

Eleftherios T Papoutsakis

List of Publications by Year in descending order

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152
papers

11,126
citations

19655

61
h-index

32838

100
g-index

158
all docs

158
docs citations

158
times ranked

6735
citing authors

#	ARTICLE	IF	CITATIONS
1	Recent advances toward the bioconversion of methane and methanol in synthetic methylotrophs. <i>Metabolic Engineering</i> , 2022, 71, 99-116.	7.0	24
2	Extracellular vesicles facilitate large-scale dynamic exchange of proteins and RNA among cultured Chinese hamster ovary and human cells. <i>Biotechnology and Bioengineering</i> , 2022, 119, 1222-1238.	3.3	18
3	Cover Image, Volume 119, Number 4, April 2022. <i>Biotechnology and Bioengineering</i> , 2022, 119, .	3.3	0
4	¹³ C-metabolic flux analysis of <i>Clostridium ljungdahlii</i> illuminates its core metabolism under mixotrophic culture conditions. <i>Metabolic Engineering</i> , 2022, 72, 161-170.	7.0	6
5	Cover Image, Volume 119, Number 5, May 2022. <i>Biotechnology and Bioengineering</i> , 2022, 119, .	3.3	0
6	miR-486-5p and miR-22-3p Enable Megakaryocytic Differentiation of Hematopoietic Stem and Progenitor Cells without Thrombopoietin. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5355.	4.1	10
7	Regulatory interventions improve the biosynthesis of limiting amino acids from methanol carbon to improve synthetic methylotrophy in <i>Escherichia coli</i> . <i>Biotechnology and Bioengineering</i> , 2021, 118, 43-57.	3.3	8
8	Adaptive laboratory evolution of methylotrophic <i>Escherichia coli</i> enables synthesis of all amino acids from methanol-derived carbon. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 869-876.	3.6	14
9	Improving the Methanol Tolerance of an <i>Escherichia coli</i> Methylotroph via Adaptive Laboratory Evolution Enhances Synthetic Methanol Utilization. <i>Frontiers in Microbiology</i> , 2021, 12, 638426.	3.5	18
10	Modeling Growth Kinetics, Interspecies Cell Fusion, and Metabolism of a <i>Clostridium acetobutylicum</i> / <i>Clostridium ljungdahlii</i> Syntrophic Coculture. <i>MSystems</i> , 2021, 6, .	3.8	6
11	Anaerobic fluorescent reporters for cell identification, microbial cell biology and high-throughput screening of microbiota and genomic libraries. <i>Current Opinion in Biotechnology</i> , 2021, 71, 151-163.	6.6	14
12	Improving synthetic methylotrophy via dynamic formaldehyde regulation of pentose phosphate pathway genes and redox perturbation. <i>Metabolic Engineering</i> , 2020, 57, 247-255.	7.0	24
13	Development of Strong Anaerobic Fluorescent Reporters for <i>Clostridium acetobutylicum</i> and <i>Clostridium ljungdahlii</i> Using HaloTag and SNAP-tag Proteins. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	21
14	Interspecies Microbial Fusion and Large-Scale Exchange of Cytoplasmic Proteins and RNA in a Syntrophic <i>Clostridium</i> Coculture. <i>MBio</i> , 2020, 11, .	4.1	36
15	Engineering <i>Escherichia coli</i> for methanol-dependent growth on glucose for metabolite production. <i>Metabolic Engineering</i> , 2020, 60, 45-55.	7.0	32
16	Triggering the stringent response enhances synthetic methanol utilization in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2020, 61, 1-10.	7.0	13
17	Human megakaryocytic microparticles induce de novo platelet biogenesis in a wild-type murine model. <i>Blood Advances</i> , 2020, 4, 804-814.	5.2	21
18	A Strongly Fluorescing Anaerobic Reporter and Protein-Tagging System for <i>Clostridium</i> Organisms Based on the Fluorescence-Activating and Absorption-Shifting Tag Protein (FAST). <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	52

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19	Extracellular vesicles: exosomes, microparticles, their parts, and their targets to enable their biomanufacturing and clinical applications. <i>Current Opinion in Biotechnology</i> , 2019, 60, 89-98.	6.6	123
20	Direct cell-to-cell exchange of matter in a synthetic <i>Clostridium</i> syntrophy enables CO ₂ fixation, superior metabolite yields, and an expanded metabolic space. <i>Metabolic Engineering</i> , 2019, 52, 9-19.	7.0	70
21	Deletion of four genes in <i>Escherichia coli</i> enables preferential consumption of xylose and secretion of glucose. <i>Metabolic Engineering</i> , 2019, 52, 168-177.	7.0	24
22	Functional Expression of the <i>Clostridium ljungdahlii</i> Acetyl-Coenzyme A Synthase in <i>Clostridium acetobutylicum</i> as Demonstrated by a Novel <i>In Vivo</i> CO Exchange Activity En Route to Heterologous Installation of a Functional Wood-Ljungdahl Pathway. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	24
23	Engineering the bioconversion of methane and methanol to fuels and chemicals in native and synthetic methylotrophs. <i>Current Opinion in Biotechnology</i> , 2018, 50, 81-93.	6.6	94
24	Expression of heterologous non-oxidative pentose phosphate pathway from <i>Bacillus methanolicus</i> and phosphoglucose isomerase deletion improves methanol assimilation and metabolite production by a synthetic <i>Escherichia coli</i> methylotroph. <i>Metabolic Engineering</i> , 2018, 45, 75-85.	7.0	74
25	Engineering human megakaryocytic microparticles for targeted delivery of nucleic acids to hematopoietic stem and progenitor cells. <i>Science Advances</i> , 2018, 4, eaau6762.	10.3	33
26	Role of p53 and transcription-independent p53-induced apoptosis in shear-stimulated megakaryocytic maturation, particle generation, and platelet biogenesis. <i>PLoS ONE</i> , 2018, 13, e0203991.	2.5	12
27	RNAseq-based transcriptome assembly of <i>Clostridium acetobutylicum</i> for functional genome annotation and discovery. <i>AIChE Journal</i> , 2018, 64, 4271-4280.	3.6	6
28	Engineering <i>Clostridium</i> organisms as microbial cell-factories: challenges & opportunities. <i>Metabolic Engineering</i> , 2018, 50, 173-191.	7.0	56
29	Small and Low but Potent: the Complex Regulatory Role of the Small RNA SolB in Solventogenesis in <i>Clostridium acetobutylicum</i> . <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	12
30	Sort-Seq Approach to Engineering a Formaldehyde-Inducible Promoter for Dynamically Regulated <i>Escherichia coli</i> Growth on Methanol. <i>ACS Synthetic Biology</i> , 2017, 6, 1584-1595.	3.8	70
31	Heterologous Expression of the <i>Clostridium carboxidivorans</i> CO Dehydrogenase Alone or Together with the Acetyl Coenzyme A Synthase Enables both Reduction of CO ₂ and Oxidation of CO by <i>Clostridium acetobutylicum</i> . <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	21
32	In vitro methanol production from methyl coenzyme M using the <i>Methanosarcina barkeri</i> MtaABC protein complex. <i>Biotechnology Progress</i> , 2017, 33, 1243-1249.	2.6	10
33	How do megakaryocytic microparticles target and deliver cargo to alter the fate of hematopoietic stem cells?. <i>Journal of Controlled Release</i> , 2017, 247, 1-18.	9.9	58
34	Engineering the biological conversion of methanol to specialty chemicals in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2017, 39, 49-59.	7.0	137
35	Megakaryocytic Maturation in Response to Shear Flow Is Mediated by the Activator Protein 1 (AP-1) Transcription Factor via Mitogen-activated Protein Kinase (MAPK) Mechanotransduction. <i>Journal of Biological Chemistry</i> , 2016, 291, 7831-7843.	3.4	21
36	Stable and enhanced gene expression in <i>Clostridium acetobutylicum</i> using synthetic untranslated regions with a stem-loop. <i>Journal of Biotechnology</i> , 2016, 230, 40-43.	3.8	13

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37	Engineering membrane and cell-wall programs for tolerance to toxic chemicals: Beyond solo genes. <i>Current Opinion in Microbiology</i> , 2016, 33, 56-66.	5.1	66
38	Scaffoldless engineered enzyme assembly for enhanced methanol utilization. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 12691-12696.	7.1	93
39	CO2 fixation by anaerobic non-photosynthetic mixotrophy for improved carbon conversion. <i>Nature Communications</i> , 2016, 7, 12800.	12.8	128
40	Whole-genome sequence of an evolved <i>Clostridium pasteurianum</i> strain reveals Spo0A deficiency responsible for increased butanol production and superior growth. <i>Biotechnology for Biofuels</i> , 2015, 8, 227.	6.2	35
41	Expression of heterologous sigma factors enables functional screening of metagenomic and heterologous genomic libraries. <i>Nature Communications</i> , 2015, 6, 7045.	12.8	55
42	The <i>Clostridium</i> Sporulation Programs: Diversity and Preservation of Endospore Differentiation. <i>Microbiology and Molecular Biology Reviews</i> , 2015, 79, 19-37.	6.6	155
43	Editorial overview: Energy biotechnology. <i>Current Opinion in Biotechnology</i> , 2015, 33, viii-xi.	6.6	2
44	Synthetic methylotrophy: engineering the production of biofuels and chemicals based on the biology of aerobic methanol utilization. <i>Current Opinion in Biotechnology</i> , 2015, 33, 165-175.	6.6	150
45	Reassessing the Progress in the Production of Advanced Biofuels in the Current Competitive Environment and Beyond: What Are the Successes and Where Progress Eludes Us and Why. <i>Industrial & Engineering Chemistry Research</i> , 2015, 54, 10170-10182.	3.7	24
46	Building cellular pathways and programs enabled by the genetic diversity of allo-genomes and meta-genomes. <i>Current Opinion in Biotechnology</i> , 2015, 36, 16-31.	6.6	1
47	Capturing the response of <i>Clostridium acetobutylicum</i> to chemical stressors using a regulated genome-scale metabolic model. <i>Biotechnology for Biofuels</i> , 2014, 7, 144.	6.2	56
48	σK of <i>Clostridium acetobutylicum</i> Is the First Known Sporulation-Specific Sigma Factor with Two Developmentally Separated Roles, One Early and One Late in Sporulation. <i>Journal of Bacteriology</i> , 2014, 196, 287-299.	2.2	48
49	Overexpression of the <i>Lactobacillus plantarum</i> peptidoglycan biosynthesis murA2 gene increases the tolerance of <i>Escherichia coli</i> to alcohols and enhances ethanol production. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 8399-8411.	3.6	23
50	Shear enhances thrombopoiesis and formation of microparticles that induce megakaryocytic differentiation of stem cells. <i>Blood</i> , 2014, 124, 2094-2103.	1.4	71
51	Exploring the Capabilities of the Geobiosphere's Microbial Genome. <i>AIChE Journal</i> , 2013, 59, 688-698.	3.6	5
52	The <i>Clostridium</i> small RNome that responds to stress: the paradigm and importance of toxic metabolite stress in <i>C. acetobutylicum</i> . <i>BMC Genomics</i> , 2013, 14, 849.	2.8	49
53	Transcription factors and genetic circuits orchestrating the complex, multilayered response of <i>Clostridium acetobutylicum</i> to butanol and butyrate stress. <i>BMC Systems Biology</i> , 2013, 7, 120.	3.0	65
54	Three-Stage <i>Ex Vivo</i> Expansion of High-Ploidy Megakaryocytic Cells: Toward Large-Scale Platelet Production. <i>Tissue Engineering - Part A</i> , 2013, 19, 998-1014.	3.1	55

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55	Workflow for quantitative proteomic analysis of <i>Clostridium acetobutylicum</i> ATCC 824 using iTRAQ tags. <i>Methods</i> , 2013, 61, 269-276.	3.8	19
56	Stemâ€Cell Niche Based Comparative Analysis of Chemical and Nanoâ€mechanical Material Properties Impacting Ex Vivo Expansion and Differentiation of Hematopoietic and Mesenchymal Stem Cells. <i>Advanced Healthcare Materials</i> , 2013, 2, 25-42.	7.6	63
57	Synthetic tolerance: three noncoding small RNAs, DsrA, ArcZ and RprA, acting supra-additively against acid stress. <i>Nucleic Acids Research</i> , 2013, 41, 8726-8737.	14.5	102
58	Novel System for Efficient Isolation of <i>Clostridium</i> Double-Crossover Allelic Exchange Mutants Enabling Markerless Chromosomal Gene Deletions and DNA Integration. <i>Applied and Environmental Microbiology</i> , 2012, 78, 8112-8121.	3.1	113
59	Proposed megakaryocytic regulon of p53: the genes engaged to control cell cycle and apoptosis during megakaryocytic differentiation. <i>Physiological Genomics</i> , 2012, 44, 638-650.	2.3	26
60	<i>Clostridia</i> : the importance of their exceptional substrate and metabolite diversity for biofuel and biorefinery applications. <i>Current Opinion in Biotechnology</i> , 2012, 23, 364-381.	6.6	364
61	Stoichiometric and energetic analyses of non-photosynthetic CO ₂ -fixation pathways to support synthetic biology strategies for production of fuels and chemicals. <i>Current Opinion in Chemical Engineering</i> , 2012, 1, 380-395.	7.8	204
62	Role of tumor suppressor p53 in megakaryopoiesis and platelet function. <i>Experimental Hematology</i> , 2012, 40, 131-142.e4.	0.4	33
63	The Bleeding Defect Exhibited by Aryl Hydrocarbon Receptor-Null Mice Is Due to Defective Collagen-Dependent Outside-in Signaling. <i>Blood</i> , 2012, 120, 3294-3294.	1.4	0
64	Inactivation of <i>Ĥf</i> ^E and <i>Ĥf</i> ^G in <i>Clostridium acetobutylicum</i> Illuminates Their Roles in Clostridial-Cell-Form Biogenesis, Granulose Synthesis, Solventogenesis, and Spore Morphogenesis. <i>Journal of Bacteriology</i> , 2011, 193, 1414-1426.	2.2	78
65	The aryl hydrocarbon receptor (AHR) transcription factor regulates megakaryocytic polyploidization. <i>British Journal of Haematology</i> , 2011, 152, 469-484.	2.5	37
66	SpoIIIE Is Necessary for Asymmetric Division, Sporulation, and Expression of <i>Ĥf</i> ^F , <i>Ĥf</i> ^E , and <i>Ĥf</i> ^G but Does Not Control Solvent Production in <i>Clostridium acetobutylicum</i> ATCC 824. <i>Journal of Bacteriology</i> , 2011, 193, 5130-5137.	2.2	47
67	Inactivation of <i>Ĥf</i> ^F in <i>Clostridium acetobutylicum</i> ATCC 824 Blocks Sporulation Prior to Asymmetric Division and Abolishes <i>Ĥf</i> ^E and <i>Ĥf</i> ^G Protein Expression but Does Not Block Solvent Formation. <i>Journal of Bacteriology</i> , 2011, 193, 2429-2440.	2.2	62
68	Small RNAs in the Genus <i>Clostridium</i> . <i>MBio</i> , 2011, 2, e00340-10.	4.1	75
69	The Aryl Hydrocarbon Receptor Influences Multiple Stages of Megakaryocyte Differentiation. <i>Blood</i> , 2011, 118, 1298-1298.	1.4	0
70	The Importance of Matrix Elasticity and Shear Force on Megakaryocytic Differentiation. <i>Blood</i> , 2011, 118, 1329-1329.	1.4	9
71	Flow cytometry for bacteria: enabling metabolic engineering, synthetic biology and the elucidation of complex phenotypes. <i>Current Opinion in Biotechnology</i> , 2010, 21, 85-99.	6.6	137
72	Metabolic flux analysis of embryonic stem cells using three distinct differentiation protocols and comparison to gene expression patterns. <i>Biotechnology Progress</i> , 2010, 26, 1222-1229.	2.6	11

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73	Metabolite stress and tolerance in the production of biofuels and chemicals: Gene expression-based systems analysis of butanol, butyrate, and acetate stresses in the anaerobe <i>Clostridium acetobutylicum</i> . <i>Biotechnology and Bioengineering</i> , 2010, 105, 1131-1147.	3.3	191
74	A genomic-library based discovery of a novel, possibly synthetic, acid-tolerance mechanism in <i>Clostridium acetobutylicum</i> involving non-coding RNAs and ribosomal RNA processing. <i>Metabolic Engineering</i> , 2010, 12, 268-281.	7.0	70
75	A comparative view of metabolite and substrate stress and tolerance in microbial bioprocessing: From biofuels and chemicals, to biocatalysis and bioremediation. <i>Metabolic Engineering</i> , 2010, 12, 307-331.	7.0	478
76	Hes1 Expression Impacts Megakaryocytic Polyploidization and May Contribute to the Platelet Defects Found In Aryl Hydrocarbon Receptor (AhR)-Null Mice.. <i>Blood</i> , 2010, 116, 2612-2612.	1.4	0
77	Mechanistic studies on the effects of nicotinamide on megakaryocytic polyploidization and the roles of NAD ⁺ levels and SIRT inhibition. <i>Experimental Hematology</i> , 2009, 37, 1340-1352.e3.	0.4	38
78	Aldehyde-alcohol dehydrogenase and/or thiolase overexpression coupled with CoA transferase downregulation lead to higher alcohol titers and selectivity in <i>Clostridium acetobutylicum</i> fermentations. <i>Biotechnology and Bioengineering</i> , 2009, 102, 38-49.	3.3	123
79	Metabolic engineering of <i>Clostridium acetobutylicum</i> M5 for highly selective butanol production. <i>Biotechnology Journal</i> , 2009, 4, 1432-1440.	3.5	117
80	Engineering solventogenic clostridia. <i>Current Opinion in Biotechnology</i> , 2008, 19, 420-429.	6.6	302
81	Genome-scale model for <i>Clostridium acetobutylicum</i> : Part II. Development of specific proton flux states and numerically determined subsystems. <i>Biotechnology and Bioengineering</i> , 2008, 101, 1053-1071.	3.3	73
82	Genome-scale model for <i>Clostridium acetobutylicum</i> : Part I. Metabolic network resolution and analysis. <i>Biotechnology and Bioengineering</i> , 2008, 101, 1036-1052.	3.3	166
83	Metabolic engineering of the non-sporulating, non-solventogenic <i>Clostridium acetobutylicum</i> strain M5 to produce butanol without acetone demonstrate the robustness of the acid-formation pathways and the importance of the electron balance. <i>Metabolic Engineering</i> , 2008, 10, 321-332.	7.0	152
84	Comparative Transcriptional Analysis of Embryoid Body Versus Two-Dimensional Differentiation of Murine Embryonic Stem Cells. <i>Tissue Engineering - Part A</i> , 2008, 14, 1603-1614.	3.1	11
85	The transcriptional program underlying the physiology of clostridial sporulation. <i>Genome Biology</i> , 2008, 9, R114.	9.6	159
86	Tumor Suppressor Protein p53 Regulates Megakaryocytic Polyploidization and Apoptosis. <i>Journal of Biological Chemistry</i> , 2008, 283, 15589-15600.	3.4	38
87	Development and Application of Flow-Cytometric Techniques for Analyzing and Sorting Endospore-Forming Clostridia. <i>Applied and Environmental Microbiology</i> , 2008, 74, 7497-7506.	3.1	73
88	Gene Ontology-driven transcriptional analysis of CD34 ⁺ cell-initiated megakaryocytic cultures identifies new transcriptional regulators of megakaryopoiesis. <i>Physiological Genomics</i> , 2008, 33, 159-169.	2.3	23
89	Tumor Suppressor Protein p53 Affects Megakaryocytic Maturation: In Vivo and Ex Vivo Post-Translational Modification Studies. <i>Blood</i> , 2008, 112, 2443-2443.	1.4	1
90	Deregulated CDC25A Expression Promotes Mammary Tumorigenesis with Genomic Instability. <i>Cancer Research</i> , 2007, 67, 984-991.	0.9	70

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91	A General Framework for Designing and Validating Oligomer-Based DNA Microarrays and Its Application to <i>Clostridium acetobutylicum</i> . <i>Applied and Environmental Microbiology</i> , 2007, 73, 4631-4638.	3.1	22
92	A systems-biology analysis of isogenic megakaryocytic and granulocytic cultures identifies new molecular components of megakaryocytic apoptosis. <i>BMC Genomics</i> , 2007, 8, 384.	2.8	18
93	Comparative, genome-scale transcriptional analysis of CHRF-288-11 and primary human megakaryocytic cell cultures provides novel insights into lineage-specific differentiation. <i>Experimental Hematology</i> , 2007, 35, 476-489.e23.	0.4	42
94	Nicotinamide (vitamin B3) increases the polyploidisation and proplatelet formation of cultured primary human megakaryocytes. <i>British Journal of Haematology</i> , 2006, 135, 554-566.	2.5	63
95	A comparative genomic view of clostridial sporulation and physiology. <i>Nature Reviews Microbiology</i> , 2005, 3, 969-978.	28.6	295
96	Transcriptional Program of Early Sporulation and Stationary-Phase Events in <i>Clostridium acetobutylicum</i> . <i>Journal of Bacteriology</i> , 2005, 187, 7103-7118.	2.2	142
97	Transcriptional Analysis of <i>spo0A</i> Overexpression in <i>Clostridium acetobutylicum</i> and Its Effect on the Cell's Response to Butanol Stress. <i>Journal of Bacteriology</i> , 2004, 186, 1959-1971.	2.2	161
98	Transcriptional Analysis of Butanol Stress and Tolerance in <i>Clostridium acetobutylicum</i> . <i>Journal of Bacteriology</i> , 2004, 186, 2006-2018.	2.2	212
99	Overexpression of <i>groESL</i> in <i>Clostridium acetobutylicum</i> Results in Increased Solvent Production and Tolerance, Prolonged Metabolism, and Changes in the Cell's Transcriptional Program. <i>Applied and Environmental Microbiology</i> , 2003, 69, 4951-4965.	3.1	339
100	Design of Antisense RNA Constructs for Downregulation of the Acetone Formation Pathway of <i>Clostridium acetobutylicum</i> . <i>Journal of Bacteriology</i> , 2003, 185, 1923-1934.	2.2	159
101	DNA Array-Based Transcriptional Analysis of Asporogenous, Nonsolventogenic <i>Clostridium acetobutylicum</i> Strains SKO1 and M5. <i>Journal of Bacteriology</i> , 2003, 185, 4539-4547.	2.2	104
102	Antisense RNA Downregulation of Coenzyme A Transferase Combined with Alcohol-Aldehyde Dehydrogenase Overexpression Leads to Predominantly Alcohologenic <i>Clostridium acetobutylicum</i> Fermentations. <i>Journal of Bacteriology</i> , 2003, 185, 3644-3653.	2.2	132
103	Northern, Morphological, and Fermentation Analysis of <i>spo0A</i> Inactivation and Overexpression in <i>Clostridium acetobutylicum</i> ATCC 824. <i>Journal of Bacteriology</i> , 2002, 184, 3586-3597.	2.2	212
104	Extracellular pH Affects the Proliferation of Cultured Human T Cells and Their Expression of the Interleukin-2 Receptor. <i>Journal of Immunotherapy</i> , 2000, 23, 669-674.	2.4	25
105	Sparging and agitation-induced injury of cultured animals cells: Do cell-to-bubble interactions in the bulk liquid injure cells?. , 2000, 51, 399-409.		42
106	Characterization of recombinant strains of the <i>Clostridium acetobutylicum</i> butyrate kinase inactivation mutant: Need for new phenomenological models for solventogenesis and butanol inhibition?. <i>Biotechnology and Bioengineering</i> , 2000, 67, 1-11.	3.3	167
107	Equations and calculations for fermentations of butyric acid bacteria. <i>Biotechnology and Bioengineering</i> , 2000, 67, 813-826.	3.3	15
108	Culture of human T cells in stirred bioreactors for cellular immunotherapy applications: Shear, proliferation, and the IL-2 receptor. , 2000, 68, 328-338.		53

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109	Characterization of recombinant strains of the Clostridium acetobutylicum butyrate kinase inactivation mutant: Need for new phenomenological models for solventogenesis and butanol inhibition?. , 2000, 67, 1.		2
110	Development and Characterization of a Gene Expression Reporter System for Clostridium acetobutylicum ATCC 824. Applied and Environmental Microbiology, 1999, 65, 3793-3799.	3.1	109
111	Regulation of the sol Locus Genes for Butanol and Acetone Formation in Clostridium acetobutylicum ATCC 824 by a Putative Transcriptional Repressor. Journal of Bacteriology, 1999, 181, 319-330.	2.2	95
112	Increased agitation intensity increases CD13 receptor surface content and mRNA levels, and alters the metabolism of HL60 cells cultured in stirred tank bioreactors. , 1998, 60, 239-250.		47
113	Effects of methocel A15LV, polyethylene glycol, and polyvinyl alcohol on CD13 and CD33 receptor surface content and metabolism of HL60 cells cultured in stirred tank bioreactors. , 1998, 60, 251-258.		5
114	Serum increases the CD13 receptor expression, reduces the transduction of fluid-mechanical forces, and alters the metabolism of HL60 cells cultured in agitated bioreactors. , 1998, 60, 259-268.		14
115	Express together and conquer. Nature Biotechnology, 1998, 16, 416-417.	17.5	17
116	Sparging and agitation-induced injury of cultured animals cells: Do cell-to-cell interactions in the bulk liquid injure cells?. Biotechnology and Bioengineering, 1996, 51, 399-409.	3.3	24
117	Analysis of cell-to-bubble attachment in sparged bioreactors in the presence of cell-protecting additives. Biotechnology and Bioengineering, 1995, 47, 407-419.	3.3	76
118	Interfacial properties of cell culture media with cell-protecting additives. Biotechnology and Bioengineering, 1995, 47, 420-430.	3.3	55
119	Ammonia affects the glycosylation patterns of recombinant mouse placental lactogen-I by chinese hamster ovary cells in a pH-dependent manner. Biotechnology and Bioengineering, 1994, 43, 505-514.	3.3	123
120	Serum-free media for cultures of primitive and mature hematopoietic cells. Biotechnology and Bioengineering, 1994, 43, 706-733.	3.3	38
121	Host-plasmid interactions in recombinant strains of Clostridium acetobutylicum ATCC 824. FEMS Microbiology Letters, 1994, 123, 335-341.	1.8	13
122	Fluid-mechanical forces in agitated bioreactors reduce the CD13 and CD33 surface protein content of HL60 cells. Biotechnology and Bioengineering, 1993, 41, 868-877.	3.3	19
123	Metabolic engineering of Clostridium acetobutylicum ATCC 824 for increased solvent production by enhancement of acetone formation enzyme activities using a synthetic acetone operon. Biotechnology and Bioengineering, 1993, 42, 1053-1060.	3.3	98
124	Expansion of Primitive Human Hematopoietic Progenitors in a Perfusion Bioreactor System with IL-3, IL-6, and Stem Cell Factor. Bio/technology, 1993, 11, 358-363.	1.5	94
125	Culture pH Affects Expression Rates and Glycosylation of Recombinant Mouse Placental Lactogen Proteins by Chinese Hamster Ovary (CHO) Cells. Nature Biotechnology, 1993, 11, 720-724.	17.5	115
126	Expression of Cloned Homologous Fermentative Genes in Clostridium Acetobutylicum ATCC 824. Nature Biotechnology, 1992, 10, 190-195.	17.5	209

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127	Vector Construction, Transformation, and Gene Amplification in <i>Clostridium acetobutylicum</i> ATCC 824. <i>Annals of the New York Academy of Sciences</i> , 1992, 665, 39-51.	3.8	32
128	FLUID-MECHANICAL DAMAGE OF FREELY-SUSPENDED ANIMAL CELLS IN AGITATED BIOREACTORS: EFFECTS OF DEXTRAN, DERIVATIZED CELLULOSES AND POLYVINYL ALCOHOL. <i>Chemical Engineering Communications</i> , 1992, 118, 341-360.	2.6	18
129	Construction of <i>Escherichia coli</i> - <i>Clostridium acetobutylicum</i> shuttle vectors and transformation of <i>Clostridium acetobutylicum</i> strains. <i>Biotechnology Letters</i> , 1992, 14, 427-432.	2.2	33
130	Agitation induced cell injury in microcarrier cultures. Protective effect of viscosity is agitation intensity dependent: Experiments and modeling. <i>Biotechnology and Bioengineering</i> , 1992, 39, 95-107.	3.3	53
131	Damaging agitation intensities increase DNA synthesis rate and alter cell-cycle phase distributions of CHO cells. <i>Biotechnology and Bioengineering</i> , 1992, 40, 978-990.	3.3	51
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#	ARTICLE	IF	CITATIONS
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