-Build-Test-Learn Cycles of Dodecanol Production in Escherichia coli Aided by Machine Learning. <i>ACS Synthetic Biology</i> , <b>2019</b> , 8, 1337-1351	5.7	53
132	MAGI: A Method for Metabolite Annotation and Gene Integration. <i>ACS Chemical Biology</i> , <b>2019</b> , 14, 704-714	14	19
131	A New Method to Correct for Habitat Filtering in Microbial Correlation Networks. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 585	5.7	11
130	Enzyme promiscuity shapes adaptation to novel growth substrates. <i>Molecular Systems Biology</i> , <b>2019</b> , 15, e8462	12.2	33
129	Regulation of Oxygenic Photosynthesis during Trophic Transitions in the Green Alga. <i>Plant Cell</i> , <b>2019</b> , 31, 579-601	11.6	35
128	Impacts of Maize Domestication and Breeding on Rhizosphere Microbial Community Recruitment from a Nutrient Depleted Agricultural Soil. <i>Scientific Reports</i> , <b>2019</b> , 9, 15611	4.9	37
127	Learning representations of microbe-metabolite interactions. <i>Nature Methods</i> , <b>2019</b> , 16, 1306-1314	21.6	79
126	Optimizing genome assembly from PCR-amplified metagenomes. <i>PeerJ</i> , <b>2019</b> , 7, e6902	3.1	14
125	Attracting and Retaining Top Scientists and Engineers at U.S. National Laboratories and Universities: Listening to the Next Generation. <i>Electrochemical Society Interface</i> , <b>2019</b> , 28, 34-36	3.6	
124	Characteristics of Wetting-Induced Bacteriophage Blooms in Biological Soil Crust. <i>MBio</i> , <b>2019</b> , 10,	7.8	23
123	Cooking shapes the structure and function of the gut microbiome. <i>Nature Microbiology</i> , <b>2019</b> , 4, 2052-2066	66	66
122	CRAGE enables rapid activation of biosynthetic gene clusters in undomesticated bacteria. <i>Nature Microbiology</i> , <b>2019</b> , 4, 2498-2510	26.6	48
121	Multilab EcoFAB study shows highly reproducible physiology and depletion of soil metabolites by a model grass. <i>New Phytologist</i> , <b>2019</b> , 222, 1149-1160	9.8	22
120	Untargeted Soil Metabolomics Using Liquid Chromatography-Mass Spectrometry and Gas Chromatography-Mass Spectrometry. <i>Methods in Molecular Biology</i> , <b>2019</b> , 1859, 97-109	1.4	5

119	Large Blooms of () Underlie the Response to Wetting of Cyanobacterial Biocrusts at Various Stages of Maturity. <i>MBio</i> , <b>2018</b> , 9,	7.8	16
118	Ecosystem Fabrication (EcoFAB) Protocols for The Construction of Laboratory Ecosystems Designed to Study Plant-microbe Interactions. <i>Journal of Visualized Experiments</i> , <b>2018</b> ,	1.6	20
117	Linking soil biology and chemistry in biological soil crust using isolate exometabolomics. <i>Nature Communications</i> , <b>2018</b> , 9, 19	17.4	83
116	A novel method to evaluate nutrient retention by biological soil crust exopolymeric matrix. <i>Plant and Soil</i> , <b>2018</b> , 429, 53-64	4.2	14
115	Dynamic root exudate chemistry and microbial substrate preferences drive patterns in rhizosphere microbial community assembly. <i>Nature Microbiology</i> , <b>2018</b> , 3, 470-480	26.6	623
114	Feed Your Friends: Do Plant Exudates Shape the Root Microbiome?. <i>Trends in Plant Science</i> , <b>2018</b> , 23, 25-41	13.1	655
113	Deciphering microbial interactions in synthetic human gut microbiome communities. <i>Molecular Systems Biology</i> , <b>2018</b> , 14, e8157	12.2	185
112	Insulator Nanostructure Desorption Ionization Mass Spectrometry. <i>Analytical Chemistry</i> , <b>2018</b> , 90, 9657-9661	9.6	1
111	Need for Laboratory Ecosystems To Unravel the Structures and Functions of Soil Microbial Communities Mediated by Chemistry. <i>MBio</i> , <b>2018</b> , 9,	7.8	23
110	Flux balance modeling to predict bacterial survival during pulsed-activity events. <i>Biogeosciences</i> , <b>2018</b> , 15, 2219-2229	4.6	5
109	Microbial Ecology on Solar Panels in Berkeley, CA, United States. <i>Frontiers in Microbiology</i> , <b>2018</b> , 9, 3043	5.7	10
108	Analysis and Interpretation of Mass Spectrometry Imaging Datasets. <i>Comprehensive Analytical Chemistry</i> , <b>2018</b> , 82, 369-386	1.9	1
107	Rapid characterization of the activities of lignin-modifying enzymes based on nanostructure-initiator mass spectrometry (NIMS). <i>Biotechnology for Biofuels</i> , <b>2018</b> , 11, 266	7.8	6
106	Web of microbes (WoM): a curated microbial exometabolomics database for linking chemistry and microbes. <i>BMC Microbiology</i> , <b>2018</b> , 18, 115	4.5	10
105	In Situ X-Ray Tomography Imaging of Soil Water and Cyanobacteria From Biological Soil Crusts Undergoing Desiccation. <i>Frontiers in Environmental Science</i> , <b>2018</b> , 6,	4.8	10
104	Extensive Turnover of Compatible Solutes in Cyanobacteria Revealed by Deuterium Oxide (DO) Stable Isotope Probing. <i>ACS Chemical Biology</i> , <b>2017</b> , 12, 674-681	4.9	18
103	Determination of glycoside hydrolase specificities during hydrolysis of plant cell walls using glycome profiling. <i>Biotechnology for Biofuels</i> , <b>2017</b> , 10, 31	7.8	13
102	Dynamic substrate preferences predict metabolic properties of a simple microbial consortium. <i>BMC Bioinformatics</i> , <b>2017</b> , 18, 57	3.6	16

101	OpenMSI Arrayed Analysis Toolkit: Analyzing Spatially Defined Samples Using Mass Spectrometry Imaging. <i>Analytical Chemistry</i> , <b>2017</b> , 89, 5818-5823	7.8	15
100	Morphology-Driven Control of Metabolite Selectivity Using Nanostructure-Initiator Mass Spectrometry. <i>Analytical Chemistry</i> , <b>2017</b> , 89, 6521-6526	7.8	15
99	On-chip integration of droplet microfluidics and nanostructure-initiator mass spectrometry for enzyme screening. <i>Lab on A Chip</i> , <b>2017</b> , 17, 323-331	7.2	33
98	Key Metabolites and Mechanistic Changes for Salt Tolerance in an Experimentally Evolved Sulfate-Reducing Bacterium,. <i>MBio</i> , <b>2017</b> , 8,	7.8	11
97	Exometabolomic Analysis of Cross-Feeding Metabolites. <i>Metabolites</i> , <b>2017</b> , 7,	5.6	8
96	Construction of Viable Soil Defined Media Using Quantitative Metabolomics Analysis of Soil Metabolites. <i>Frontiers in Microbiology</i> , <b>2017</b> , 8, 2618	5.7	12
95	Smartphone Analytics: Mobilizing the Lab into the Cloud for Omic-Scale Analyses. <i>Analytical Chemistry</i> , <b>2016</b> , 88, 9753-9758	7.8	13
94	Comparative Community Proteomics Demonstrates the Unexpected Importance of Actinobacterial Glycoside Hydrolase Family 12 Protein for Crystalline Cellulose Hydrolysis. <i>MBio</i> , <b>2016</b> , 7,	7.8	12
93	A robust gene-stacking method utilizing yeast assembly for plant synthetic biology. <i>Nature Communications</i> , <b>2016</b> , 7, 13215	17.4	42
92	Bacteria increase arid-land soil surface temperature through the production of sunscreens. <i>Nature Communications</i> , <b>2016</b> , 7, 10373	17.4	107
91	High-throughput platforms for metabolomics. <i>Current Opinion in Chemical Biology</i> , <b>2016</b> , 30, 7-13	9.7	52
90	Exometabolomics Assisted Design and Validation of Synthetic Obligate Mutualism. <i>ACS Synthetic Biology</i> , <b>2016</b> , 5, 569-76	5.7	18
89	Application of Black Silicon for Nanostructure-Initiator Mass Spectrometry. <i>Analytical Chemistry</i> , <b>2016</b> , 88, 1625-30	7.8	25
88	New insight into the role of MMP14 in metabolic balance. <i>PeerJ</i> , <b>2016</b> , 4, e2142	3.1	16
87	Proteogenomic analyses indicate bacterial methylotrophy and archaeal heterotrophy are prevalent below the grass root zone. <i>PeerJ</i> , <b>2016</b> , 4, e2687	3.1	72
86	Belowground Response to Drought in a Tropical Forest Soil. II. Change in Microbial Function Impacts Carbon Composition. <i>Frontiers in Microbiology</i> , <b>2016</b> , 7, 323	5.7	37
85	Belowground Response to Drought in a Tropical Forest Soil. I. Changes in Microbial Functional Potential and Metabolism. <i>Frontiers in Microbiology</i> , <b>2016</b> , 7, 525	5.7	63
84	Exometabolomics for Linking Soil Carbon Dynamics to Microbial Communities <b>2016</b> , 119-145		

83	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. <i>Nature Biotechnology</i> , <b>2016</b> , 34, 828-837	44.5	1566
82	Gut microbiota mediate caffeine detoxification in the primary insect pest of coffee. <i>Nature Communications</i> , <b>2015</b> , 6, 7618	17.4	194
81	Identification of metabolic signatures linked to anti-inflammatory effects of <i>Faecalibacterium prausnitzii</i> . <i>MBio</i> , <b>2015</b> , 6,	7.8	128
80	Isolation of a significant fraction of non-phototroph diversity from a desert Biological Soil Crust. <i>Frontiers in Microbiology</i> , <b>2015</b> , 6, 277	5.7	37
79	Unraveling heterogeneous susceptibility and the evolution of breast cancer using a systems biology approach. <i>Genome Biology</i> , <b>2015</b> , 16, 40	18.3	11
78	Exometabolomics and MSI: deconstructing how cells interact to transform their small molecule environment. <i>Current Opinion in Biotechnology</i> , <b>2015</b> , 34, 209-16	11.4	40
77	Competitive sorption of microbial metabolites on an iron oxide mineral. <i>Soil Biology and Biochemistry</i> , <b>2015</b> , 90, 34-41	7.5	28
76	Untargeted soil metabolomics methods for analysis of extractable organic matter. <i>Soil Biology and Biochemistry</i> , <b>2015</b> , 80, 189-198	7.5	87
75	High throughput screening of enzyme activity with mass spectrometry imaging. <i>Current Opinion in Biotechnology</i> , <b>2015</b> , 31, 1-9	11.4	41
74	Lineage-specific chromatin signatures reveal a regulator of lipid metabolism in microalgae. <i>Nature Plants</i> , <b>2015</b> , 1, 15107	11.5	60
73	Exometabolite niche partitioning among sympatric soil bacteria. <i>Nature Communications</i> , <b>2015</b> , 6, 8289	17.4	120
72	Development of a High Throughput Platform for Screening Glycoside Hydrolases Based on Oxime-NIMS. <i>Frontiers in Bioengineering and Biotechnology</i> , <b>2015</b> , 3, 153	5.8	14
71	Use of Nanostructure-Initiator Mass Spectrometry to Deduce Selectivity of Reaction in Glycoside Hydrolases. <i>Frontiers in Bioengineering and Biotechnology</i> , <b>2015</b> , 3, 165	5.8	4
70	Analysis of Metabolomics Datasets with High-Performance Computing and Metabolite Atlases. <i>Metabolites</i> , <b>2015</b> , 5, 431-42	5.6	28
69	Multifunctional cellulase catalysis targeted by fusion to different carbohydrate-binding modules. <i>Biotechnology for Biofuels</i> , <b>2015</b> , 8, 220	7.8	38
68	Nanostructure imaging mass spectrometry: the role of fluorocarbons in metabolite analysis and yoctomole level sensitivity. <i>Methods in Molecular Biology</i> , <b>2015</b> , 1203, 141-9	1.4	8
67	Rapid kinetic characterization of glycosyl hydrolases based on oxime derivatization and nanostructure-initiator mass spectrometry (NIMS). <i>ACS Chemical Biology</i> , <b>2014</b> , 9, 1470-9	4.9	30
66	Phylogenomically guided identification of industrially relevant GH1 $\beta$ glucosidases through DNA synthesis and nanostructure-initiator mass spectrometry. <i>ACS Chemical Biology</i> , <b>2014</b> , 9, 2082-91	4.9	45

65	Noninvasive Mapping of Photosynthetic Heterogeneity in Biological Soil Crusts by Positron Emission Tomography: Carbon Fixation. <i>Environmental Science and Technology Letters</i> , <b>2014</b> , 1, 393-398	11	3
64	The lethal cargo of <i>Myxococcus xanthus</i> outer membrane vesicles. <i>Frontiers in Microbiology</i> , <b>2014</b> , 5, 474	5.7	86
63	Metabolic imaging using nanostructure-initiator mass spectrometry (NIMS). <i>Methods in Molecular Biology</i> , <b>2014</b> , 1198, 313-29	1.4	6
62	Versatile synthesis of probes for high-throughput enzyme activity screening. <i>Analytical and Bioanalytical Chemistry</i> , <b>2013</b> , 405, 4969-73	4.4	14
61	Dynamic cyanobacterial response to hydration and dehydration in a desert biological soil crust. <i>ISME Journal</i> , <b>2013</b> , 7, 2178-91	11.9	156
60	Robust automated mass spectra interpretation and chemical formula calculation using mixed integer linear programming. <i>Analytical Chemistry</i> , <b>2013</b> , 85, 9777-84	7.8	6
59	Mass spectrometry imaging for in situ kinetic histochemistry. <i>Scientific Reports</i> , <b>2013</b> , 3, 1656	4.9	47
58	Stable-isotope probing reveals that hydrogen isotope fractionation in proteins and lipids in a microbial community are different and species-specific. <i>ACS Chemical Biology</i> , <b>2013</b> , 8, 1755-63	4.9	24
57	"Replica-extraction-transfer" nanostructure-initiator mass spectrometry imaging of acoustically printed bacteria. <i>Analytical Chemistry</i> , <b>2013</b> , 85, 10856-62	7.8	36
56	Metabolic footprinting of mutant libraries to map metabolite utilization to genotype. <i>ACS Chemical Biology</i> , <b>2013</b> , 8, 189-99	4.9	23
55	OpenMSI: a high-performance web-based platform for mass spectrometry imaging. <i>Analytical Chemistry</i> , <b>2013</b> , 85, 10354-61	7.8	68
54	Metabolites associated with adaptation of microorganisms to an acidophilic, metal-rich environment identified by stable-isotope-enabled metabolomics. <i>MBio</i> , <b>2013</b> , 4, e00484-12	7.8	63
53	Lipids as tumoricidal components of human $\alpha$ 2-macroglobulin made lethal to tumor cells (HAMLET): unique and shared effects on signaling and death. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 17460-71	5.4	18
52	From soil to structure, a novel dimeric $\beta$ -glucosidase belonging to glycoside hydrolase family 3 isolated from compost using metagenomic analysis. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 14985-92	5.4	35
51	High throughput nanostructure-initiator mass spectrometry screening of microbial growth conditions for maximal $\beta$ -glucosidase production. <i>Frontiers in Microbiology</i> , <b>2013</b> , 4, 365	5.7	11
50	Functional genomics of novel secondary metabolites from diverse cyanobacteria using untargeted metabolomics. <i>Marine Drugs</i> , <b>2013</b> , 11, 3617-31	6	45
49	Cyanobacteria as Biocatalysts for Carbonate Mineralization. <i>Minerals (Basel, Switzerland)</i> , <b>2012</b> , 2, 338-364	6.4	83
48	Resolving brain regions using nanostructure initiator mass spectrometry imaging of phospholipids. <i>Integrative Biology (United Kingdom)</i> , <b>2012</b> , 4, 693-9	3.7	32



47	Low-dose ionizing radiation-induced blood plasma metabolic response in a diverse genetic mouse population. <i>Radiation Research</i> , <b>2012</b> , 178, 551-5	3.1	17
46	Encoding substrates with mass tags to resolve stereospecific reactions using Nimzyme. <i>Rapid Communications in Mass Spectrometry</i> , <b>2012</b> , 26, 611-5	2.2	18
45	Deuterium-exchange metabolomics identifies N-methyl lyso phosphatidylethanolamines as abundant lipids in acidophilic mixed microbial communities. <i>Metabolomics</i> , <b>2012</b> , 8, 566-578	4.7	12
44	Acoustic deposition with NIMS as a high-throughput enzyme activity assay. <i>Analytical and Bioanalytical Chemistry</i> , <b>2012</b> , 403, 707-11	4.4	30
43	Retinoic acid induces a metabolic switch in SH-SY5Y cells from glycolysis to oxidative phosphorylation. <i>FASEB Journal</i> , <b>2012</b> , 26, 967.1	0.9	
42	Nanostructure-Initiator Mass Spectrometry (NIMS) for the Analysis of Enzyme Activities. <i>Current Protocols in Chemical Biology</i> , <b>2012</b> , 4, 123-142	1.8	3
41	Conserved features of cancer cells define their sensitivity to HAMLET-induced death; c-Myc and glycolysis. <i>Oncogene</i> , <b>2011</b> , 30, 4765-79	9.2	31
40	The Small-Molecule Dimension: Mass-Spectrometry-Based Metabolomics, Enzyme Assays, and Imaging <b>2011</b> , 675-682		
39	Improved genome annotation through untargeted detection of pathway-specific metabolites. <i>BMC Genomics</i> , <b>2011</b> , 12 Suppl 1, S6	4.5	12
38	Untargeted metabolic footprinting reveals a surprising breadth of metabolite uptake and release by <i>Synechococcus</i> sp. PCC 7002. <i>Molecular BioSystems</i> , <b>2011</b> , 7, 3200-6		40
37	Meta-analysis of global metabolomics and proteomics data to link alterations with phenotype. <i>Spectroscopy</i> , <b>2011</b> , 26, 151-154		
36	Colloid-based multiplexed screening for plant biomass-degrading glycoside hydrolase activities in microbial communities. <i>Energy and Environmental Science</i> , <b>2011</b> , 4, 2884	35.4	28
35	Multivariate analysis of a 3D mass spectral image for examining tissue heterogeneity. <i>Integrative Biology (United Kingdom)</i> , <b>2011</b> , 3, 460-7	3.7	17
34	Mass spectrometry-based metabolomics, analysis of metabolite-protein interactions, and imaging. <i>BioTechniques</i> , <b>2010</b> , 49, 557-65	2.5	51
33	Metabolome-proteome differentiation coupled to microbial divergence. <i>MBio</i> , <b>2010</b> , 1,	7.8	26
32	Rapid screening of fatty acids using nanostructure-initiator mass spectrometry. <i>Analytical Chemistry</i> , <b>2010</b> , 82, 3751-5	7.8	28
31	Metabolite identification in <i>Synechococcus</i> sp. PCC 7002 using untargeted stable isotope assisted metabolite profiling. <i>Analytical Chemistry</i> , <b>2010</b> , 82, 9034-42	7.8	59
30	Calcifying cyanobacteria--the potential of biomineralization for carbon capture and storage. <i>Current Opinion in Biotechnology</i> , <b>2010</b> , 21, 365-71	11.4	124

29	Large scale physiological readjustment during growth enables rapid, comprehensive and inexpensive systems analysis. <i>BMC Systems Biology</i> , <b>2010</b> , 4, 64	3.5	21
28	Dealing with the unknown: metabolomics and metabolite atlases. <i>Journal of the American Society for Mass Spectrometry</i> , <b>2010</b> , 21, 1471-6	3.5	135
27	Phosphonium labeling for increasing metabolomic coverage of neutral lipids using electrospray ionization mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , <b>2009</b> , 23, 1849-55	2.2	20
26	Mass spectrometry based metabolomics and enzymatic assays for functional genomics. <i>Current Opinion in Microbiology</i> , <b>2009</b> , 12, 547-52	7.9	57
25	Nanostructure initiator mass spectrometry: tissue imaging and direct biofluid analysis. <i>Analytical Chemistry</i> , <b>2009</b> , 81, 2969-75	7.8	110
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16	Biofuels and biomaterials from microbes315-335		
15	Cervicovaginal microbiome composition drives metabolic profiles in healthy pregnancy		2
14	Linking soil biology and chemistry using bacterial isolate exometabolite profiles		1
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11	Study of Oak Ridge soils using BONCAT-FACS-Seq reveals that a large fraction of the soil microbiome is active	3
10	Multi-lab EcoFAB study shows highly reproducible physiology and depletion of soil metabolites by a model grass	3
9	Processing of grassland soil C-N compounds into soluble and volatile molecules is depth stratified and mediated by genomically novel bacteria and archaea	4
8	A New Method to Correct for Habitat Filtering in Microbial Correlation Networks	1
7	Physiological adaptations of leaf litter microbial communities to long-term drought	3
6	Root morphology and exudate availability is shaped by particle size and chemistry in <i>Brachypodium distachyon</i>	2
5	Dynamic substrate preferences and predicted metabolic properties of a simple microbial consortium	1
4	GNPS Dashboard: Collaborative Analysis of Mass Spectrometry Data in the Web Browser	2
3	Long-read metagenomics of soil communities reveals phylum-specific secondary metabolite dynamics	2
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