

Ralf Takors

List of Publications by Year in descending order

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95
papers

3,190
citations

172207

29
h-index

168136

53
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98
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98
docs citations

98
times ranked

3227
citing authors

#	ARTICLE	IF	CITATIONS
1	Simultaneous determination of multiple intracellular metabolites in glycolysis, pentose phosphate pathway and tricarboxylic acid cycle by liquid chromatography–mass spectrometry. <i>Journal of Chromatography A</i> , 2007, 1147, 153-164.	1.8	430
2	Quantification of Intracellular Metabolites in <i>Escherichia coli</i> K12 Using Liquid Chromatographic-Electrospray Ionization Tandem Mass Spectrometric Techniques. <i>Analytical Biochemistry</i> , 2001, 295, 129-137.	1.1	210
3	Metabolomics: quantification of intracellular metabolite dynamics. <i>New Biotechnology</i> , 2002, 19, 5-15.	2.7	194
4	Genome reduction boosts heterologous gene expression in <i>Pseudomonas putida</i> . <i>Microbial Cell Factories</i> , 2015, 14, 23.	1.9	142
5	Using gas mixtures of CO, CO ₂ and H ₂ as microbial substrates: the do's and don'ts of successful technology transfer from laboratory to production scale. <i>Microbial Biotechnology</i> , 2018, 11, 606-625.	2.0	126
6	High Substrate Uptake Rates Empower <i>Vibrio natriegens</i> as Production Host for Industrial Biotechnology. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	1.4	112
7	Platform Engineering of <i>Corynebacterium glutamicum</i> with Reduced Pyruvate Dehydrogenase Complex Activity for Improved Production of L-Lysine, L-Valine, and 2-Ketoisovalerate. <i>Applied and Environmental Microbiology</i> , 2013, 79, 5566-5575.	1.4	98
8	Grand Research Challenges for Sustainable Industrial Biotechnology. <i>Trends in Biotechnology</i> , 2019, 37, 1042-1050.	4.9	94
9	Process strategies to enhance pyruvate production with recombinant <i>Escherichia coli</i> : From repetitive fed-batch to in situ product recovery with fully integrated electrodialysis. <i>Biotechnology and Bioengineering</i> , 2004, 85, 638-646.	1.7	83
10	Alkaline conditions in hydrophilic interaction liquid chromatography for intracellular metabolite quantification using tandem mass spectrometry. <i>Analytical Biochemistry</i> , 2015, 475, 4-13.	1.1	72
11	Engineering <i>E. coli</i> for large-scale production – Strategies considering ATP expenses and transcriptional responses. <i>Metabolic Engineering</i> , 2016, 38, 73-85.	3.6	72
12	CO ₂ /HCO ₃ ⁻ perturbations of simulated large scale gradients in a scale-down device cause fast transcriptional responses in <i>Corynebacterium glutamicum</i> . <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 8563-8572.	1.7	63
13	Bioprocess scale-up/down as integrative enabling technology: from fluid mechanics to systems biology and beyond. <i>Microbial Biotechnology</i> , 2017, 10, 1267-1274.	2.0	55
14	Simplified absolute metabolite quantification by gas chromatography–isotope dilution mass spectrometry on the basis of commercially available source material. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2011, 879, 3859-3870.	1.2	53
15	Monitoring and Modeling of the Reaction Dynamics in the Valine/Leucine Synthesis Pathway in <i>Corynebacterium glutamicum</i> . <i>Biotechnology Progress</i> , 2006, 22, 1071-1083.	1.3	45
16	CO ₂ – Intrinsic Product, Essential Substrate, and Regulatory Trigger of Microbial and Mammalian Production Processes. <i>Frontiers in Bioengineering and Biotechnology</i> , 2015, 3, 108.	2.0	45
17	<i>Pseudomonas putida</i> KT2440 is naturally endowed to withstand industrial-scale stress conditions. <i>Microbial Biotechnology</i> , 2020, 13, 1145-1161.	2.0	42
18	Transcriptional response of <i>Escherichia coli</i> to ammonia and glucose fluctuations. <i>Microbial Biotechnology</i> , 2017, 10, 858-872.	2.0	41

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19	Metabolic engineering to guide evolution – Creating a novel mode for L-valine production with <i>Corynebacterium glutamicum</i> . <i>Metabolic Engineering</i> , 2018, 47, 31-41.	3.6	41
20	Impact of different CO ₂ /HCO ₃ ⁻ levels on metabolism and regulation in <i>Corynebacterium glutamicum</i> . <i>Journal of Biotechnology</i> , 2013, 168, 331-340.	1.9	40
21	<i>Escherichia coli</i> HGT: Engineered for high glucose throughput even under slowly growing or resting conditions. <i>Metabolic Engineering</i> , 2017, 40, 93-103.	3.6	39
22	Dynamics of benzoate metabolism in <i>Pseudomonas putida</i> KT2440. <i>Metabolic Engineering Communications</i> , 2016, 3, 97-110.	1.9	37
23	Lagrangian Trajectories to Predict the Formation of Population Heterogeneity in Large-Scale Bioreactors. <i>Bioengineering</i> , 2017, 4, 27.	1.6	37
24	Compartment-specific metabolomics for CHO reveals that ATP pools in mitochondria are much lower than in cytosol. <i>Biotechnology Journal</i> , 2015, 10, 1639-1650.	1.8	36
25	Valorization of pyrolysis water: a biorefinery side stream, for 1,2-propanediol production with engineered <i>Corynebacterium glutamicum</i> . <i>Biotechnology for Biofuels</i> , 2017, 10, 277.	6.2	35
26	Improving the carbon balance of fermentations by total carbon analyses. <i>Biochemical Engineering Journal</i> , 2014, 90, 162-169.	1.8	34
27	Modular systems metabolic engineering enables balancing of relevant pathways for l-histidine production with <i>Corynebacterium glutamicum</i> . <i>Biotechnology for Biofuels</i> , 2019, 12, 65.	6.2	34
28	Harnessing novel chromosomal integration loci to utilize an organosolv-derived hemicellulose fraction for isobutanol production with engineered <i>Corynebacterium glutamicum</i> . <i>Microbial Biotechnology</i> , 2018, 11, 257-263.	2.0	33
29	HILIC-Enabled ¹³ C Metabolomics Strategies: Comparing Quantitative Precision and Spectral Accuracy of QTOF High- and QQQ Low-Resolution Mass Spectrometry. <i>Metabolites</i> , 2019, 9, 63.	1.3	32
30	Experimentally Validated Model Enables Debottlenecking of <i>in Vitro</i> Protein Synthesis and Identifies a Control Shift under <i>in Vivo</i> Conditions. <i>ACS Synthetic Biology</i> , 2017, 6, 1913-1921.	1.9	30
31	Continuous Adaptive Evolution of a Fast-Growing <i>Corynebacterium glutamicum</i> Strain Independent of Protocatechuate. <i>Frontiers in Microbiology</i> , 2019, 10, 1648.	1.5	29
32	The impact of CO gradients on <i>C. ljungdahlii</i> in a 125 m ³ bubble column: Mass transfer, circulation time and lifeline analysis. <i>Chemical Engineering Science</i> , 2019, 207, 410-423.	1.9	29
33	Exploiting <i>Hydrogenophaga pseudoflava</i> for aerobic syngas-based production of chemicals. <i>Metabolic Engineering</i> , 2019, 55, 220-230.	3.6	28
34	Predictability of <i>k_La</i> in stirred tank reactors under multiple operating conditions using an Euler-Lagrange approach. <i>Engineering in Life Sciences</i> , 2016, 16, 633-642.	2.0	27
35	A guide to gene regulatory network inference for obtaining predictive solutions: Underlying assumptions and fundamental biological and data constraints. <i>BioSystems</i> , 2018, 174, 37-48.	0.9	27
36	Electron availability in CO ₂ , CO and H ₂ mixtures constrains flux distribution, energy management and product formation in <i>Clostridium ljungdahlii</i> . <i>Microbial Biotechnology</i> , 2020, 13, 1831-1846.	2.0	27

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37	Zero-growth bioprocesses: A challenge for microbial production strains and bioprocess engineering. <i>Engineering in Life Sciences</i> , 2017, 17, 27-35.	2.0	26
38	Phosphate limited fed-batch processes: Impact on carbon usage and energy metabolism in <i>Escherichia coli</i> . <i>Journal of Biotechnology</i> , 2014, 190, 96-104.	1.9	25
39	Changes in intracellular ATP-content of CHO cells as response to hyperosmolality. <i>Biotechnology Progress</i> , 2015, 31, 1212-1216.	1.3	24
40	Physiological Response of <i>Corynebacterium glutamicum</i> to Increasingly Nutrient-Rich Growth Conditions. <i>Frontiers in Microbiology</i> , 2018, 9, 2058.	1.5	24
41	Simulated oxygen and glucose gradients as a prerequisite for predicting industrial scale performance a priori. <i>Biotechnology and Bioengineering</i> , 2020, 117, 2760-2770.	1.7	24
42	The identification of enzyme targets for the optimization of a valine producing <i>Corynebacterium glutamicum</i> strain using a kinetic model. <i>Biotechnology Progress</i> , 2009, 25, 754-762.	1.3	23
43	Applying systems biology tools to study <i>n</i> -butanol degradation in <i>Pseudomonas putida</i> KT2440. <i>Engineering in Life Sciences</i> , 2015, 15, 760-771.	2.0	23
44	Hyperosmotic stimulus study discloses benefits in ATP supply and reveals miRNA/mRNA targets to improve recombinant protein production of CHO cells. <i>Biotechnology Journal</i> , 2016, 11, 1037-1047.	1.8	23
45	Repetitive Short-Term Stimuli Imposed in Poor Mixing Zones Induce Long-Term Adaptation of <i>E. coli</i> Cultures in Large-Scale Bioreactors: Experimental Evidence and Mathematical Model. <i>Frontiers in Microbiology</i> , 2017, 8, 1195.	1.5	23
46	Switching between nitrogen and glucose limitation: Unraveling transcriptional dynamics in <i>Escherichia coli</i> . <i>Journal of Biotechnology</i> , 2017, 258, 2-12.	1.9	19
47	Deciphering the Adaptation of <i>Corynebacterium glutamicum</i> in Transition from Aerobiosis via Microaerobiosis to Anaerobiosis. <i>Genes</i> , 2018, 9, 297.	1.0	19
48	Protein production in <i>Escherichia coli</i> is guided by the trade-off between intracellular substrate availability and energy cost. <i>Microbial Cell Factories</i> , 2019, 18, 8.	1.9	19
49	From nutritional wealth to autophagy: In vivo metabolic dynamics in the cytosol, mitochondrion and shuttles of IgG producing CHO cells. <i>Metabolic Engineering</i> , 2019, 54, 145-159.	3.6	18
50	Engineering <i>Pseudomonas putida</i> KT2440 for the production of isobutanol. <i>Engineering in Life Sciences</i> , 2020, 20, 148-159.	2.0	18
51	Production of <i>n</i> -Octanol from <i>n</i> -Octane by <i>Pseudomonas putida</i> KT2440. <i>Chemie-Ingenieur-Technik</i> , 2013, 85, 841-848.	0.4	16
52	Subpopulation-proteomics reveal growth rate, but not cell cycling, as a major impact on protein composition in <i>Pseudomonas putida</i> KT2440. <i>AMB Express</i> , 2014, 4, 71.	1.4	16
53	Environmental stress speeds up DNA replication in <i>Pseudomonas putida</i> in chemostat cultivations. <i>Biotechnology Journal</i> , 2016, 11, 155-163.	1.8	16
54	Engineering of a robust <i>Escherichia coli</i> chassis and exploitation for large-scale production processes. <i>Metabolic Engineering</i> , 2021, 67, 75-87.	3.6	15

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55	In Silico Prediction of Large-Scale Microbial Production Performance: Constraints for Getting Proper Data-Driven Models. Computational and Structural Biotechnology Journal, 2018, 16, 246-256.	1.9	13
56	Perfusion cultures require optimum respiratory ATP supply to maximize cell-specific and volumetric productivities. Biotechnology and Bioengineering, 2019, 116, 951-960.	1.7	12
57	Identifying the Growth Modulon of <i>Corynebacterium glutamicum</i> . Frontiers in Microbiology, 2019, 10, 974.	1.5	12
58	Predicting By-Product Gradients of Baker's Yeast Production at Industrial Scale: A Practical Simulation Approach. Processes, 2020, 8, 1554.	1.3	12
59	<i>Pseudomonas putida</i> KT2440 endures temporary oxygen limitations. Biotechnology and Bioengineering, 2021, 118, 4735-4750.	1.7	12
60	Scaling down biopharmaceutical production processes via a single multi-compartment bioreactor (SMCB). Engineering in Life Sciences, 2023, 23, .	2.0	12
61	Dynamic modeling reveals a three-step response of <i>Saccharomyces cerevisiae</i> to high CO ₂ levels accompanied by increasing ATP demands. FEMS Yeast Research, 2017, 17, .	1.1	11
62	CO ₂ /HCO ₃ ⁻ Accelerates Iron Reduction through Phenolic Compounds. MBio, 2020, 11, .	1.8	11
63	Identifying and Engineering Bottlenecks of Autotrophic Isobutanol Formation in Recombinant <i>C. ljungdahlii</i> by Systemic Analysis. Frontiers in Bioengineering and Biotechnology, 2021, 9, 647853.	2.0	10
64	Microaerobic production of isobutanol with engineered <i>Pseudomonas putida</i> . Engineering in Life Sciences, 2021, 21, 475-488.	2.0	9
65	Towards valorization of pectin-rich agro-industrial residues: Engineering of <i>Saccharomyces cerevisiae</i> for co-fermentation of d-galacturonic acid and glycerol. Metabolic Engineering, 2022, 69, 1-14.	3.6	9
66	The Less the Better: How Suppressed Base Addition Boosts Production of Monoclonal Antibodies With Chinese Hamster Ovary Cells. Frontiers in Bioengineering and Biotechnology, 2019, 7, 76.	2.0	8
67	Synergistically applying 1D modeling and CFD for designing industrial scale bubble column syngas bioreactors. Engineering in Life Sciences, 2020, 20, 239-251.	2.0	8
68	Tracking dipeptides at work-up take and intracellular fate in CHO culture. AMB Express, 2016, 6, 48.	1.4	7
69	Modeling Cell-Free Protein Synthesis Systems' Approaches and Applications. Frontiers in Bioengineering and Biotechnology, 2020, 8, 584178.	2.0	7
70	Revisiting the Growth Modulon of <i>Corynebacterium glutamicum</i> Under Glucose Limited Chemostat Conditions. Frontiers in Bioengineering and Biotechnology, 2020, 8, 584614.	2.0	7
71	Prediction of novel non-coding RNAs relevant for the growth of <i>Pseudomonas putida</i> in a bioreactor. Microbiology (United Kingdom), 2020, 166, 149-156.	0.7	7
72	Data-driven in silico prediction of regulation heterogeneity and ATP demands of <i>Escherichia coli</i> in large-scale bioreactors. Biotechnology and Bioengineering, 2021, 118, 265-278.	1.7	6

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73	CRISPRi enables fast growth followed by stable aerobic pyruvate formation in <i>Escherichia coli</i> without auxotrophy. <i>Engineering in Life Sciences</i> , 2022, 22, 70-84.	2.0	6
74	Quantitative Profiling of Endogenous Metabolites Using Hydrophilic Interaction Liquid Chromatography-Tandem Mass Spectrometry (HILIC-MS/MS). <i>Methods in Molecular Biology</i> , 2019, 1859, 185-207.	0.4	5
75	Comparison of tyrosine containing dipeptides reveals maximum ATP availability for prolyl-tyrosine in CHO cells. <i>Engineering in Life Sciences</i> , 2020, 20, 384-394.	2.0	5
76	Streamlining the Analysis of Dynamic ¹³ C-Labeling Patterns for the Metabolic Engineering of <i>Corynebacterium glutamicum</i> as l-Histidine Production Host. <i>Metabolites</i> , 2020, 10, 458.	1.3	5
77	Balancing glucose and oxygen uptake rates to enable high amorphadiene production in <i>Escherichia coli</i> via the methylerythritol phosphate pathway. <i>Biotechnology and Bioengineering</i> , 2021, 118, 1317-1329.	1.7	5
78	Monitoring Intracellular Metabolite Dynamics in <i>Saccharomyces cerevisiae</i> during Industrially Relevant Famine Stimuli. <i>Metabolites</i> , 2022, 12, 263.	1.3	5
79	Editorial: How can we ensure the successful transfer from lab to large scale production?. <i>Engineering in Life Sciences</i> , 2016, 16, 587-587.	2.0	4
80	S-adenosylmethionine and methylthioadenosine boost cellular productivities of antibody forming Chinese hamster ovary cells. <i>Biotechnology and Bioengineering</i> , 2020, 117, 3239-3247.	1.7	4
81	Reduced and Minimal Cell Factories in Bioprocesses: Towards a Streamlined Chassis. , 2020, , 1-44.		4
82	FAIR research data management as community approach in bioengineering. <i>Engineering in Life Sciences</i> , 2023, 23, .	2.0	4
83	Monitoring intracellular protein degradation in antibody-producing Chinese hamster ovary cells. <i>Engineering in Life Sciences</i> , 2015, 15, 499-508.	2.0	3
84	Biochemical engineering provides mindset, tools and solutions for the driving questions of a sustainable future. <i>Engineering in Life Sciences</i> , 2020, 20, 5-6.	2.0	3
85	Compartment-specific metabolome labeling enables the identification of subcellular fluxes that may serve as promising metabolic engineering targets in CHO cells. <i>Bioprocess and Biosystems Engineering</i> , 2021, 44, 2567-2578.	1.7	3
86	Wachstumskinetik. , 2018, , 45-70.		3
87	Editorial overview: Microbial systems biology: systems biology prepares the ground for successful synthetic biology. <i>Current Opinion in Microbiology</i> , 2016, 33, viii-x.	2.3	2
88	Methylthioadenosine (MTA) boosts cell-specific productivities of Chinese hamster ovary cultures: dosage effects on proliferation, cell cycle and gene expression. <i>FEBS Open Bio</i> , 2020, 10, 2791-2804.	1.0	2
89	Comprehensive Analysis of <i>C. glutamicum</i> Anaplerotic Deletion Mutants Under Defined d-Glucose Conditions. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020, 8, 602936.	2.0	2
90	Transcriptional profiling of the stringent response mutant strain <i>E. coli</i> SR reveals enhanced robustness to large scale conditions. <i>Microbial Biotechnology</i> , 2021, 14, 993-1010.	2.0	2

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91	Compartment-specific ¹³ C metabolic flux analysis reveals boosted NADPH availability coinciding with increased cell-specific productivity for IgG1 producing CHO cells after MTA treatment. <i>Engineering in Life Sciences</i> , 2021, 21, 832-847.	2.0	2
92	Synthetic mutualism in engineered <i>E. coli</i> mutant strains as functional basis for microbial production consortia. <i>Engineering in Life Sciences</i> , 0, , .	2.0	2
93	Euler-Lagrangian Simulations: A Proper Tool for Predicting Cellular Performance in Industrial Scale Bioreactors. <i>Advances in Biochemical Engineering/Biotechnology</i> , 2020, 177, 229-254.	0.6	1
94	A transhydrogenase-like mechanism in CHO cells comprising concerted cytosolic reaction and mitochondrial shuttling activities. <i>Biochemical Engineering Journal</i> , 2021, 170, 107986.	1.8	0
95	Systembiologie in der Bioverfahrenstechnik. , 2018, , 545-569.		0