Wolfgang Wiechert

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.

80 7,149 194 44 h-index g-index citations papers 8,218 6.15 209 4.7 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
194	Bayesian calibration, process modeling and uncertainty quantification in biotechnology <i>PLoS Computational Biology</i> , 2022 , 18, e1009223	5	3
193	bletl - A Python package for integrating BioLector microcultivation devices in the Design-Build-Test-Learn cycle <i>Engineering in Life Sciences</i> , 2022 , 22, 242-259	3.4	3
192	Hot isopropanol quenching procedure for automated microtiter plate scale C-labeling experiments <i>Microbial Cell Factories</i> , 2022 , 21, 78	6.4	O
191	High-Quality Genome-Scale Reconstruction of ATCC 13032. Frontiers in Microbiology, 2021, 12, 750206	5.7	0
190	A fully automated pipeline for the dynamic at-line morphology analysis of microscale Aspergillus cultivation. <i>Fungal Biology and Biotechnology</i> , 2021 , 8, 2	7.5	3
189	Microbioreactor-assisted cultivation workflows for time-efficient phenotyping of protein producing Aspergillus niger in batch and fed-batch mode. <i>Biotechnology Progress</i> , 2021 , 37, e3144	2.8	
188	Need for speed: evaluation of dilute and shoot-mass spectrometry for accelerated metabolic phenotyping in bioprocess development. <i>Analytical and Bioanalytical Chemistry</i> , 2021 , 413, 3253-3268	4.4	1
187	Robotic integration enables autonomous operation of laboratory scale stirred tank bioreactors with model-driven process analysis. <i>Biotechnology and Bioengineering</i> , 2021 , 118, 2759-2769	4.9	2
186	Quantitative Metabolic Flux Analysis Based on Isotope Labeling 2021 , 73-136		O
185	Microaerobic growth-decoupled production of Eketoglutarate and succinate from xylose in a one-pot process using Corynebacterium glutamicum. <i>Biotechnology Journal</i> , 2021 , 16, e2100043	5.6	2
184	pyFOOMB: Python framework for object oriented modeling of bioprocesses. <i>Engineering in Life Sciences</i> , 2021 , 21, 242-257	3.4	6
183	Construction and comprehensive characterization of an EcLDCc-CatIB set-varying linkers and aggregation inducing tags. <i>Microbial Cell Factories</i> , 2021 , 20, 49	6.4	6
182	Scaling production of GFP1-10 detector protein in E. coli for secretion screening by split GFP assay. <i>Microbial Cell Factories</i> , 2021 , 20, 191	6.4	1
181	Revisiting the Growth Modulon of Under Glucose Limited Chemostat Conditions. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 584614	5.8	2
180	Simulation of differential-algebraic equation systems with optimization criteria embedded in Modelica. <i>Computers and Chemical Engineering</i> , 2020 , 140, 106920	4	2
179	Parallelized microscale fed-batch cultivation in online-monitored microtiter plates: implications of media composition and feed strategies for process design and performance. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2020 , 47, 35-47	4.2	4
178	Comprehensive analysis of metabolic sensitivity of 1,4-butanediol producing Escherichia coli toward substrate and oxygen availability. <i>Biotechnology Progress</i> , 2020 , 36, e2917	2.8	2

(2018-2020)

177	Analysis of the local well-posedness of optimization-constrained differential equations by local optimality conditions. <i>AICHE Journal</i> , 2020 , 66, e16548	3.6	2
176	Screening of a genome-reduced Corynebacterium glutamicum strain library for improved heterologous cutinase secretion. <i>Microbial Biotechnology</i> , 2020 , 13, 2020-2031	6.3	5
175	Comprehensive Analysis of Anaplerotic Deletion Mutants Under Defined d-Glucose Conditions. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 602936	5.8	2
174	A closer look at: online monitoring via scattered light enables reproducible phenotyping. <i>Fungal Biology and Biotechnology</i> , 2019 , 6, 11	7.5	5
173	mycelyso - high-throughput analysis of Streptomyces mycelium live cell imaging data. <i>BMC Bioinformatics</i> , 2019 , 20, 452	3.6	O
172	The Design of FluxML: A Universal Modeling Language for C Metabolic Flux Analysis. <i>Frontiers in Microbiology</i> , 2019 , 10, 1022	5.7	14
171	Analyzing Microbial Population Heterogeneity-Expanding the Toolbox of Microfluidic Single-Cell Cultivations. <i>Journal of Molecular Biology</i> , 2019 , 431, 4569-4588	6.5	22
170	Multiscale dynamic modeling and simulation of a biorefinery. <i>Biotechnology and Bioengineering</i> , 2019 , 116, 2561-2574	4.9	5
169	An Enzymatic 2-Step Cofactor and Co-Product Recycling Cascade towards a Chiral 1,2-Diol. Part I: Cascade Design. <i>Advanced Synthesis and Catalysis</i> , 2019 , 361, 2607-2615	5.6	7
168	An Enzymatic 2-Step Cofactor and Co-Product Recycling Cascade towards a Chiral 1,2-Diol. Part II: Catalytically Active Inclusion Bodies. <i>Advanced Synthesis and Catalysis</i> , 2019 , 361, 2616	5.6	5
167	FeedER: a feedback-regulated enzyme-based slow-release system for fed-batch cultivation in microtiter plates. <i>Bioprocess and Biosystems Engineering</i> , 2019 , 42, 1843-1852	3.7	6
166	A FRET-based biosensor for the quantification of glucose in culture supernatants of mL scale microbial cultivations. <i>Microbial Cell Factories</i> , 2019 , 18, 143	6.4	10
165	Rapid and comprehensive evaluation of microalgal fatty acids via untargeted gas chromatography and time-of-flight mass spectrometry. <i>Engineering in Life Sciences</i> , 2019 , 19, 1006-1011	3.4	0
164	Combinatorial impact of Sec signal peptides from Bacillus subtilis and bioprocess conditions on heterologous cutinase secretion by Corynebacterium glutamicum. <i>Biotechnology and Bioengineering</i> , 2019 , 116, 644-655	4.9	10
163	Less Sacrifice, More Insight: Repeated Low-Volume Sampling of Microbioreactor Cultivations Enables Accelerated Deep Phenotyping of Microbial Strain Libraries. <i>Biotechnology Journal</i> , 2019 , 14, e1800428	5.6	19
162	Microbioreactor Systems for Accelerated Bioprocess Development. <i>Biotechnology Journal</i> , 2018 , 13, e1700141	5.6	75
161	Laboratory-scale photobiotechnology-current trends and future perspectives. <i>FEMS Microbiology Letters</i> , 2018 , 365,	2.9	5
160	Quantitative measurements in single-cell analysis: towards scalability in microbial bioprocess development. <i>Current Opinion in Biotechnology</i> , 2018 , 54, 121-127	11.4	10

159	Microbial single-cell analysis in picoliter-sized batch cultivation chambers. <i>New Biotechnology</i> , 2018 , 47, 50-59	6.4	14
158	Differential transcriptomic analysis reveals hidden light response in Streptomyces lividans. <i>Biotechnology Progress</i> , 2018 , 34, 287-292	2.8	1
157	Multi-Omics and Targeted Approaches to Determine the Role of Cellular Proteases in Protein Secretion. <i>Frontiers in Microbiology</i> , 2018 , 9, 1174	5.7	13
156	Improved microscale cultivation of for clonal screening. Fungal Biology and Biotechnology, 2018, 5, 8	7.5	10
155	Corynebacterium glutamicum Chassis C1*: Building and Testing a Novel Platform Host for Synthetic Biology and Industrial Biotechnology. <i>ACS Synthetic Biology</i> , 2018 , 7, 132-144	5.7	43
154	Tailor-made catalytically active inclusion bodies for different applications in biocatalysis. <i>Catalysis Science and Technology</i> , 2018 , 8, 5816-5826	5.5	19
153	A Pareto approach to resolve the conflict between information gain and experimental costs: Multiple-criteria design of carbon labeling experiments. <i>PLoS Computational Biology</i> , 2018 , 14, e100653	ı3 ⁵	9
152	Characterization of Sigma Factor Genes in TK24 Using a Genomic Library-Based Approach for Multiple Gene Deletions. <i>Frontiers in Microbiology</i> , 2018 , 9, 3033	5.7	9
151	Germination and Growth Analysis of at the Single-Cell Level Under Varying Medium Compositions. <i>Frontiers in Microbiology</i> , 2018 , 9, 2680	5.7	3
150	The structural index of sensitivity equation systems. <i>Mathematical and Computer Modelling of Dynamical Systems</i> , 2018 , 24, 573-592	1	2
149	Real-time monitoring of fungal growth and morphogenesis at single-cell resolution. <i>Engineering in Life Sciences</i> , 2017 , 17, 86-92	3.4	9
148	Comparative evaluation of phototrophic microtiter plate cultivation against laboratory-scale photobioreactors. <i>Bioprocess and Biosystems Engineering</i> , 2017 , 40, 663-673	3.7	8
147	Dynamic flux balance analysis with nonlinear objective function. <i>Journal of Mathematical Biology</i> , 2017 , 75, 1487-1515	2	10
146	Miniaturized and automated adaptive laboratory evolution: Evolving Corynebacterium glutamicum towards an improved d-xylose utilization. <i>Bioresource Technology</i> , 2017 , 245, 1377-1385	11	57
145	Fast and reliable strain characterization of Streptomyces lividans through micro-scale cultivation. <i>Biotechnology and Bioengineering</i> , 2017 , 114, 2011-2022	4.9	18
144	Homogenizing bacterial cell factories: Analysis and engineering of phenotypic heterogeneity. <i>Metabolic Engineering</i> , 2017 , 42, 145-156	9.7	63
143	Automated growth rate determination in high-throughput microbioreactor systems. <i>BMC Research Notes</i> , 2017 , 10, 617	2.3	10
142	pH fluctuations imperil the robustness of C. glutamicum to short term oxygen limitation. <i>Journal of Biotechnology</i> , 2017 , 259, 248-260	3.7	8

(2016-2017)

To be certain about the uncertainty: Bayesian statistics for C metabolic flux analysis. <i>Biotechnology and Bioengineering</i> , 2017 , 114, 2668-2684	4.9	14	
Coarse-graining bacteria colonies for modelling critical solute distributions in picolitre bioreactors for bacterial studies on single-cell level. <i>Microbial Biotechnology</i> , 2017 , 10, 845-857	6.3	9	
Kriging with trend functions nonlinear in their parameters: Theory and application in enzyme kinetics. <i>Engineering in Life Sciences</i> , 2017 , 17, 916-922	3.4	6	
The linkage between nutrient supply, intracellular enzyme abundances and bacterial growth: New evidences from the central carbon metabolism of Corynebacterium glutamicum. <i>Journal of Biotechnology</i> , 2017 , 258, 13-24	3.7	8	
A framework for accelerated phototrophic bioprocess development: integration of parallelized microscale cultivation, laboratory automation and Kriging-assisted experimental design. <i>Biotechnology for Biofuels</i> , 2017 , 10, 26	7.8	9	
Design and validation of a parallelized micro-photobioreactor enabling phototrophic bioprocess development at elevated throughput. <i>Biotechnology and Bioengineering</i> , 2017 , 114, 122-131	4.9	13	
Metabolic profile of 1,5-diaminopentane producing Corynebacterium glutamicum under scale-down conditions: Blueprint for robustness to bioreactor inhomogeneities. <i>Biotechnology and Bioengineering</i> , 2017 , 114, 560-575	4.9	34	
Generic Protocol for Optimization of Heterologous Protein Production Using Automated Microbioreactor Technology. <i>Journal of Visualized Experiments</i> , 2017 ,	1.6	4	
Artificial fluorogenic substrates in microfluidic devices for bacterial diagnostics in biotechnology. <i>Journal of Flow Chemistry</i> , 2016 , 6, 3-7	3.3	1	
Time-resolved, single-cell analysis of induced and programmed cell death via non-invasive propidium iodide and counterstain perfusion. <i>Scientific Reports</i> , 2016 , 6, 32104	4.9	37	
Adaptation of Microscale Cultivation for Optimization of Protein Production in Pichia pastoris. <i>Chemie-Ingenieur-Technik</i> , 2016 , 88, 1406-1406	0.8		
Exploring the Sequence-Function Space of ThDP-Dependent Enzymes. <i>Chemie-Ingenieur-Technik</i> , 2016 , 88, 1246-1246	0.8		
Cutting the Gordian Knot: Identifiability of anaplerotic reactions in Corynebacterium glutamicum by means of (13) C-metabolic flux analysis. <i>Biotechnology and Bioengineering</i> , 2016 , 113, 661-74	4.9	16	
Photocaged Arabinose: A Novel Optogenetic Switch for Rapid and Gradual Control of Microbial Gene Expression. <i>ChemBioChem</i> , 2016 , 17, 296-9	3.8	23	
Plug flow versus stirred tank reactor flow characteristics in two-compartment scale-down bioreactor: Setup-specific influence on the metabolic phenotype and bioprocess performance of Corynebacterium glutamicum. <i>Engineering in Life Sciences</i> , 2016 , 16, 610-619	3.4	25	
Framework for Kriging-based iterative experimental analysis and design: Optimization of secretory protein production inCorynebacterium glutamicum. <i>Engineering in Life Sciences</i> , 2016 , 16, 538-549	3.4	20	
Discrete-continuous reaction-diffusion model with mobile point-like sources and sinks. <i>European Physical Journal E</i> , 2016 , 39, 11	1.5	6	
Automation of a Nile red staining assay enables high throughput quantification of microalgal lipid production. <i>Microbial Cell Factories</i> , 2016 , 15, 34	6.4	21	
	Coarse-graining bacteria colonies for modelling critical solute distributions in picolitre bioreactors for bacterial studies on single-cell level. <i>Microbial Biotechnology</i> , 2017, 10, 845-857 Kriging with trend functions nonlinear in their parameters: Theory and application in enzyme kinetics. <i>Engineering in Life Sciences</i> , 2017, 17, 916-922 The linkage between nutrient supply, intracellular enzyme abundances and bacterial growth: New evidences from the central carbon metabolism of Corynebacterium glutamicum. <i>Journal of Biotechnology</i> , 2017, 258, 13-24 A framework for accelerated phototrophic bioprocess development: integration of parallelized microscale cultivation, laboratory automation and Kriging-assisted experimental design. <i>Biotechnology for Biofuels</i> , 2017, 10, 26 Design and validation of a parallelized micro-photobioreactor enabling phototrophic bioprocess development at elevated throughput. <i>Biotechnology and Bioengineering</i> , 2017, 114, 122-131 Metabolic profile of 1,5-diaminopentane producing Corynebacterium glutamicum under scale-down conditions: Blueprint for robustness to bioreactor inhomogeneities. <i>Biotechnology and Bioengineering</i> , 2017, 114, 560-575 Generic Protocol for Optimization of Heterologous Protein Production Using Automated Microbioreactor Technology. <i>Journal of Visualized Experiments</i> , 2017, Artificial fluorogenic substrates in microfluidic devices for bacterial diagnostics in biotechnology. <i>Journal of Flow Chemistry</i> , 2016, 6, 3-7 Time-resolved, single-cell analysis of induced and programmed cell death via non-invasive propidium iodide and counterstain perfusion. <i>Scientific Reports</i> , 2016, 6, 32104 Adaptation of Microscale Cultivation for Optimization of Protein Production in Pichia pastoris. <i>Chemie-Ingenieur-Technik</i> , 2016, 88, 1406-1406 Exploring the Sequence-Function Space of ThDP-Dependent Enzymes. <i>Chemie-Ingenieur-Technik</i> , 2016, 88, 1246-1246 Cutting the Gordian Knot: Identifiability of anaplerotic reactions in Corynebacterium glutamicum by means of (13) C-me	Coarse-graining bacteria colonies for modelling critical solute distributions in picolitre bioreactors for bacterial studies on single-cell level. <i>Microbial Biotechnology</i> , 2017, 10, 845-857 Kriging with trend functions nonlinear in their parameters: Theory and application in enzyme kinetics. <i>Engineering in Life Sciences</i> , 2017, 17, 916-922 The linkage between nutrient supply, intracellular enzyme abundances and bacterial growth: New evidences from the central carbon metabolism of Corynebacterium glutamicum. <i>Journal of Biotechnology</i> , 2017, 258, 13-24 A framework for accelerated phototrophic bioprocess development: integration of parallelized microscale cultivation, laboratory automation and Kriging-assisted experimental design. Biotechnology for Biofuels, 2017, 10, 26 Design and validation of a parallelized micro-photobioreactor enabling phototrophic bioprocess development at elevated throughput. <i>Biotechnology and Bioengineering</i> , 2017, 114, 122-131 Metabolic profile of 1,5-diaminopentane producing Corynebacterium glutamicum under scale-down conditions: Blueprine for robustness to bioreactor inhomogeneities. <i>Biotechnology and Bioengineering</i> , 2017, 114, 560-575 Generic Protocol for Optimization of Heterologous Protein Production Using Automated Microbioreactor Technology. <i>Journal of Visualized Experiments</i> , 2017, Artificial fluorogenic substrates in microfluidic devices for bacterial diagnostics in biotechnology. <i>Journal of Flow Chemistry</i> , 2016, 6, 3-7 Time-resolved, single-cell analysis of induced and programmed cell death via non-invasive propidium iodide and counterstain perfusion. <i>Scientific Reports</i> , 2016, 6, 32104 Adaptation of Microscale Cultivation for Optimization of Protein Production in Pichia pastoris. <i>Chemie-Ingenieur-Technik</i> , 2016, 88, 1406-1406 Exploring the Sequence-Function Space of ThDP-Dependent Enzymes. <i>Chemie-Ingenieur-Technik</i> , 2016, 88, 1266-1246 Cutting the Gordian Knot: Identifiability of anaplerotic reactions in Corynebacterium glutamicum by means of (13) C-met	Coarse-graining bacteria colonies for modelling critical solute distributions in picolitre bioreactors for bacterial studies on single-cell level. Microbiol Biotechnology, 2017, 10, 845-857 Kriging with trend functions nonlinear in their parameters: Theory and application in enzyme kinetics. Engineering in Life Sciences, 2017, 17, 916-922 The linkage between nutrient supply, intracellular enzyme abundances and bacterial growth: New evidences from the central carbon metabolism of Corynebacterium glutamicum. Journal of Biotechnology, 2017, 258, 13-24 A framework for accelerated phototrophic bioprocess development: integration of parallelized microscale cultivation, laboratory automation and Kriging-assisted experimental design. Biotechnology 76 Biolyieis, 2017, 10, 26 Design and validation of a parallelized micro-photobioreactor enabling phototrophic bioprocess development at elevated throughput. Biotechnology and Bioengineering, 2017, 114, 122-131 Metabolic profile of 1,5-diaminopentane producing Corynebacterium glutamicum under scale-down conditions: Blueprint for robustness to bioreactor inhomogeneities. Biotechnology and Bioengineering, 2017, 114, 560-575 Generic Protocol for Optimization of Heterologous Protein Production Using Automated Microbioreactor Technology. Journal of Visualized Experiments, 2017. Artificial fluorogenic substrates in microfluidic devices for bacterial diagnostics in biotechnology. Journal of Flow Chemistry, 2016, 6, 3-7 Time-resolved, single-cell analysis of induced and programmed cell death via non-invasive propidium iodice and counterstain perfusion. Scientific Reports, 2016, 6, 32104 Adaptation of Microscale Cultivation for Optimization of Protein Production in Pichia pastoris. O.8 Chemie-Ingenieur-Technik, 2016, 88, 1406-1406 Exploring the Sequence-Function Space of ThDP-Dependent Enzymes. Chemie-Ingenieur-Technik, 2016, 17, 296-9 Plug flow versus stirred tank reactor flow characteristics in two-compartment scale-down bioreactor: Setup-specific influence on the meta

123	A scientific workflow framework for (13)C metabolic flux analysis. <i>Journal of Biotechnology</i> , 2016 , 232, 12-24	3.7	14
122	Image-Based Single Cell Profiling: High-Throughput Processing of Mother Machine Experiments. <i>PLoS ONE</i> , 2016 , 11, e0163453	3.7	23
121	A Toolbox of Genetically Encoded FRET-Based Biosensors for Rapid l-Lysine Analysis. <i>Sensors</i> , 2016 , 16,	3.8	20
120	Use of a Sec signal peptide library from Bacillus subtilis for the optimization of cutinase secretion in Corynebacterium glutamicum. <i>Microbial Cell Factories</i> , 2016 , 15, 208	6.4	31
119	MORE: A microfluidic magnetic oscillation reactor for accelerated parameter optimization in biocatalysis. <i>Journal of Biotechnology</i> , 2016 , 231, 174-182	3.7	10
118	Simplified cryopreservation of the microalga Chlorella vulgaris integrating a novel concept for cell viability estimation. <i>Engineering in Life Sciences</i> , 2016 , 16, 36-44	3.4	13
117	Technical bias of microcultivation environments on single-cell physiology. <i>Lab on A Chip</i> , 2015 , 15, 1822	-3/42	28
116	Chassis organism from Corynebacterium glutamicuma top-down approach to identify and delete irrelevant gene clusters. <i>Biotechnology Journal</i> , 2015 , 10, 290-301	5.6	87
115	Computational Systems Biology Theues Fach in den Lebenswissenschaften. <i>BioSpektrum</i> , 2015 , 21, 46-48	0.1	
114	Bioprocess automation on a Mini Pilot Plant enables fast quantitative microbial phenotyping. <i>Microbial Cell Factories</i> , 2015 , 14, 32	6.4	56
113	Modeling and CFD simulation of nutrient distribution in picoliter bioreactors for bacterial growth studies on single-cell level. <i>Lab on A Chip</i> , 2015 , 15, 4177-86	7.2	28
112	Vizardous: interactive analysis of microbial populations with single cell resolution. <i>Bioinformatics</i> , 2015 , 31, 3875-7	7.2	5
111	Rapid inoculation of single bacteria into parallel picoliter fermentation chambers. <i>Analytical Methods</i> , 2015 , 7, 91-98	3.2	24
110	How to measure metabolic fluxes: a taxonomic guide for (13)C fluxomics. <i>Current Opinion in Biotechnology</i> , 2015 , 34, 82-90	11.4	79
109	Absolute quantification of Corynebacterium glutamicum glycolytic and anaplerotic enzymes by QconCAT. <i>Journal of Proteomics</i> , 2015 , 113, 366-77	3.9	11
108	Visual workflows for 13C-metabolic flux analysis. <i>Bioinformatics</i> , 2015 , 31, 346-54	7.2	23
107	A Primer to 13C Metabolic Flux Analysis 2015 , 97-142		7
106	Live cell imaging of SOS and prophage dynamics in isogenic bacterial populations. <i>Molecular Microbiology</i> , 2015 , 98, 636-50	4.1	26

(2013-2015)

105	Spatiotemporal microbial single-cell analysis using a high-throughput microfluidics cultivation platform. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2015, 87, 110	04f5	68
104	(13)C Tracers for Glucose Degrading Pathway Discrimination in Gluconobacter oxydans 621H. <i>Metabolites</i> , 2015 , 5, 455-74	5.6	
103	The effect of composition on diffusion of macromolecules in a crowded environment. <i>Physical Biology</i> , 2015 , 12, 046003	3	23
102	Non-Invasive Microbial Metabolic Activity Sensing at Single Cell Level by Perfusion of Calcein Acetoxymethyl Ester. <i>PLoS ONE</i> , 2015 , 10, e0141768	3.7	11
101	Growth and Production Capabilities of Corynebacterium glutamicum: Interrogating a Genome-scale Metabolic Network Model 2015 , 39-56		7
100	Effective Production of (S)-Hydroxy ketones: An Reaction Engineering Approach. <i>Topics in Catalysis</i> , 2014 , 57, 401-411	2.3	8
99	Process inhomogeneity leads to rapid side product turnover in cultivation of Corynebacterium glutamicum. <i>Microbial Cell Factories</i> , 2014 , 13, 6	6.4	48
98	Single-cell microfluidics: opportunity for bioprocess development. <i>Current Opinion in Biotechnology</i> , 2014 , 29, 15-23	11.4	128
97	Quantitative metabolomics: a phantom?. Trends in Biotechnology, 2014, 32, 238-44	15.1	36
96	Assessment of robustness against dissolved oxygen/substrate oscillations for C. glutamicum DM1933 in two-compartment bioreactor. <i>Bioprocess and Biosystems Engineering</i> , 2014 , 37, 1151-62	3.7	35
95	Engineering of Corynebacterium glutamicum for minimized carbon loss during utilization of D-xylose containing substrates. <i>Journal of Biotechnology</i> , 2014 , 192 Pt A, 156-60	3.7	65
94	Rapid assessment of oxygen transfer impact for Corynebacterium glutamicum. <i>Bioprocess and Biosystems Engineering</i> , 2014 , 37, 2567-77	3.7	16
93	Beyond growth rate 0.6: What drives Corynebacterium glutamicum to higher growth rates in defined medium. <i>Biotechnology and Bioengineering</i> , 2014 , 111, 359-71	4.9	85
92	An evaluation of genetically encoded FRET-based biosensors for quantitative metabolite analyses in vivo. <i>Journal of Biotechnology</i> , 2014 , 191, 250-9	3.7	24
91	Light-responsive control of bacterial gene expression: precise triggering of the lac promoter activity using photocaged IPTG. <i>Integrative Biology (United Kingdom)</i> , 2014 , 6, 755-65	3.7	34
90	Developing a new production host from a blueprint: Bacillus pumilus as an industrial enzyme producer. <i>Microbial Cell Factories</i> , 2014 , 13, 46	6.4	12
89	13C-flux spectral analysis of host-pathogen metabolism reveals a mixed diet for intracellular Mycobacterium tuberculosis. <i>Chemistry and Biology</i> , 2013 , 20, 1012-21		106
88	Combined fluxomics and transcriptomics analysis of glucose catabolism via a partially cyclic pentose phosphate pathway in Gluconobacter oxydans 621H. <i>Applied and Environmental Microbiology</i> 2013 , 79, 2336-48	4.8	49

87	Omix IA Visualization Tool for Metabolic Networks with Highest Usability and Customizability in Focus. <i>Chemie-Ingenieur-Technik</i> , 2013 , 85, 849-862	0.8	21
86	Simultaneous utilization of glucose and gluconate in Penicillium chrysogenum during overflow metabolism. <i>Biotechnology and Bioengineering</i> , 2013 , 110, 3235-43	4.9	13
85	Cloud MapReduce for Monte Carlo bootstrap applied to Metabolic Flux Analysis. <i>Future Generation Computer Systems</i> , 2013 , 29, 582-590	7.5	10
84	Metabolic isotopomer labeling systems. Part III: path tracing. <i>Mathematical Biosciences</i> , 2013 , 244, 1-12	3.9	5
83	Isotopically non-stationary metabolic flux analysis: complex yet highly informative. <i>Current Opinion in Biotechnology</i> , 2013 , 24, 979-86	11.4	78
82	13CFLUX2high-performance software suite for (13)C-metabolic flux analysis. <i>Bioinformatics</i> , 2013 , 29, 143-5	7.2	150
81	Microfluidic growth chambers with optical tweezers for full spatial single-cell control and analysis of evolving microbes. <i>Journal of Microbiological Methods</i> , 2013 , 95, 470-6	2.8	34
80	Beyond growth rate 0.6: Corynebacterium glutamicum cultivated in highly diluted environments. <i>Biotechnology and Bioengineering</i> , 2013 , 110, 220-8	4.9	50
79	Microfluidic picoliter bioreactor for microbial single-cell analysis: fabrication, system setup, and operation. <i>Journal of Visualized Experiments</i> , 2013 , 50560	1.6	38
78	Polydimethylsiloxane (PDMS) Sub-Micron Traps for Single-Cell Analysis of Bacteria. <i>Micromachines</i> , 2013 , 4, 357-369	3.3	37
77	Collisional fragmentation of central carbon metabolites in LC-MS/MS increases precision of IIC metabolic flux analysis. <i>Biotechnology and Bioengineering</i> , 2012 , 109, 763-71	4.9	79
76	Surface bound adsorption in a microfluidic T-sensor: Numerical comparison and optimization of 2D and 3D models and of sensor designs. <i>Sensors and Actuators B: Chemical</i> , 2012 , 170, 75-81	8.5	3
75	Extensive exometabolome analysis reveals extended overflow metabolism in various microorganisms. <i>Microbial Cell Factories</i> , 2012 , 11, 122	6.4	176
74	A disposable picolitre bioreactor for cultivation and investigation of industrially relevant bacteria on the single cell level. <i>Lab on A Chip</i> , 2012 , 12, 2060-8	7.2	90
73	Error propagation analysis for quantitative intracellular metabolomics. <i>Metabolites</i> , 2012 , 2, 1012-30	5.6	16
72	Tailoring a stabilized variant of hydroxynitrile lyase from Arabidopsis thaliana. <i>ChemBioChem</i> , 2012 , 13, 797-802	3.8	18
71	Expressionsoptimierung in Mikroorganismen. <i>BioSpektrum</i> , 2012 , 18, 449-451	0.1	
70	Semi-automatic drawing of metabolic networks. <i>Information Visualization</i> , 2012 , 11, 171-187	2.4	2

69	Influence of Organic Solvents on Enzymatic Asymmetric Carboligations. <i>Advanced Synthesis and Catalysis</i> , 2012 , 354, 2805-2820	5.6	39
68	Accuracy of Parameter Sensitivities of DAE Systems using Finite Difference Methods. <i>IFAC Postprint Volumes IPPV / International Federation of Automatic Control</i> , 2012 , 45, 136-142		
67	Optical sensors for monitoring dynamic changes of intracellular metabolite levels in mammalian cells. <i>Nature Protocols</i> , 2011 , 6, 1818-33	18.8	97
66	Determination of flux directions by thermodynamic network analysis: Computing informative metabolite pools. <i>Mathematics and Computers in Simulation</i> , 2011 , 82, 460-470	3.3	3
65	Mechanistic pathway modeling for industrial biotechnology: challenging but worthwhile. <i>Current Opinion in Biotechnology</i> , 2011 , 22, 604-10	11.4	40
64	The benefits of being transient: isotope-based metabolic flux analysis at the short time scale. <i>Applied Microbiology and Biotechnology</i> , 2011 , 91, 1247-65	5.7	64
63	An Efficient Route to Both Enantiomers of allo-Threonine by Simultaneous Amino Acid Racemase-Catalyzed Isomerization of Threonine and Crystallization. <i>Advanced Synthesis and Catalysis</i> , 2011 , 353, 2431-2438	5.6	9
62	Hydroxynitrile Lyase from Arabidopsis thaliana: Identification of Reaction Parameters for Enantiopure Cyanohydrin Synthesis by Pure and Immobilized Catalyst. <i>Advanced Synthesis and Catalysis</i> , 2011 , 353, 2399-2408	5.6	31
61	An Online Provenance Service for Distributed Metabolic Flux Analysis Workflows 2011,		3
60	Visualizing multi-omics data in metabolic networks with the software Omix: a case study. <i>BioSystems</i> , 2011 , 105, 154-61	1.9	78
59	Stationary versus non-stationary (13)C-MFA: a comparison using a consistent dataset. <i>Journal of Biotechnology</i> , 2011 , 154, 179-90	3.7	55
58	Comparative 13C metabolic flux analysis of pyruvate dehydrogenase complex-deficient, L-valine-producing Corynebacterium glutamicum. <i>Applied and Environmental Microbiology</i> , 2011 , 77, 664	1 4 :82	63
57	A Hybrid Parallelization Approach for Cloud-enabled Metabolic Flux Analysis Simulation Workflows. <i>PARS Parallel-Algorithmen -Rechnerstrukturen Und -Systemsoftware</i> , 2011 , 28, 210-213		
56	Workflows for Metabolic Flux Analysis: Data Integration and Human Interaction. <i>Lecture Notes in Computer Science</i> , 2010 , 261-275	0.9	6
55	Metabolic Flux Analysis in the Cloud 2010 ,		7
54	Modeling languages for biochemical network simulation: reaction vs equation based approaches. <i>Advances in Biochemical Engineering/Biotechnology</i> , 2010 , 121, 109-38	1.7	1
53	Surface bound adsorption in a microfluidic T-sensor: Numerical comparison and optimization of 2D and 3D models. <i>Procedia Engineering</i> , 2010 , 5, 1272-1275		
52	Customizable Visualization on Demand for Hierarchically Organized Information in Biochemical Networks. <i>Lecture Notes in Computer Science</i> , 2010 , 163-174	0.9	2

51	Translating biochemical network models between different kinetic formats. <i>Metabolic Engineering</i> , 2009 , 11, 87-100	9.7	24
50	Customizable Visualization of Multi-omics Data in the Context of Biochemical Networks 2009,		2
49	Atomic Force Microscope Cantilevers Used as Sensors for Monitoring Microdrop Evaporation. <i>Nanoscience and Technology</i> , 2009 , 17-38	0.6	
48	13C labeling experiments at metabolic nonstationary conditions: an exploratory study. <i>BMC Bioinformatics</i> , 2008 , 9, 152	3.6	45
47	Visualizing regulatory interdependencies and parameter sensitivities in biochemical network models. <i>Mathematics and Computers in Simulation</i> , 2008 , 79, 991-998	3.3	4
46	Automatic Sensitivity Analysis of DAE-systems Generated from Equation-Based Modeling Languages. <i>Lecture Notes in Computational Science and Engineering</i> , 2008 , 235-246	0.3	5
45	The thermodynamic meaning of metabolic exchange fluxes. <i>Biophysical Journal</i> , 2007 , 93, 2255-64	2.9	36
44	Visualizing regulatory interactions in metabolic networks. <i>BMC Biology</i> , 2007 , 5, 46	7.3	13
43	The topology of metabolic isotope labeling networks. <i>BMC Bioinformatics</i> , 2007 , 8, 315	3.6	24
42	Fluxomics: mass spectrometry versus quantitative imaging. <i>Current Opinion in Plant Biology</i> , 2007 , 10, 323-30	9.9	68
41	Event driven algorithms applied to a high energy ball mill simulation. <i>Granular Matter</i> , 2007 , 9, 251-266	2.6	6
40	Evaporation of Solvent Microdrops on Polymer Substrates: From Well Controlled Experiments To Mathematical Models and Back. <i>Nanoscale and Microscale Thermophysical Engineering</i> , 2007 , 11, 31-41	3.7	7
39	Nondestructive and noncontact method for determining the spring constant of rectangular cantilevers. <i>Review of Scientific Instruments</i> , 2007 , 78, 043705	1.7	23
38	Stress and failure at mechanical contacts of microspheres under uniaxial compression. <i>Journal of Applied Physics</i> , 2007 , 101, 084908	2.5	3
37	Metabolic flux analysis at ultra short time scale: isotopically non-stationary 13C labeling experiments. <i>Journal of Biotechnology</i> , 2007 , 129, 249-67	3.7	178
36	Experimental design principles for isotopically instationary 13C labeling experiments. <i>Biotechnology and Bioengineering</i> , 2006 , 94, 234-51	4.9	81
35	Interpretation of metabolic flux maps by limitation potentials and constrained limitation sensitivities. <i>Biotechnology and Bioengineering</i> , 2006 , 94, 263-72	4.9	2
34	Sessile-drop-induced bending of atomic force microscope cantilevers: a model system for monitoring microdrop evaporation. <i>Journal of Micromechanics and Microengineering</i> , 2006 , 16, 2273-228	3 6	30

(2002-2006)

33	Modelling and simulation of micro-well formation. <i>Mathematical and Computer Modelling of Dynamical Systems</i> , 2006 , 12, 263-276	1	7
32	Pressure distribution in a mechanical microcontact. Applied Physics Letters, 2006, 88, 234101	3.4	8
31	A general framework for large-scale model selection. Optimization Methods and Software, 2006, 21, 901	-9. 37	2
30	Emerging Corynebacterium glutamicum systems biology. <i>Journal of Biotechnology</i> , 2006 , 124, 74-92	3.7	93
29	Computational tools for isotopically instationary 13C labeling experiments under metabolic steady state conditions. <i>Metabolic Engineering</i> , 2006 , 8, 554-77	9.7	89
28	From Enzyme Kinetics to Metabolic Network Modeling LV isualization Tool for Enhanced Kinetic Analysis of Biochemical Network Models. <i>Engineering in Life Sciences</i> , 2006 , 6, 155-162	3.4	6
27	From stationary to instationary metabolic flux analysis. <i>Advances in Biochemical Engineering/Biotechnology</i> , 2005 , 92, 145-72	1.7	42
26	The time-dependent reorderable matrix method for visualizing evolving tabular data 2005 , 5669, 199		1
25	Investigating the dynamic behavior of biochemical networks using model families. <i>Bioinformatics</i> , 2005 , 21, 1617-25	7.2	31
24	A Multi-Scale Modeling Concept and Computational Tools for the Integrative Analysis of Stationary Metabolic Data. <i>Journal of Integrative Bioinformatics</i> , 2004 , 1, 38-51	3.8	1
23	Parallel Solution of Cascaded ODE Systems Applied to 13C-Labeling Experiments. <i>Lecture Notes in Computer Science</i> , 2004 , 594-597	0.9	1
22	Serial 13C-based flux analysis of an L-phenylalanine-producing E. coli strain using the sensor reactor. <i>Biotechnology Progress</i> , 2004 , 20, 706-14	2.8	32
21	New tools for mass isotopomer data evaluation in (13)C flux analysis: mass isotope correction, data consistency checking, and precursor relationships. <i>Biotechnology and Bioengineering</i> , 2004 , 85, 259-68	4.9	140
20	Serial flux mapping of Corynebacterium glutamicum during fed-batch L-lysine production using the sensor reactor approach. <i>Biotechnology and Bioengineering</i> , 2004 , 85, 497-505	4.9	27
19	The role of modeling in computational science education. <i>Future Generation Computer Systems</i> , 2003 , 19, 1363-1374	7.5	4
18	Metabolic isotopomer labeling systems. Part II: structural flux identifiability analysis. <i>Mathematical Biosciences</i> , 2003 , 183, 175-214	3.9	39
17	An introduction to 13C metabolic flux analysis. <i>Genetic Engineering</i> , 2002 , 24, 215-38		17
16	Modeling and simulation: tools for metabolic engineering. <i>Journal of Biotechnology</i> , 2002 , 94, 37-63	3.7	161

MMT--a pathway modeling tool for data from rapid sampling experiments. In Silico Biology, 2002, 2, 467-84 15 4 13C metabolic flux analysis. *Metabolic Engineering*, **2001**, 3, 195-206 14 9.7 641 A universal framework for 13C metabolic flux analysis. Metabolic Engineering, 2001, 3, 265-83 296 13 9.7 Metabolic isotopomer labeling systems. Part I: global dynamic behavior. Mathematical Biosciences, 12 3.9 54 **2001**, 169, 173-205 In vivo quantification of parallel and bidirectional fluxes in the anaplerosis of Corynebacterium 11 5.4 155 glutamicum. Journal of Biological Chemistry, 2000, 275, 35932-41 Determination of full 13C isotopomer distributions for metabolic flux analysis using heteronuclear 58 10 3.7 spin echo difference NMR spectroscopy. Journal of Biotechnology, 2000, 77, 25-35 Metabolic state of Zymomonas mobilis in glucose-, fructose-, and xylose-fed continuous cultures as 81 9 3 analysed by 13C- and 31P-NMR spectroscopy. Archives of Microbiology, 1999, 171, 371-85 Bidirectional reaction steps in metabolic networks: III. Explicit solution and analysis of isotopomer 4.9 245 labeling systems. Biotechnology and Bioengineering, 1999, 66, 69-85 Bidirectional reaction steps in metabolic networks: IV. Optimal design of isotopomer labeling 163 4.9 experiments. Biotechnology and Bioengineering, 1999, 66, 86-103 13C tracer experiments and metabolite balancing for metabolic flux analysis: comparing two 62 4.9 approaches. Biotechnology and Bioengineering, 1998, 58, 254-7 Bidirectional reaction steps in metabolic networks: I. Modeling and simulation of carbon isotope 5 4.9 206 labeling experiments. Biotechnology and Bioengineering, 1997, 55, 101-17 Bidirectional reaction steps in metabolic networks: II. Flux estimation and statistical analysis. 198 4.9 Biotechnology and Bioengineering, 1997, 55, 118-35 Determination of the fluxes in the central metabolism of Corynebacterium glutamicum by nuclear magnetic resonance spectroscopy combined with metabolite balancing. Biotechnology and 3 4.9 375 Bioengineering, **1996**, 49, 111-29 Stochastic simulation of biotechnical processes. Mathematics and Computers in Simulation, 1996, 42, 171:178 Bayesian calibration, process modeling and uncertainty quantification in biotechnology 1 1