Robert Winkler

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.

99 2,208 25 44 g-index

125 2,660 5.1 5.12
ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
99	MoBiMS: A modular miniature mass analyzer for the real-time monitoring of gases and volatile compounds in biological systems. <i>Microchemical Journal</i> , 2022 , 175, 107090	4.8	1
98	Loss of Sensory Cup Quality: Physiological and Chemical Changes during Green Coffee Storage <i>Plant Foods for Human Nutrition</i> , 2022 , 77, 1	3.9	1
97	Analyzing the Distribution of Specialized Metabolites from Plant Native Tissues with Laser Desorption Low-Temperature Plasma Mass Spectrometry Imaging <i>Methods in Molecular Biology</i> , 2022 , 2469, 145-154	1.4	
96	Mass Fingerprinting for High-Throughput Analyses of Food: Authentication and Quality Control 2022 , 3-16		
95	Evaluation of Metabolomic Profile and Growth of Moringa oleifera L. Cultivated with Vermicompost under Different Soil Types. <i>Agronomy</i> , 2021 , 11, 2061	3.6	O
94	CHD1 controls H3.3 incorporation in adult brain chromatin to maintain metabolic homeostasis and normal lifespan. <i>Cell Reports</i> , 2021 , 37, 109769	10.6	0
93	Novel UV filters from Pentacalia pulchella extracts with photoprotective properties and antioxidant activity. <i>Photochemical and Photobiological Sciences</i> , 2021 , 20, 1585-1597	4.2	2
92	Effects of the Developmental Regulator BOLITA on the Plant Metabolome. <i>Genes</i> , 2021 , 12,	4.2	1
91	Contrast optimization of mass spectrometry imaging (MSI) data visualization by threshold intensity quantization (TrIQ). <i>PeerJ Computer Science</i> , 2021 , 7, e585	2.7	
90	ProtyQuant: Comparing label-free shotgun proteomics datasets using accumulated peptide probabilities. <i>Journal of Proteomics</i> , 2021 , 230, 103985	3.9	0
89	The emerging role of 3D-printing in ion mobility spectrometry and mass spectrometry. <i>Analytical Methods</i> , 2021 , 13, 852-861	3.2	5
88	MeteoMex: open infrastructure for networked environmental monitoring and agriculture 4.0. <i>PeerJ Computer Science</i> , 2021 , 7, e343	2.7	4
87	Protein extract of Bromelia karatas L. rich in cysteine proteases (ananain- and bromelain-like) has antibacterial activity against foodborne pathogens Listeria monocytogenes and Salmonella Typhimurium. <i>Folia Microbiologica</i> , 2021 , 1	2.8	O
86	Metabolic response to larval herbivory in three species. <i>Plant Signaling and Behavior</i> , 2021 , 1962050	2.5	0
85	Modulation of Aleurone Peroxidases in Kernels of Insect-Resistant Maize (L.; Pob84-C3R) After Mechanical and Insect Damage. <i>Frontiers in Plant Science</i> , 2020 , 11, 781	6.2	1
84	Prediction of the antioxidant capacity of maize (Zea mays) hybrids using mass fingerprinting and data mining. <i>Food Bioscience</i> , 2020 , 37, 100647	4.9	1
83	Mass spectrometry imaging of thin-layer chromatography plates using laser desorption/low-temperature plasma ionisation. <i>Analyst, The</i> , 2020 , 145, 3885-3891	5	8

(2018-2020)

82	Distinct gene expression and secondary metabolite profiles in tomato mutants having impaired mycorrhizal colonization. <i>PeerJ</i> , 2020 , 8, e8888	3.1	2
81	Proteomic analysis and interactions network in leaves of mycorrhizal and nonmycorrhizal sorghum plants under water deficit. <i>PeerJ</i> , 2020 , 8, e8991	3.1	4
80	Metabolomic effects of the colonization of Medicago truncatula by the facultative endophyte Arthrobacter agilis UMCV2 in a foliar inoculation system. <i>Scientific Reports</i> , 2020 , 10, 8426	4.9	10
79	Open LabBot and RmsiGUI: Community development kit for sampling automation and ambient imaging. <i>Microchemical Journal</i> , 2020 , 152, 104343	4.8	9
78	Mass Fingerprints of Tomatoes Fertilized with Different Nitrogen Sources Reveal Potential Biomarkers of Organic Farming. <i>Plant Foods for Human Nutrition</i> , 2019 , 74, 247-254	3.9	3
77	Software solutions for evaluation and visualization of laser ablation inductively coupled plasma mass spectrometry imaging (LA-ICP-MSI) data: a short overview. <i>Journal of Cheminformatics</i> , 2019 , 11, 16	8.6	21
76	Plasma protein adsorption on FeO-PEG nanoparticles activates the complement system and induces an inflammatory response. <i>International Journal of Nanomedicine</i> , 2019 , 14, 2055-2067	7.3	20
75	Screening for Green Coffee with Sensorial Defects Due to Aging During Storage by MALDI-ToF Mass Fingerprinting. <i>Food Analytical Methods</i> , 2019 , 12, 1571-1576	3.4	3
74	A Mass Spectrometry-Based Study Shows that Volatiles Emitted by UMCV2 Increase the Content of Brassinosteroids in in Response to Iron Deficiency Stress. <i>Molecules</i> , 2019 , 24,	4.8	3
73	Effects of Water Availability in the Soil on Tropane Alkaloid Production in Cultivated. <i>Metabolites</i> , 2019 , 9,	5.6	6
72	Metabolomic Markers for the Early Selection of Plants with Desirable Cup Quality Traits. <i>Metabolites</i> , 2019 , 9,	5.6	10
71	Low-Temperature Plasma Ionization 2019 , 105-124		O
70	from Predator to Prey: Role of the Mitogen-Activated Protein Kinase Tmk3 in Fungal Chemical Defense against Fungivory by Larvae. <i>Applied and Environmental Microbiology</i> , 2019 , 85,	4.8	11
69	Elucidating the Distribution of Plant Metabolites from Native Tissues with Laser Desorption Low-Temperature Plasma Mass Spectrometry Imaging. <i>Analytical Chemistry</i> , 2019 , 91, 2734-2743	7.8	26
68	Increase of peroxidase activity in tropical maize after recurrent selection to storage pest resistance. <i>Journal of Stored Products Research</i> , 2018 , 75, 47-55	2.5	5
67	In vivo monitoring of nicotine biosynthesis in tobacco leaves by low-temperature plasma mass spectrometry. <i>Talanta</i> , 2018 , 185, 324-327	6.2	9
66	Postharvest insect resistance in maize. Journal of Stored Products Research, 2018, 77, 66-76	2.5	19
65	Funaria hygrometrica Hedw. elevated tolerance to DO: its use for the production of highly deuterated metabolites. <i>Planta</i> , 2018 , 247, 405-412	4.7	3

64	In vitro UV absorption properties and radical scavenging capacity of Morella parvifolia (Benth.) Parra-Os. extracts. <i>Brazilian Journal of Pharmaceutical Sciences</i> , 2018 , 54,	1.8	4
63	Modulation of steroidal glycoalkaloid biosynthesis in tomato (Solanum lycopersicum) by jasmonic acid. <i>Plant Science</i> , 2018 , 277, 155-165	5.3	14
62	The organophosphate pesticide methamidophos opens the blood-testis barrier and covalently binds to ZO-2 in mice. <i>Toxicology and Applied Pharmacology</i> , 2018 , 360, 257-272	4.6	16
61	Down-regulation of PvTRX1h increases nodule number and affects auxin, starch, and metabolic fingerprints in the common bean (Phaseolus vulgaris L.). <i>Plant Science</i> , 2018 , 274, 45-58	5.3	9
60	Low-temperature plasma (LTP) jets for mass spectrometry (MS): Ion processes, instrumental set-ups, and application examples. <i>TrAC - Trends in Analytical Chemistry</i> , 2017 , 89, 133-145	14.6	32
59	Automated chemical fingerprinting of Mexican spirits derived from Agave (tequila and mezcal) using direct-injection electrospray ionisation (DIESI) and low-temperature plasma (LTP) mass spectrometry. <i>Analytical Methods</i> , 2017 , 9, 5023-5028	3.2	19
58	Lipidomic profiles of Drosophila melanogaster and cactophilic fly species: models of human metabolic diseases. <i>Integrative Biology (United Kingdom)</i> , 2017 , 9, 885-891	3.7	4
57	A comparison of the human and mouse protein corona profiles of functionalized SiO nanocarriers. <i>Nanoscale</i> , 2017 , 9, 13651-13660	7.7	36
56	Genomic history of the origin and domestication of common bean unveils its closest sister species. <i>Genome Biology</i> , 2017 , 18, 60	18.3	79
55	Rapid Classification of Coffee Products by Data Mining Models from Direct Electrospray and Plasma-Based Mass Spectrometry Analyses. <i>Food Analytical Methods</i> , 2017 , 10, 1359-1368	3.4	20
54	The Flavonoid Fraction from Rhoeo discolor Leaves Acts Antiviral Against Influenza A Virus. <i>Records of Natural Products</i> , 2017 , 11, 532-546	1.9	10
53	Improved Representation of Biological Information by Using Correlation as Distance Function for Heatmap Cluster Analysis. <i>American Journal of Plant Sciences</i> , 2017 , 08, 502-516	0.5	11
52	Photoprotective Potential of Baccharis antioquensis (Asteraceae) as Natural Sunscreen. <i>Photochemistry and Photobiology</i> , 2016 , 92, 742-52	3.6	17
51	Template for 3D Printing a Low-Temperature Plasma Probe. <i>Analytical Chemistry</i> , 2016 , 88, 6976-80	7.8	40
50	Popper and the Omics. Frontiers in Plant Science, 2016 , 7, 195	6.2	2
49	Structural Basis for Redox Regulation of Cytoplasmic and Chloroplastic Triosephosphate Isomerases from. <i>Frontiers in Plant Science</i> , 2016 , 7, 1817	6.2	17
48	MSI.R scripts reveal volatile and semi-volatile features in low-temperature plasma mass spectrometry imaging (LTP-MSI) of chilli (Capsicum annuum). <i>Analytical and Bioanalytical Chemistry</i> , 2015 , 407, 5673-84	4.4	17
47	SpiderMass: Semantic database creation and tripartite metabolite identification strategy. <i>Journal of Mass Spectrometry</i> , 2015 , 50, 538-41	2.2	13

(2012-2015)

GridMass: a fast two-dimensional feature detection method for LC/MS. <i>Journal of Mass Spectrometry</i> , 2015 , 50, 165-74	2.2	34	
Metabolic fingerprinting of Arabidopsis thaliana accessions. <i>Frontiers in Plant Science</i> , 2015 , 6, 365	6.2	13	
The Phaseolus vulgaris PvTRX1h gene regulates plant hormone biosynthesis in embryogenic callus from common bean. <i>Frontiers in Plant Science</i> , 2015 , 6, 577	6.2	16	
Identification of B6T173 (ZmPrx35) as the prevailing peroxidase in highly insect-resistant maize (Zea mays, p84C3) kernels by activity-directed purification. <i>Frontiers in Plant Science</i> , 2015 , 6, 670	6.2	7	
Metabolic profiling of plant extracts using direct-injection electrospray ionization mass spectrometry allows for high-throughput phenotypic characterization according to genetic and environmental effects. <i>Journal of Agricultural and Food Chemistry</i> , 2015 , 63, 1042-52	5.7	25	
Reduction of aflatoxin B1 during tortilla production and identification of degradation by-products by direct-injection electrospray mass spectrometry (DIESI-MS). <i>Salud Publica De Mexico</i> , 2015 , 57, 50-7	1.7	9	
An evolving computational platform for biological mass spectrometry: workflows, statistics and data mining with MASSyPup64. <i>PeerJ</i> , 2015 , 3, e1401	3.1	16	
Preventive and therapeutic potential of peptides from cereals against cancer. <i>Journal of Proteomics</i> , 2014 , 111, 165-83	3.9	70	
MASSyPupan Tout of the boxTsolution for the analysis of mass spectrometry data. <i>Journal of Mass Spectrometry</i> , 2014 , 49, 37-42	2.2	14	
Plug and PlayTassembly of a low-temperature plasma ionization mass spectrometry imaging (LTP-MSI) system. <i>Journal of Proteomics</i> , 2014 , 102, 60-5	3.9	36	
Proteomic analysis of Entamoeba histolytica in vivo assembled pre-mRNA splicing complexes. Journal of Proteomics, 2014 , 111, 30-45	3.9	10	
Vinylogous chain branching catalysed by a dedicated polyketide synthase module. <i>Nature</i> , 2013 , 502, 124-8	50.4	93	
Design of a low-temperature plasma (LTP) probe with adjustable output temperature and variable beam diameter for the direct detection of organic molecules. <i>Rapid Communications in Mass Spectrometry</i> , 2013 , 27, 629-34	2.2	25	
Lysine-directed staining of proteins for MS-based analyses. <i>Electrophoresis</i> , 2013 , 34, 401-4	3.6	1	
Metabolic phenotyping for the classification of coffee trees and the exploration of selection markers. <i>Molecular BioSystems</i> , 2013 , 9, 693-9		25	
Phenotypic comparison of samdc and spe mutants reveals complex relationships of polyamine metabolism in Ustilago maydis. <i>Microbiology (United Kingdom)</i> , 2012 , 158, 674-684	2.9	10	
Fungal Earabinofuranosidases of glycosyl hydrolase families 51 and 54 show a dual arabinofuranosyl- and galactofuranosyl-hydrolyzing activity. <i>Biological Chemistry</i> , 2012 , 393, 767-75	4.5	10	
Evaluating the physiological state of maize (Zea mays L.) plants by direct-injection electrospray mass spectrometry (DIESI-MS). <i>Molecular BioSystems</i> , 2012 , 8, 1658-60		19	
	Metabolic fingerprinting of Arabidopsis thaliana accessions. Frontiers in Plant Science, 2015, 6, 365 The Phaseolus vulgaris PVTRX1h gene regulates plant hormone biosynthesis in embryogenic callus from common bean. Frontiers in Plant Science, 2015, 6, 577 Identification of B6T173 (ZmPrx35) as the prevailing peroxidase in highly insect-resistant maize (Zea mays, p84C3) kernels by activity-directed purification. Frontiers in Plant Science, 2015, 6, 670 Metabolic profiling of plant extracts using direct-injection electrospray ionization mass spectrometry allows for high-throughput phenotypic characterization according to genetic and environmental effects. Journal of Agricultural and Food Chemistry, 2015, 63, 1042-52 Reduction of aflatoxin B1 during tortilla production and identification of degradation by-products by direct-injection electrospray mass spectrometry (DIESI-MS). Salud Publica De Mexica, 2015, 57, 50-7 An evolving computational platform for biological mass spectrometry: workflows, statistics and data mining with MASSyPup64. Peer.J, 2015, 3, e1401 Preventive and therapeutic potential of peptides from cereals against cancer. Journal of Proteomics, 2014, 111, 165-83 MASSyPup-an Tout of the boxTsolution for the analysis of mass spectrometry data. Journal of Mass Spectrometry, 2014, 49, 37-42 Plug and PlayTassembly of a low-temperature plasma ionization mass spectrometry imaging (LTP-MSI) system. Journal of Proteomics, 2014, 102, 60-5 Proteomic analysis of Entamoeba histolytica in vivo assembled pre-mRNA splicing complexes. Journal of Proteomics, 2014, 111, 30-45 Vinylogous chain branching catalysed by a dedicated polyketide synthase module. Nature, 2013, 502, 124-8 Design of a low-temperature plasma (LTP) probe with adjustable output temperature and variable beam diameter for the direct detection of organic molecules. Rapid Communications in Mass Spectrometry, 2013, 27, 629-34 Lysine-directed staining of proteins for MS-based analyses. Electrophoresis, 2013, 34, 401-4 Metabolic phenotyping for	Metabolic Fingerprinting of Arabidopsis thaliana accessions. Frontiers in Plant Science, 2015, 6, 365 6.2 The Phaseolus vulgaris PVTRX1h gene regulates plant hormone biosynthesis in embryogenic callus from common bean. Frontiers in Plant Science, 2015, 6, 577 Identification of B6T173 (ZmPrx35) as the prevailing peroxidase in highly insect-resistant maize (Zea mays, p84G3) kernels by activity-directed purification. Frontiers in Plant Science, 2015, 6, 670 Metabolic profiling of plant extracts using direct-injection electrospray ionization mass spectrometry allows for high-throughput phenotypic characterization according to genetic and environmental effects. Journal of Agricultural and Food Chemistry, 2015, 63, 1042-52 Reduction of aflatoxin B1 during tortilla production and identification of degradation by-products by direct-injection electrospray mass spectrometry (DIESI-MS). Salud Publica De Mexico, 2015, 57, 50-7 An evolving computational platform for biological mass spectrometry: workflows, statistics and data mining with MASSyPup64. PeerJ, 2015, 3, e1401 Preventive and therapeutic potential of peptides from cereals against cancer. Journal of Proteomics, 2014, 111, 165-83 MASSyPup-an Dut of the boxTsolution for the analysis of mass spectrometry data. Journal of Mass Spectrometry, 2014, 49, 37-42 Plug and PlayTassembly of a low-temperature plasma ionization mass spectrometry imaging (LTP-MSI) system. Journal of Proteomics, 2014, 102, 60-5 Proteomic analysis of Entamoeba histolytica in vivo assembled pre-mRNA splicing complexes. Journal of Proteomics, 2014, 111, 30-45 Vinylogous chain branching catalysed by a dedicated polyketide synthase module. Nature, 2013, 50-4 Design of a low-temperature plasma (LTP) probe with adjustable output temperature and variable beam diameter for the direct detection of organic molecules. Rapid Communications in Mass Spectrometry, 2013, 27, 629-34 Lysine-directed staining of proteins for MS-based analyses. Electrophoresis, 2013, 34, 401-4 Metabolic phenotyping for th	Spectrometry, 2015, 50, 165-74 Metabolic fingerprinting of Arabidopsis thaliana accessions. Frontiers in Plant Science, 2015, 6, 365 The Phaseolus vulgaris PVTRX1h gene regulates plant hormone biosynthesis in embryogenic callus from common bean. Frontiers in Plant Science, 2015, 6, 570 Identification of B6T173 (ZmPrx35) as the prevailing peroxidase in highly insect-resistant maize (Zea mays, BaC3) kernels by activity-directed purification. Frontiers in Plant Science, 2015, 6, 670 Metabolic profiling of plant extracts using direct-injection electrospray ionization mass spectrometry allows for high-throughput phenotypic characterization according to genetic and environmental effects. Journal of Agricultural and Food Chemistry, 2015, 6, 1042-52 Reduction of aflatoxin B1 during tortilla production and identification of degradation by-products by direct-injection electrospray mass spectrometry (DIESHMS). Solud Publica De Mexico, 2015, 57, 50-7 An evolving computational platform for biological mass spectrometry: workflows, statistics and data mining with MASSyPup64. PeerJ, 2015, 3, e1401 3.1 16 Preventive and therapeutic potential of peptides from cereals against cancer. Journal of Proteomics 2014, 111, 165-83 MASSyPup-an Tout of the boxTsolution for the analysis of mass spectrometry data. Journal of Mass Spectrometry, 2014, 49, 37-42 Flug and PlayTassembly of a low-temperature plasma ionization mass spectrometry imaging (LTP-MS) system. Journal of Proteomics, 2014, 111, 30-45 Vinylogous chain branching catalysed by a dedicated polyketide synthase module. Nature, 2013, 50-2 102-124-8 Design of a low-temperature plasma (LTP) probe with adjustable output temperature and variable beam diameter for the direct detection of organic molecules. Rapid Communications in Mass 50-2, 124-8 Design of a low-temperature plasma (LTP) probe with adjustable output temperature and variable beam diameter for the direct detection of organic molecules. Rapid Communications in Mass 50-2, 124-18 Design of a low-temperature pla

28	Proteome analysis of the farnesol-induced stress response in Aspergillus nidulansThe role of a putative dehydrin. <i>Journal of Proteomics</i> , 2012 , 75, 4038-49	3.9	21
27	Accelerated identification of proteins by mass spectrometry by employing covalent pre-gel staining with Uniblue A. <i>PLoS ONE</i> , 2012 , 7, e31438	3.7	7
26	Comparative and functional genomics provide insights into the pathogenicity of dermatophytic fungi. <i>Genome Biology</i> , 2011 , 12, R7	18.3	147
25	Mass spectometry-based protein patterns in the diagnosis of sepsis/systemic inflammatory response syndrome. <i>Shock</i> , 2011 , 36, 560-9	3.4	16
24	Colorimetric protein quantification in aqueous two-phase systems. <i>Process Biochemistry</i> , 2011 , 46, 413-4	14.7 8	20
23	A ribonucleotide reductase-like electron transfer system in the nitroaryl-forming N-oxygenase AurF. <i>ChemBioChem</i> , 2011 , 12, 1832-5	3.8	9
22	Analysis of the Aspergillus fumigatus proteome reveals metabolic changes and the activation of the pseurotin A biosynthesis gene cluster in response to hypoxia. <i>Journal of Proteome Research</i> , 2011 , 10, 2508-24	5.6	103
21	An FMN hydrolase of the haloacid dehalogenase superfamily is active in plant chloroplasts. <i>Journal of Biological Chemistry</i> , 2011 , 286, 42091-42098	5.4	14
20	Activity-directed identification of maize kernel peroxidases associated with postharvest insect resistance. <i>Molecular BioSystems</i> , 2010 , 6, 1810-2		11
19	Structural and biochemical basis for the firm chemo- and regioselectivity of the nitro-forming N-oxygenase AurF. <i>Chemical Communications</i> , 2010 , 46, 7760-2	5.8	6
18	ESIprot: a universal tool for charge state determination and molecular weight calculation of proteins from electrospray ionization mass spectrometry data. <i>Rapid Communications in Mass Spectrometry</i> , 2010 , 24, 285-94	2.2	48
17	A novel immune evasion strategy of candida albicans: proteolytic cleavage of a salivary antimicrobial peptide. <i>PLoS ONE</i> , 2009 , 4, e5039	3.7	100
16	Production of pyomelanin, a second type of melanin, via the tyrosine degradation pathway in Aspergillus fumigatus. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 493-503	4.8	158
15	Two-dimensional proteome reference maps for the human pathogenic filamentous fungus Aspergillus fumigatus. <i>Proteomics</i> , 2009 , 9, 1407-15	4.8	66
14	Proteomic analysis of human papillomavirus-related oral squamous cell carcinoma: identification of thioredoxin and epidermal-fatty acid binding protein as upregulated protein markers in microdissected tumor tissue. <i>Proteomics</i> , 2009 , 9, 2193-201	4.8	27
13	Global distribution and evolution of a toxinogenic Burkholderia-Rhizopus symbiosis. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 2982-6	4.8	65
12	Evolution of host resistance in a toxin-producing bacterial-fungal alliance. ISME Journal, 2008, 2, 632-41	11.9	66
11	Biosynthesis of nitro compounds. <i>ChemBioChem</i> , 2007 , 8, 973-7	3.8	75

LIST OF PUBLICATIONS

10	A binuclear manganese cluster that catalyzes radical-mediated N-oxygenation. <i>Angewandte Chemie - International Edition</i> , 2007 , 46, 8605-8	16.4	28
9	A Binuclear Manganese Cluster That Catalyzes Radical-Mediated N-Oxygenation. <i>Angewandte Chemie</i> , 2007 , 119, 8759-8762	3.6	6
8	Monitoring phosphatase reactions of multiple phosphorylated substrates by reversed-phase HPLC. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2007, 853, 204-	33 ²	3
7	Structure and action of the N-oxygenase AurF from Streptomyces thioluteus. <i>Journal of Molecular Biology</i> , 2007 , 373, 65-74	6.5	44
6	Regio- and chemoselective enzymatic N-oxygenation in vivo, in vitro, and in flow. <i>Angewandte Chemie - International Edition</i> , 2006 , 45, 8016-8	16.4	39
5	Regio- and Chemoselective Enzymatic N-Oxygenation In Vivo, In Vitro, and in Flow. <i>Angewandte Chemie</i> , 2006 , 118, 8184-8186	3.6	14
4	The expression of selected non-ribosomal peptide synthetases in Aspergillus fumigatus is controlled by the availability of free iron. <i>FEMS Microbiology Letters</i> , 2005 , 248, 83-91	2.9	42
3	Sequential enzymatic oxidation of aminoarenes to nitroarenes via hydroxylamines. <i>Angewandte Chemie - International Edition</i> , 2005 , 44, 4083-7	16.4	64
2	Sequenzielle enzymatische Oxidation von Aminoarenen zu Nitroarenen Ber Hydroxylamine. <i>Angewandte Chemie</i> , 2005 , 117, 4152-4155	3.6	22
1	Formatting Open Science: agilely creating multiple document formats for academic manuscripts with Pandoc Scholar. <i>PeerJ Computer Science</i> , 3, e112	2.7	4