George N Bennett

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.

192
papers

8,377
citations

51
h-index

83
g-index

195
ext. papers

9,053
ext. citations

5.7
avg, IF

L-index

#	Paper	IF	Citations
192	Metabolic engineering of Escherichia coli for quinolinic acid production by assembling L-aspartate oxidase and quinolinate synthase as an enzyme complex. <i>Metabolic Engineering</i> , 2021 , 67, 164-172	9.7	2
191	100th Anniversary of Macromolecular Science Viewpoint: Soft Materials for Microbial Bioelectronics. <i>ACS Macro Letters</i> , 2020 , 9, 1590-1603	6.6	9
190	phage ferredoxin: structural characterization and electron transfer to cyanobacterial sulfite reductases. <i>Journal of Biological Chemistry</i> , 2020 , 295, 10610-10623	5.4	3
189	Localized mandibular infection affects remote in vivo bioreactor bone generation. <i>Biomaterials</i> , 2020 , 256, 120185	15.6	3
188	Improving the organization and interactivity of metabolic pathfinding with precomputed pathways. <i>BMC Bioinformatics</i> , 2020 , 21, 13	3.6	7
187	Metabolic engineering of Escherichia coli to produce succinate from woody hydrolysate under anaerobic conditions. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2020 , 47, 223-232	4.2	4
186	Combinatorial design of chemical-dependent protein switches for controlling intracellular electron transfer. <i>AICHE Journal</i> , 2020 , 66, e16796	3.6	3
185	Improved succinate production from galactose-rich feedstocks by engineered Escherichia coli under anaerobic conditions. <i>Biotechnology and Bioengineering</i> , 2020 , 117, 1082-1091	4.9	3
184	Metabolic engineering of Escherichia coli for malate production with a temperature sensitive malate dehydrogenase. <i>Biochemical Engineering Journal</i> , 2020 , 164, 107762	4.2	1
183	Recombination of 2Fe-2S Ferredoxins Reveals Differences in the Inheritance of Thermostability and Midpoint Potential. <i>ACS Synthetic Biology</i> , 2020 , 9, 3245-3253	5.7	O
182	Single cell protein production from food waste using purple non-sulfur bacteria shows economically viable protein products have higher environmental impacts. <i>Journal of Cleaner Production</i> , 2020 , 276, 123114	10.3	8
181	Genetic sensor-regulators functional in Clostridia. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2020 , 47, 609-620	4.2	1
180	Evolutionary Relationships Between Low Potential Ferredoxin and Flavodoxin Electron Carriers. <i>Frontiers in Energy Research</i> , 2019 , 7,	3.8	13
179	De novo design of symmetric ferredoxins that shuttle electrons in vivo. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 14557-14562	11.5	23
178	Biosynthesis of Medium-Chain EHydroxy Fatty Acids by AlkBGT of GPo1 With Native FadL in Engineered. <i>Frontiers in Bioengineering and Biotechnology</i> , 2019 , 7, 273	5.8	8
177	Metalloprotein switches that display chemical-dependent electron transfer in cells. <i>Nature Chemical Biology</i> , 2019 , 15, 189-195	11.7	19
176	Metabolic engineering of Escherichia coli to produce succinate from soybean hydrolysate under anaerobic conditions. <i>Biotechnology and Bioengineering</i> , 2018 , 115, 1743-1754	4.9	12

175	Ratiometric Gas Reporting: A Nondisruptive Approach To Monitor Gene Expression in Soils. <i>ACS Synthetic Biology</i> , 2018 , 7, 903-911	5.7	8
174	Improvement of butanol production in Clostridium acetobutylicum through enhancement of NAD(P)H availability. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2018 , 45, 993-1002	4.2	20
173	Bioconversion of methane to C-4 carboxylic acids using carbon flux through acetyl-CoA in engineered Methylomicrobium buryatense 5GB1C. <i>Metabolic Engineering</i> , 2018 , 48, 175-183	9.7	22
172	High yield production of four-carbon dicarboxylic acids by metabolically engineered Escherichia coli. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2018 , 45, 53-60	4.2	16
171	Genome analysis of a hyper acetone-butanol-ethanol (ABE) producing Clostridium acetobutylicum BKM19. <i>Biotechnology Journal</i> , 2017 , 12, 1600457	5.6	9
170	Effects of Local Antibiotic Delivery from Porous Space Maintainers on Infection Clearance and Induction of an Osteogenic Membrane in an Infected Bone Defect. <i>Tissue Engineering - Part A</i> , 2017 , 23, 91-100	3.9	22
169	Strategies for manipulation of oxygen utilization by the electron transfer chain in microbes for metabolic engineering purposes. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2017 , 44, 647-658	34.2	8
168	Role of Clostridial Nitroreductases in Bioremediation 2017 , 175-186		2
167	Use of transposase and ends of IS608 enables precise and scarless genome modification for modulating gene expression and metabolic engineering applications in Escherichia coli. <i>Biotechnology Journal</i> , 2016 , 11, 80-90	5.6	2
166	Cellular Assays for Ferredoxins: A Strategy for Understanding Electron Flow through Protein Carriers That Link Metabolic Pathways. <i>Biochemistry</i> , 2016 , 55, 7047-7064	3.2	23
165	Volatile Gas Production by Methyl Halide Transferase: An In Situ Reporter Of Microbial Gene Expression In Soil. <i>Environmental Science & Expression In Soil. Environmental Expression In Soil Environmental Expression </i>	10.3	10
164	A rapid, flexible method for incorporating controlled antibiotic release into porous polymethylmethacrylate space maintainers for craniofacial reconstruction. <i>Biomaterials Science</i> , 2016 , 4, 121-9	7.4	7
163	Polymer-Based Local Antibiotic Delivery for Prevention of Polymicrobial Infection in Contaminated Mandibular Implants. <i>ACS Biomaterials Science and Engineering</i> , 2016 , 2, 558-566	5.5	15
162	Direct bioconversion of sorghum extract sugars to free fatty acids using metabolically engineered Escherichia coli strains: Value addition to the sorghum bioenergy crop. <i>Biomass and Bioenergy</i> , 2016 , 93, 217-226	5.3	3
161	Metabolic transistor strategy for controlling electron transfer chain activity in Escherichia coli. <i>Metabolic Engineering</i> , 2015 , 28, 159-168	9.7	15
160	Metabolic control of respiratory levels in coenzyme Q biosynthesis-deficient Escherichia coli strains leading to fine-tune aerobic lactate fermentation. <i>Biotechnology and Bioengineering</i> , 2015 , 112, 1720-6	4.9	9
159	Metabolic engineering of carbon and redox flow in the production of small organic acids. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2015 , 42, 403-22	4.2	38
158	Efficient production of free fatty acids from soybean meal carbohydrates. <i>Biotechnology and Bioengineering</i> , 2015 , 112, 2324-33	4.9	13

157	Efficient free fatty acid production in engineered Escherichia coli strains using soybean oligosaccharides as feedstock. <i>Biotechnology Progress</i> , 2015 , 31, 686-94	2.8	10
156	Proteomic analyses of the phase transition from acidogenesis to solventogenesis using solventogenic and non-solventogenic Clostridium acetobutylicum strains. <i>Applied Microbiology and Biotechnology</i> , 2014 , 98, 5105-15	5.7	26
155	Effects of antibiotic physicochemical properties on their release kinetics from biodegradable polymer microparticles. <i>Pharmaceutical Research</i> , 2014 , 31, 3379-89	4.5	33
154	Increased Biofuel Production by Metabolic Engineering of Clostridium acetobutylicum 2014 , 361-376		1
153	Soybean Carbohydrates as a Renewable Feedstock for the Fermentative Production of Succinic Acid and Ethanol. <i>ACS Symposium Series</i> , 2014 , 81-107	0.4	0
152	Metabolic engineering of Escherichia coli to minimize byproduct formate and improving succinate productivity through increasing NADH availability by heterologous expression of NAD(+)-dependent formate dehydrogenase. <i>Metabolic Engineering</i> , 2013 , 20, 1-8	9.7	81
151	Characterization and evaluation of corn steep liquid in acetone-butanol-ethanol production by Clostridium acetobutylicum. <i>Biotechnology and Bioprocess Engineering</i> , 2013 , 18, 266-271	3.1	9
150	Evaluation of antibiotic releasing porous polymethylmethacrylate space maintainers in an infected composite tissue defect model. <i>Acta Biomaterialia</i> , 2013 , 9, 8832-9	10.8	20
149	Improvement of NADPH bioavailability in Escherichia coli by replacing NAD(+)-dependent glyceraldehyde-3-phosphate dehydrogenase GapA with NADP (+)-dependent GapB from Bacillus subtilis and addition of NAD kinase. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2013 , 40, 1449	4.2 -60	20
148	Production of succinic acid by engineered E. coli strains using soybean carbohydrates as feedstock under aerobic fermentation conditions. <i>Bioresource Technology</i> , 2013 , 130, 398-405	11	47
147	Metabolic engineering and transhydrogenase effects on NADPH availability in Escherichia coli. <i>Biotechnology Progress</i> , 2013 , 29, 1124-30	2.8	31
146	Cofactor engineering for advancing chemical biotechnology. <i>Current Opinion in Biotechnology</i> , 2013 , 24, 994-9	11.4	105
145	Improvement of NADPH bioavailability in Escherichia coli through the use of phosphofructokinase deficient strains. <i>Applied Microbiology and Biotechnology</i> , 2013 , 97, 6883-93	5.7	22
144	Analysis of redox responses during TNT transformation by Clostridium acetobutylicum ATCC 824 and mutants exhibiting altered metabolism. <i>Applied Microbiology and Biotechnology</i> , 2013 , 97, 4651-63	5.7	8
143	Metabolic engineering of Clostridium acetobutylicum ATCC 824 for isopropanol-butanol-ethanol fermentation. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 1416-23	4.8	190
142	Succinate production in Escherichia coli. <i>Biotechnology Journal</i> , 2012 , 7, 213-24	5.6	138
141	Manipulating respiratory levels in Escherichia coli for aerobic formation of reduced chemical products. <i>Metabolic Engineering</i> , 2011 , 13, 704-12	9.7	23
140	Improving the Clostridium acetobutylicum butanol fermentation by engineering the strain for co-production of riboflavin. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2011 , 38, 1013-25	4.2	32

(2008-2011)

139	Effect of culture operating conditions on succinate production in a multiphase fed-batch bioreactor using an engineered Escherichia coli strain. <i>Applied Microbiology and Biotechnology</i> , 2011 , 92, 499-508	5.7	21
138	Culture conditionsSimpact on succinate production by a high succinate producing Escherichia coli strain. <i>Biotechnology Progress</i> , 2011 , 27, 1225-31	2.8	8
137	Succinate production from sucrose by metabolic engineered Escherichia coli strains under aerobic conditions. <i>Biotechnology Progress</i> , 2011 , 27, 1242-7	2.8	10
136	Heterologous pyc gene expression under various natural and engineered promoters in Escherichia coli for improved succinate production. <i>Journal of Biotechnology</i> , 2011 , 155, 236-43	3.7	26
135	Succinate production from different carbon sources under anaerobic conditions by metabolic engineered Escherichia coli strains. <i>Metabolic Engineering</i> , 2011 , 13, 328-35	9.7	45
134	An algorithm for efficient identification of branched metabolic pathways. <i>Journal of Computational Biology</i> , 2011 , 18, 1575-97	1.7	11
133	Finding metabolic pathways using atom tracking. <i>Bioinformatics</i> , 2010 , 26, 1548-55	7.2	47
132	Structural correlations of activity of Clostridium acetobutylicum ATCC 824 butyrate kinase isozymes. <i>Enzyme and Microbial Technology</i> , 2010 , 46, 118-124	3.8	6
131	Metabolic impact of the level of aeration during cell growth on anaerobic succinate production by an engineered Escherichia coli strain. <i>Metabolic Engineering</i> , 2010 , 12, 499-509	9.7	40
130	Metabolic flux analysis of Escherichia coli creB and arcA mutants reveals shared control of carbon catabolism under microaerobic growth conditions. <i>Journal of Bacteriology</i> , 2009 , 191, 5538-48	3.5	37
129	Metabolic engineering of the anaerobic central metabolic pathway in Escherichia coli for the simultaneous anaerobic production of isoamyl acetate and succinic acid. <i>Biotechnology Progress</i> , 2009 , 25, 1304-9	2.8	7
128	Microbial formation of esters. Applied Microbiology and Biotechnology, 2009, 85, 13-25	5.7	88
127	Environmentally-modulated changes in fluorescence distribution in cells with oscillatory genetic network dynamics. <i>Journal of Biotechnology</i> , 2009 , 140, 203-17	3.7	4
126	Clostridium taeniosporum is a close relative of the Clostridium botulinum Group II. <i>Anaerobe</i> , 2008 , 14, 318-24	2.8	6
125	Activity of abrB310 promoter in wild type and spo0A-deficient strains of Clostridium acetobutylicum. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2008 , 35, 743-50	4.2	6
124	Engineering poly(3-hydroxybutyrate-co-3-hydroxyvalerate) copolymer composition in E. coli. <i>Biotechnology and Bioengineering</i> , 2008 , 99, 919-28	4.9	31
123	Reduction of acetate accumulation in Escherichia coli cultures for increased recombinant protein production. <i>Metabolic Engineering</i> , 2008 , 10, 97-108	9.7	49
122	Replacing Escherichia coli NAD-dependent glyceraldehyde 3-phosphate dehydrogenase (GAPDH) with a NADP-dependent enzyme from Clostridium acetobutylicum facilitates NADPH dependent pathways. <i>Metabolic Engineering</i> , 2008 , 10, 352-9	9.7	102

121	The YfiD protein contributes to the pyruvate formate-lyase flux in an Escherichia coli arcA mutant strain. <i>Biotechnology and Bioengineering</i> , 2007 , 97, 138-43	4.9	17
120	Analysis of the clostridial hydrophobic with a conserved tryptophan family (ChW) of proteins in Clostridium acetobutylicum with emphasis on ChW14 and ChW16/17. <i>Enzyme and Microbial Technology</i> , 2007 , 42, 29-43	3.8	7
119	Characterization of a novel ferredoxin with N-terminal extension from Clostridium acetobutylicum ATCC 824. <i>Archives of Microbiology</i> , 2007 , 187, 161-9	3	3
118	Characterization of alcohol dehydrogenase 1 and 3 from Neurospora crassa FGSC2489. <i>Applied Microbiology and Biotechnology</i> , 2007 , 76, 349-56	5.7	14
117	Cell population heterogeneity in expression of a gene-switching network with fluorescent markers of different half-lives. <i>Journal of Biotechnology</i> , 2007 , 128, 362-75	3.7	8
116	Acetyl-CoA synthetase overexpression in Escherichia coli demonstrates more efficient acetate assimilation and lower acetate accumulation: a potential tool in metabolic engineering. <i>Applied Microbiology and Biotechnology</i> , 2006 , 71, 870-4	5.7	95
115	A kinetic model of oxygen regulation of cytochrome production in Escherichia coli. <i>Journal of Theoretical Biology</i> , 2006 , 242, 547-63	2.3	12
114	Characterization of D-ribose biosynthesis in Bacillus subtilis JY200 deficient in transketolase gene. Journal of Biotechnology, 2006 , 121, 508-16	3.7	10
113	Expression of the pfl gene and resulting metabolite flux distribution in nuo and ackA-pta E. coli mutant strains. <i>Biotechnology Progress</i> , 2006 , 22, 898-902	2.8	2
112	Effect of overexpression of a soluble pyridine nucleotide transhydrogenase (UdhA) on the production of poly(3-hydroxybutyrate) in Escherichia coli. <i>Biotechnology Progress</i> , 2006 , 22, 420-5	2.8	81
111	Ester production in E. coli and C. acetobutylicum. Enzyme and Microbial Technology, 2006, 38, 937-943	3.8	24
110	Development of a metabolic network design and optimization framework incorporating implementation constraints: a succinate production case study. <i>Metabolic Engineering</i> , 2006 , 8, 46-57	9.7	38
109	Batch culture characterization and metabolic flux analysis of succinate-producing Escherichia coli strains. <i>Metabolic Engineering</i> , 2006 , 8, 209-26	9.7	73
108	Effect of the global redox sensing/regulation networks on Escherichia coli and metabolic flux distribution based on C-13 labeling experiments. <i>Metabolic Engineering</i> , 2006 , 8, 619-27	9.7	33
107	Proteome analysis and comparison of Clostridium acetobutylicum ATCC 824 and Spo0A strain variants. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2006 , 33, 298-308	4.2	46
106	Studies on inhibition of transformation of 2,4,6-trinitrotoluene catalyzed by Fe-only hydrogenase from Clostridium acetobutylicum. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2006 , 33, 368-76	4.2	7
105	Molecular cloning and characterization of the alcohol dehydrogenase ADH1 gene of Candida utilis ATCC 9950. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2006 , 33, 1032-6	4.2	26
104	Efficient succinic acid production from glucose through overexpression of pyruvate carboxylase in an Escherichia coli alcohol dehydrogenase and lactate dehydrogenase mutant. <i>Biotechnology</i>	2.8	99

(2005-2005)

103	Redistribution of metabolic fluxes in the central aerobic metabolic pathway of E. coli mutant strains with deletion of the ackA-pta and poxB pathways for the synthesis of isoamyl acetate. <i>Biotechnology Progress</i> , 2005 , 21, 627-31	2.8	57
102	Characterization of the acetate-producing pathways in Escherichia coli. <i>Biotechnology Progress</i> , 2005 , 21, 1062-7	2.8	99
101	Effect of different levels of NADH availability on metabolic fluxes of Escherichia coli chemostat cultures in defined medium. <i>Journal of Biotechnology</i> , 2005 , 117, 395-405	3.7	58
100	Metabolic engineering of aerobic succinate production systems in Escherichia coli to improve process productivity and achieve the maximum theoretical succinate yield. <i>Metabolic Engineering</i> , 2005 , 7, 116-27	9.7	161
99	Novel pathway engineering design of the anaerobic central metabolic pathway in Escherichia coli to increase succinate yield and productivity. <i>Metabolic Engineering</i> , 2005 , 7, 229-39	9.7	202
98	Chemostat culture characterization of Escherichia coli mutant strains metabolically engineered for aerobic succinate production: a study of the modified metabolic network based on metabolite profile, enzyme activity, and gene expression profile. <i>Metabolic Engineering</i> , 2005 , 7, 337-52	9.7	36
97	Effect of oxygen, and ArcA and FNR regulators on the expression of genes related to the electron transfer chain and the TCA cycle in Escherichia coli. <i>Metabolic Engineering</i> , 2005 , 7, 364-74	9.7	100
96	Genetically constrained metabolic flux analysis. <i>Metabolic Engineering</i> , 2005 , 7, 445-56	9.7	20
95	Enhanced lycopene productivity by manipulation of carbon flow to isopentenyl diphosphate in Escherichia coli. <i>Biotechnology Progress</i> , 2005 , 21, 1558-61	2.8	67
94	Genetic reconstruction of the aerobic central metabolism in Escherichia coli for the absolute aerobic production of succinate. <i>Biotechnology and Bioengineering</i> , 2005 , 89, 148-56	4.9	98
93	Effect of oxygen on the Escherichia coli ArcA and FNR regulation systems and metabolic responses. <i>Biotechnology and Bioengineering</i> , 2005 , 89, 556-64	4.9	99
92	Fed-batch culture of a metabolically engineered Escherichia coli strain designed for high-level succinate production and yield under aerobic conditions. <i>Biotechnology and Bioengineering</i> , 2005 , 90, 775-9	4.9	102
91	Effect of ArcA and FNR on the expression of genes related to the oxygen regulation and the glycolysis pathway in Escherichia coli under microaerobic growth conditions. <i>Biotechnology and Bioengineering</i> , 2005 , 92, 147-59	4.9	100
90	Effect of Sorghum vulgare phosphoenolpyruvate carboxylase and Lactococcus lactis pyruvate carboxylase coexpression on succinate production in mutant strains of Escherichia coli. <i>Applied Microbiology and Biotechnology</i> , 2005 , 67, 515-23	5.7	61
89	Biodegradation of xenobiotics by anaerobic bacteria. <i>Applied Microbiology and Biotechnology</i> , 2005 , 67, 600-18	5.7	114
88	Biochemical characterization of trinitrotoluene transforming oxygen-insensitive nitroreductases from Clostridium acetobutylicum ATCC 824. <i>Archives of Microbiology</i> , 2005 , 184, 158-67	3	39
87	Characterization of thermostable Xyn10A enzyme from mesophilic Clostridium acetobutylicum ATCC 824. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2005 , 32, 12-8	4.2	18
86	Effect of carbon sources differing in oxidation state and transport route on succinate production in metabolically engineered Escherichia coli. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2005, 32, 87-93	4.2	44

85	Intracellular butyryl phosphate and acetyl phosphate concentrations in Clostridium acetobutylicum and their implications for solvent formation. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 530-7	4.8	85
84	SpoIIE regulates sporulation but does not directly affect solventogenesis in Clostridium acetobutylicum ATCC 824. <i>Journal of Bacteriology</i> , 2005 , 187, 1930-6	3.5	42
83	Expression of abrB310 and SinR, and effects of decreased abrB310 expression on the transition from acidogenesis to solventogenesis, in Clostridium acetobutylicum ATCC 824. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 1987-95	4.8	38
82	Enhanced isoamyl acetate production upon manipulation of the acetyl-CoA node in Escherichia coli. <i>Biotechnology Progress</i> , 2004 , 20, 692-7	2.8	14
81	Increasing the acetyl-CoA pool in the presence of overexpressed phosphoenolpyruvate carboxylase or pyruvate carboxylase enhances succinate production in Escherichia coli. <i>Biotechnology Progress</i> , 2004 , 20, 1599-604	2.8	63
80	Thermostable xylanase10B from Clostridium acetobutylicum ATCC824. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2004 , 31, 229-34	4.2	17
79	Production of isoamyl acetate in ackA-pta and/or ldh mutants of Escherichia coli with overexpression of yeast ATF2. <i>Applied Microbiology and Biotechnology</i> , 2004 , 63, 698-704	5.7	34
78	Effect of different levels of NADH availability on metabolite distribution in Escherichia coli fermentation in minimal and complex media. <i>Applied Microbiology and Biotechnology</i> , 2004 , 65, 426-32	5.7	44
77	Cofactor engineering of intracellular CoA/acetyl-CoA and its effect on metabolic flux redistribution in Escherichia coli. <i>Metabolic Engineering</i> , 2004 , 6, 133-9	9.7	61
76	Applicability of CoA/acetyl-CoA manipulation system to enhance isoamyl acetate production in Escherichia coli. <i>Metabolic Engineering</i> , 2004 , 6, 294-9	9.7	41
75	2,4,6-trinitrotoluene reduction by an Fe-only hydrogenase in Clostridium acetobutylicum. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 1542-7	4.8	44
74	Mutagenicity of nitroaromatic degradation compounds. <i>Environmental Toxicology and Chemistry</i> , 2003 , 22, 2293-7	3.8	75
73	The effect of carbon sources and lactate dehydrogenase deletion on 1,2-propanediol production in Escherichia coli. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2003 , 30, 34-40	4.2	43
72	Sequences affecting the regulation of solvent production in Clostridium acetobutylicum. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2003 , 30, 414-20	4.2	11
71	Heterologous expression of the Saccharomyces cerevisiae alcohol acetyltransferase genes in Clostridium acetobutylicum and Escherichia coli for the production of isoamyl acetate. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2003 , 30, 427-32	4.2	29
70	Role of hydroxylamine intermediates in the phytotransformation of 2,4,6-trinitrotoluene by Myriophyllum aquaticum. <i>Environmental Science & Environmental Science & Environmen</i>	10.3	42
69	Expression of a cloned cyclopropane fatty acid synthase gene reduces solvent formation in Clostridium acetobutylicum ATCC 824. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 2831-41	4.8	87
68	Metabolic engineering through cofactor manipulation and its effects on metabolic flux redistribution in Escherichia coli. <i>Metabolic Engineering</i> , 2002 , 4, 182-92	9.7	203

(1999-2002)

67	Metabolic engineering of Escherichia coli: increase of NADH availability by overexpressing an NAD(+)-dependent formate dehydrogenase. <i>Metabolic Engineering</i> , 2002 , 4, 217-29	9.7	224
66	The effect of increasing NADH availability on the redistribution of metabolic fluxes in Escherichia coli chemostat cultures. <i>Metabolic Engineering</i> , 2002 , 4, 230-7	9.7	116
65	The effects of feed and intracellular pyruvate levels on the redistribution of metabolic fluxes in Escherichia coli. <i>Metabolic Engineering</i> , 2001 , 3, 115-23	9.7	59
64	Genome sequence and comparative analysis of the solvent-producing bacterium Clostridium acetobutylicum. <i>Journal of Bacteriology</i> , 2001 , 183, 4823-38	3.5	656
63	Effect of variation of Klebsiella pneumoniae acetolactate synthase expression on metabolic flux redistribution in Escherichia coli. <i>Biotechnology and Bioengineering</i> , 2000 , 69, 150-9	4.9	24
62	Effect of glucose analog supplementation on metabolic flux distribution in anaerobic chemostat cultures of Escherichia coli. <i>Metabolic Engineering</i> , 2000 , 2, 149-54	9.7	8
61	Mutagenicity of trinitrotoluene and metabolites formed during anaerobic degradation by Clostridium acetobutylicum ATCC 824. <i>Environmental Toxicology and Chemistry</i> , 2000 , 19, 2871-2875	3.8	15
60	Cloning, Sequencing, and Characterization of the Gene Encoding Flagellin, flaC, and the Post-translational Modification of Flagellin, FlaC, from Clostridium acetobutylicum ATCC824. <i>Anaerobe</i> , 2000 , 6, 69-79	2.8	19
59	2,4,6-trinitrotoluene reduction by carbon monoxide dehydrogenase from Clostridium thermoaceticum. <i>Applied and Environmental Microbiology</i> , 2000 , 66, 1474-8	4.8	66
58	. Environmental Toxicology and Chemistry, 2000 , 19, 2871	3.8	4
57	Improvement of biomass yield and recombinant gene expression in Escherichia coli by using fructose as the primary carbon source. <i>Biotechnology Progress</i> , 1999 , 15, 140-5	2.8	34
56	Metabolic flux analysis of Escherichia coli deficient in the acetate production pathway and expressing the Bacillus subtilis acetolactate synthase. <i>Metabolic Engineering</i> , 1999 , 1, 26-34	9.7	65
55	Redistribution of metabolic fluxes in Escherichia coli with fermentative lactate dehydrogenase overexpression and deletion. <i>Metabolic Engineering</i> , 1999 , 1, 141-52	9.7	62
54	Metabolic flux analysis of Escherichia coli expressing the Bacillus subtilis acetolactate synthase in batch and continuous cultures. <i>Biotechnology and Bioengineering</i> , 1999 , 63, 737-49	4.9	28
53	Effect of inactivation of nuo and ackA-pta on redistribution of metabolic fluxes in Escherichia coli. <i>Biotechnology and Bioengineering</i> , 1999 , 65, 291-297	4.9	45
52	Overexpression, purification, and characterization of the thermostable mevalonate kinase from Methanococcus jannaschii. <i>Protein Expression and Purification</i> , 1999 , 17, 33-40	2	35
51	Characterization of methylglyoxal synthase from Clostridium acetobutylicum ATCC 824 and its use in the formation of 1, 2-propanediol. <i>Applied and Environmental Microbiology</i> , 1999 , 65, 3244-7	4.8	35
50	Regulation of the sol locus genes for butanol and acetone formation in Clostridium acetobutylicum ATCC 824 by a putative transcriptional repressor. <i>Journal of Bacteriology</i> , 1999 , 181, 319-30	3.5	83

49	Reduction of 2,4,6-trinitrotoluene by Clostridium acetobutylicum through hydroxylamino-nitrotoluene intermediates. <i>Environmental Toxicology and Chemistry</i> , 1998 , 17, 343-348	3.8	51
48	Genetic manipulation of acid and solvent formation in clostridium acetobutylicum ATCC 824. <i>Biotechnology and Bioengineering</i> , 1998 , 58, 215-21	4.9	22
47	Complementation of an Escherichia coli polypeptide deformylase mutant with a gene from Clostridium acetobutylicum ATCC 824. <i>Current Microbiology</i> , 1998 , 36, 248-9	2.4	4
46	Cloning, sequence, and expression of the phosphofructokinase gene of Clostridium acetobutylicum ATCC 824 in Escherichia coli. <i>Current Microbiology</i> , 1998 , 37, 17-22	2.4	14
45	. Environmental Toxicology and Chemistry, 1998 , 17, 343	3.8	6
44	A method for construction of E. coli strains with multiple DNA insertions in the chromosome. <i>Gene</i> , 1997 , 187, 231-8	3.8	37
43	Cloning and assembly of PCR products using modified primers and DNA repair enzymes. <i>BioTechniques</i> , 1997 , 23, 858-62, 864	2.5	8
42	Escherichia coli strain for thermoinducible T7 RNA polymerase-driven expression. <i>Gene</i> , 1996 , 177, 267-	8 3.8	3
41	Inactivation of an aldehyde/alcohol dehydrogenase gene from Clostridium acetobutylicum ATCC 824. <i>Applied Biochemistry and Biotechnology</i> , 1996 , 57-58, 213-21	3.2	46
40	Recombination-induced variants of Clostridium acetobutylicum ATCC 824 with increased solvent production. <i>Current Microbiology</i> , 1996 , 32, 349-56	2.4	5
39	Genetic manipulation of stationary-phase genes to enhance recombinant protein production in Escherichia coli. <i>Biotechnology and Bioengineering</i> , 1996 , 50, 636-42	4.9	18
38	Molecular characterization of adiY, a regulatory gene which affects expression of the biodegradative acid-induced arginine decarboxylase gene (adiA) of Escherichia coli. <i>Microbiology (United Kingdom)</i> , 1996 , 142 (Pt 5), 1311-1320	2.9	62
37	The central metabolic pathway from acetyl-CoA to butyryl-CoA inClostridium acetobutylicum. <i>FEMS Microbiology Reviews</i> , 1995 , 17, 241-249	15.1	46
36	Sequence and arrangement of genes encoding sigma factors in Clostridium acetobutylicum ATCC 824. <i>Gene</i> , 1995 , 153, 89-92	3.8	20
35	Characterization of an acetyl-CoA C-acetyltransferase (thiolase) gene from Clostridium acetobutylicum ATCC 824. <i>Gene</i> , 1995 , 154, 81-5	3.8	37
34	Metabolic engineering of Escherichia coli to enhance recombinant protein production through acetate reduction. <i>Biotechnology Progress</i> , 1995 , 11, 475-8	2.8	58
33	Characterization of a pH-inducible promoter system for high-level expression of recombinant proteins in Escherichia coli. <i>Biotechnology and Bioengineering</i> , 1995 , 47, 186-92	4.9	26
32	Modification of central metabolic pathway in escherichia coli to reduce acetate accumulation by heterologous expression of the bacillus subtilis acetolactate synthase gene. <i>Biotechnology and Bioengineering</i> , 1994 , 44, 944-51	4.9	75

31	Effect of modified glucose uptake using genetic engineering techniques on high-level recombinant protein production in escherichia coli dense cultures. <i>Biotechnology and Bioengineering</i> , 1994 , 44, 952-6	o ^{4.9}	84
30	Effect of modulated glucose uptake on high-level recombinant protein production in a dense Escherichia coli culture. <i>Biotechnology Progress</i> , 1994 , 10, 644-7	2.8	40
29	Genetic and metabolic engineering of Clostridium acetobutylicum ATCC 824. <i>Annals of the New York Academy of Sciences</i> , 1994 , 721, 54-68	6.5	22
28	Intracellular Concentrations of Coenzyme A and Its Derivatives from Clostridium acetobutylicum ATCC 824 and Their Roles in Enzyme Regulation. <i>Applied and Environmental Microbiology</i> , 1994 , 60, 39-4	14 ^{1.8}	48
27	Sequence and arrangement of two genes of the butyrate-synthesis pathway of Clostridium acetobutylicum ATCC 824. <i>Gene</i> , 1993 , 134, 107-11	3.8	57
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25	Metabolic engineering of Clostridium acetobutylicum ATCC 824 for increased solvent production by enhancement of acetone formation enzyme activities using a synthetic acetone operon. <i>Biotechnology and Bioengineering</i> , 1993 , 42, 1053-60	4.9	88
24	Isolation of mutants ofClostridium acetobutylicum ATCC 824 deficient in protease activity. <i>Current Microbiology</i> , 1993 , 26, 151-154	2.4	8
23	Expression of cloned homologous fermentative genes in Clostridium acetobutylicum ATCC 824. <i>Nature Biotechnology</i> , 1992 , 10, 190-5	44.5	165
22	Vector construction, transformation, and gene amplification in Clostridium acetobutylicum ATCC 824. <i>Annals of the New York Academy of Sciences</i> , 1992 , 665, 39-51	6.5	28
21	Construction of Escherichia coli-Clostridium acetobutylicum shuttle vectors and transformation of Clostridium acetobutylicum strains. <i>Biotechnology Letters</i> , 1992 , 14, 427-432	3	27
20	Effects of rifampicin and chloramphenicol on product and enzyme levels of the acid- and solvent-producing pathways of Clostridium acetobutylicum (ATCC 824). <i>Enzyme and Microbial Technology</i> , 1992 , 14, 277-283	3.8	10
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15	Isolation and Characterization of Mutants of Clostridium acetobutylicum ATCC 824 Deficient in Acetoacetyl-Coenzyme A:Acetate/Butyrate:Coenzyme A-Transferase (EC 2.8.3.9) and in Other Solvent Pathway Enzymes. <i>Applied and Environmental Microbiology</i> , 1989 , 55, 970-6	4.8	76
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