

# Brian F Pflieger

## List of Publications by Year in descending order

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92  
papers

5,865  
citations

87886

38  
h-index

79691

73  
g-index

101  
all docs

101  
docs citations

101  
times ranked

6108  
citing authors

#	ARTICLE	IF	CITATIONS
1	Nonenzymatic Sugar Production from Biomass Using Biomass-Derived $\hat{\text{I}}^3$ -Valerolactone. <i>Science</i> , 2014, 343, 277-280.	12.6	607
2	Combinatorial engineering of intergenic regions in operons tunes expression of multiple genes. <i>Nature Biotechnology</i> , 2006, 24, 1027-1032.	17.5	492
3	Common principles and best practices for engineering microbiomes. <i>Nature Reviews Microbiology</i> , 2019, 17, 725-741.	28.6	324
4	A process for microbial hydrocarbon synthesis: Overproduction of fatty acids in <i>Escherichia coli</i> and catalytic conversion to alkanes. <i>Biotechnology and Bioengineering</i> , 2010, 106, 193-202.	3.3	223
5	Impact of synthetic biology and metabolic engineering on industrial production of fine chemicals. <i>Biotechnology Advances</i> , 2015, 33, 1395-1402.	11.7	195
6	Synthetic Biology Toolbox for Controlling Gene Expression in the Cyanobacterium <i>Synechococcus</i> sp. strain PCC 7002. <i>ACS Synthetic Biology</i> , 2015, 4, 595-603.	3.8	176
7	Engineering <i>Escherichia coli</i> to synthesize free fatty acids. <i>Trends in Biotechnology</i> , 2012, 30, 659-667.	9.3	174
8	Microbial production of fatty acid-derived fuels and chemicals. <i>Current Opinion in Biotechnology</i> , 2013, 24, 1044-1053.	6.6	174
9	Application of Functional Genomics to Pathway Optimization for Increased Isoprenoid Production. <i>Applied and Environmental Microbiology</i> , 2008, 74, 3229-3241.	3.1	171
10	Modular Synthase-Encoding Gene Involved in $\hat{\text{I}}^{\pm}$ -Olefin Biosynthesis in <i>Synechococcus</i> sp. Strain PCC 7002. <i>Applied and Environmental Microbiology</i> , 2011, 77, 4264-4267.	3.1	170
11	CRISPR interference as a titratable, trans-acting regulatory tool for metabolic engineering in the cyanobacterium <i>Synechococcus</i> sp. strain PCC 7002. <i>Metabolic Engineering</i> , 2016, 38, 170-179.	7.0	160
12	Metabolic engineering strategies for microbial synthesis of oleochemicals. <i>Metabolic Engineering</i> , 2015, 29, 1-11.	7.0	152
13	Membrane Stresses Induced by Overproduction of Free Fatty Acids in <i>Escherichia coli</i> . <i>Applied and Environmental Microbiology</i> , 2011, 77, 8114-8128.	3.1	135
14	Biosynthetic Analysis of the Petrobactin Siderophore Pathway from <i>Bacillus anthracis</i> . <i>Journal of Bacteriology</i> , 2007, 189, 1698-1710.	2.2	133
15	Identification of Transport Proteins Involved in Free Fatty Acid Efflux in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2013, 195, 135-144.	2.2	116
16	Genetic tools for reliable gene expression and recombineering in <i>Pseudomonas putida</i> . <i>Journal of Industrial Microbiology and Biotechnology</i> , 2018, 45, 517-527.	3.0	108
17	Production of medium chain length fatty alcohols from glucose in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2013, 20, 177-186.	7.0	98
18	Directed Evolution of AraC for Improved Compatibility of Arabinose- and Lactose-Inducible Promoters. <i>Applied and Environmental Microbiology</i> , 2007, 73, 5711-5715.	3.1	97

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19	A metabolic pathway for catabolizing levulinic acid in bacteria. <i>Nature Microbiology</i> , 2017, 2, 1624-1634.	13.3	86
20	Efflux systems in bacteria and their metabolic engineering applications. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 9381-9393.	3.6	85
21	Characterization and Analysis of Early Enzymes for Petrobactin Biosynthesis in <i>Bacillus anthracis</i> . <i>Biochemistry</i> , 2007, 46, 4147-4157.	2.5	82
22	Microbial sensors for small molecules: Development of a mevalonate biosensor. <i>Metabolic Engineering</i> , 2007, 9, 30-38.	7.0	80
23	Revisiting metabolic engineering strategies for microbial synthesis of oleochemicals. <i>Metabolic Engineering</i> , 2020, 58, 35-46.	7.0	80
24	Computational Redesign of Acyl-ACP Thioesterase with Improved Selectivity toward Medium-Chain-Length Fatty Acids. <i>ACS Catalysis</i> , 2017, 7, 3837-3849.	11.2	77
25	Construction of new synthetic biology tools for the control of gene expression in the cyanobacterium <i>Synechococcus</i> sp. strain PCC 7002. <i>Biotechnology and Bioengineering</i> , 2016, 113, 424-432.	3.3	73
26	Transcription control engineering and applications in synthetic biology. <i>Synthetic and Systems Biotechnology</i> , 2017, 2, 176-191.	3.7	70
27	Modulating Membrane Composition Alters Free Fatty Acid Tolerance in <i>Escherichia coli</i> . <i>PLoS ONE</i> , 2013, 8, e54031.	2.5	68
28	An Organic Acid Based Counter Selection System for Cyanobacteria. <i>PLoS ONE</i> , 2013, 8, e76594.	2.5	62
29	Engineering <i>Escherichia coli</i> for production of C <sub>12</sub> -C <sub>14</sub> polyhydroxyalkanoate from glucose. <i>Metabolic Engineering</i> , 2012, 14, 705-713.	7.0	61
30	Highly Active C <sub>8</sub> -Acyl-ACP Thioesterase Variant Isolated by a Synthetic Selection Strategy. <i>ACS Synthetic Biology</i> , 2018, 7, 2205-2215.	3.8	60
31	Structural and functional analysis of AsbF: Origin of the stealth 3,4-dihydroxybenzoic acid subunit for petrobactin biosynthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 17133-17138.	7.1	58
32	RNA Sequencing Identifies New RNase III Cleavage Sites in <i>Escherichia coli</i> and Reveals Increased Regulation of mRNA. <i>MBio</i> , 2017, 8, .	4.1	56
33	Reassessing <i>Escherichia coli</i> as a cell factory for biofuel production. <i>Current Opinion in Biotechnology</i> , 2017, 45, 92-103.	6.6	53
34	Anaerobic production of medium-chain fatty alcohols via a $\hat{I}^2$ -reduction pathway. <i>Metabolic Engineering</i> , 2018, 48, 63-71.	7.0	53
35	Freshwater diatoms as a source of lipids for biofuels. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2012, 39, 419-428.	3.0	51
36	Machine learning-guided acyl-ACP reductase engineering for improved in vivo fatty alcohol production. <i>Nature Communications</i> , 2021, 12, 5825.	12.8	50

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37	Synthetic biology strategies for synthesizing polyhydroxyalkanoates from unrelated carbon sources. <i>Chemical Engineering Science</i> , 2013, 103, 58-67.	3.8	48
38	A roadmap for the synthesis of separation networks for the recovery of bio-based chemicals: Matching biological and process feasibility. <i>Biotechnology Advances</i> , 2016, 34, 1362-1383.	11.7	43
39	Light-optimized growth of cyanobacterial cultures: Growth phases and productivity of biomass and secreted molecules in light-limited batch growth. <i>Metabolic Engineering</i> , 2018, 47, 230-242.	7.0	43
40	Artificial repressors for controlling gene expression in bacteria. <i>Chemical Communications</i> , 2013, 49, 4325-4327.	4.1	42
41	Engineering photosynthetic production of L-lysine. <i>Metabolic Engineering</i> , 2017, 44, 273-283.	7.0	36
42	Kinetic modeling of free fatty acid production in <i>Escherichia coli</i> based on continuous cultivation of a plasmid free strain. <i>Biotechnology and Bioengineering</i> , 2012, 109, 1518-1527.	3.3	34
43	Bacterial production of free fatty acids from freshwater macroalgal cellulose. <i>Applied Microbiology and Biotechnology</i> , 2011, 91, 435-446.	3.6	31
44	Application of TALEs, CRISPR/Cas and sRNAs as trans-acting regulators in prokaryotes. <i>Current Opinion in Biotechnology</i> , 2014, 29, 46-54.	6.6	31
45	Flux balance analysis indicates that methane is the lowest cost feedstock for microbial cell factories. <i>Metabolic Engineering Communications</i> , 2017, 5, 26-33.	3.6	31
46	Functional and Structural Analysis of the Siderophore Synthetase AsbB through Reconstitution of the Petrobactin Biosynthetic Pathway from <i>Bacillus anthracis</i> . <i>Journal of Biological Chemistry</i> , 2012, 287, 16058-16072.	3.4	30
47	Solvent-Enabled Nonenzymatic Sugar Production from Biomass for Chemical and Biological Upgrading. <i>ChemSusChem</i> , 2015, 8, 1317-1322.	6.8	30
48	A transcription activator-like effector (TALE) induction system mediated by proteolysis. <i>Nature Chemical Biology</i> , 2016, 12, 254-260.	8.0	30
49	A translation-coupling DNA cassette for monitoring protein translation in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2012, 14, 298-305.	7.0	28
50	Optimization of DsRed production in <i>Escherichia coli</i> : Effect of ribosome binding site sequestration on translation efficiency. <i>Biotechnology and Bioengineering</i> , 2005, 92, 553-558.	3.3	27
51	Free fatty acid production in <i>Escherichia coli</i> under phosphate-limited conditions. <i>Applied Microbiology and Biotechnology</i> , 2013, 97, 5149-5159.	3.6	26
52	High-CO <sub>2</sub> Requirement as a Mechanism for the Containment of Genetically Modified Cyanobacteria. <i>ACS Synthetic Biology</i> , 2018, 7, 384-391.	3.8	26
53	Regulatory Tools for Controlling Gene Expression in Cyanobacteria. <i>Advances in Experimental Medicine and Biology</i> , 2018, 1080, 281-315.	1.6	26
54	Rewiring yeast metabolism to synthesize products beyond ethanol. <i>Current Opinion in Chemical Biology</i> , 2020, 59, 182-192.	6.1	25

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55	Metabolic engineering of $\beta^2$ -oxidation to leverage thioesterases for production of 2-heptanone, 2-nonanone and 2-undecanone. <i>Metabolic Engineering</i> , 2020, 61, 335-343.	7.0	24
56	Genetic and genomic analysis of RNases in model cyanobacteria. <i>Photosynthesis Research</i> , 2015, 126, 171-183.	2.9	23
57	Production of 1-octanol in <i>Escherichia coli</i> by a high flux thioesterase route. <i>Metabolic Engineering</i> , 2020, 61, 352-359.	7.0	22
58	A Desaturase Gene Involved in the Formation of 1,14-Nonadecadiene in <i>Synechococcus</i> sp. Strain PCC 7002. <i>Applied and Environmental Microbiology</i> , 2014, 80, 6073-6079.	3.1	18
59	Insights into the industrial growth of cyanobacteria from a model of the carbon concentrating mechanism. <i>AIChE Journal</i> , 2014, 60, 1269-1277.	3.6	18
60	Isolation of improved free fatty acid overproducing strains of <i>Escherichia coli</i> via Nile red based high-throughput screening. <i>Environmental Progress and Sustainable Energy</i> , 2012, 31, 17-23.	2.3	16
61	Directed Evolution Reveals the Functional Sequence Space of an Adenylation Domain Specificity Code. <i>ACS Chemical Biology</i> , 2019, 14, 2044-2054.	3.4	16
62	Growth-coupled bioconversion of levulinic acid to butanone. <i>Metabolic Engineering</i> , 2019, 55, 92-101.	7.0	16
63	Stepwise genetic engineering of <i>Pseudomonas putida</i> enables robust heterologous production of prodigiosin and glidobactin A. <i>Metabolic Engineering</i> , 2021, 67, 112-124.	7.0	16
64	Introduction of NADH-dependent nitrate assimilation in <i>Synechococcus</i> sp. PCC 7002 improves photosynthetic production of 2-methyl-1-butanol and isobutanol. <i>Metabolic Engineering</i> , 2022, 69, 87-97.	7.0	14
65	Inhibition of Cyanobacterial Growth on a Municipal Wastewater Sidestream Is Impacted by Temperature. <i>MSphere</i> , 2018, 3, .	2.9	13
66	Leveraging synthetic biology for producing bioactive polyketides and non-ribosomal peptides in bacterial heterologous hosts. <i>MedChemComm</i> , 2019, 10, 668-681.	3.4	13
67	Model-driven analysis of mutant fitness experiments improves genome-scale metabolic models of <i>Zymomonas mobilis</i> ZM4. <i>PLoS Computational Biology</i> , 2020, 16, e1008137.	3.2	12
68	Microbes paired for biological gas-to-liquids (Bio-GTL) process. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 3717-3719.	7.1	11
69	Biological synthesis unbounded?. <i>Nature Biotechnology</i> , 2015, 33, 1148-1149.	17.5	10
70	Enabling commercial success of industrial biotechnology. <i>Science</i> , 2021, 374, 1563-1565.	12.6	10
71	Genome sequence and analysis of <i>Escherichia coli</i> production strain LS5218. <i>Metabolic Engineering Communications</i> , 2017, 5, 78-83.	3.6	9
72	Distinct and redundant functions of three homologs of RNase III in the cyanobacterium <i>Synechococcus</i> sp. strain PCC 7002. <i>Nucleic Acids Research</i> , 2018, 46, 1984-1997.	14.5	9

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73	Enhancing Photosynthetic Production of Glycogen-Rich Biomass for Use as a Fermentation Feedstock. <i>Frontiers in Energy Research</i> , 2020, 8, .	2.3	9
74	Renewable linear alpha-olefins by base-catalyzed dehydration of biologically-derived fatty alcohols. <i>Green Chemistry</i> , 2021, 23, 4338-4354.	9.0	9
75	IPro+: Computational Protein Design Tool Allowing for Insertions and Deletions. <i>Structure</i> , 2020, 28, 1344-1357.e4.	3.3	8
76	Optimization of a T7-RNA polymerase system in <i>Synechococcus</i> sp. PCC 7002 mirrors the protein overproduction phenotype from <i>E. coli</i> BL21(DE3). <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 1147-1158.	3.6	8
77	Accelerating strain phenotyping with desorption electrospray ionization-imaging mass spectrometry and untargeted analysis of intact microbial colonies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	8
78	Metabolic engineering strategies to produce medium-chain oleochemicals via acyl-ACP:CoA transacylase activity. <i>Nature Communications</i> , 2022, 13, 1619.	12.8	8
79	EnZymClass: Substrate specificity prediction tool of plant acyl-ACP thioesterases based on ensemble learning. <i>Current Research in Biotechnology</i> , 2022, 4, 1-9.	3.7	7
80	Genome-Wide Analysis of RNA Decay in the Cyanobacterium <i>Synechococcus</i> sp. Strain PCC 7002. <i>MSystems</i> , 2020, 5, .	3.8	6
81	Optimization of Synthetic Operons Using Libraries of Post-Transcriptional Regulatory Elements. <i>Methods in Molecular Biology</i> , 2011, 765, 99-111.	0.9	5
82	Functional genomics analysis of free fatty acid production under continuous phosphate limiting conditions. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2017, 44, 759-772.	3.0	5
83	Infrastructures for Phosphorus Recovery from Livestock Waste Using Cyanobacteria: Transportation, Techno-Economic, and Policy Implications. <i>ACS Sustainable Chemistry and Engineering</i> , 2021, 9, 11416-11426.	6.7	4
84	Structural and Biosynthetic Analysis of the Fabrubactins, Unusual Siderophores from <i>Agrobacterium fabrum</i> Strain C58. <i>ACS Chemical Biology</i> , 2021, 16, 125-135.	3.4	4
85	Comparative functional genomics identifies an iron-limited bottleneck in a <i>Saccharomyces cerevisiae</i> strain with a cytosolic-localized isobutanol pathway. <i>Synthetic and Systems Biotechnology</i> , 2022, 7, 738-749.	3.7	4
86	Production of Fatty Acids and Derivatives by Metabolic Engineering of Bacteria. , 2016, , 1-24.		2
87	Bypassing the refinery for production of high-value BTX derivatives. <i>Biotechnology Journal</i> , 2013, 8, 1375-1376.	3.5	0
88	Editorial: Biochemical and molecular engineering. <i>Biotechnology Journal</i> , 2014, 9, 587-588.	3.5	0
89	Editorial overview: Energy biotechnology. <i>Current Opinion in Biotechnology</i> , 2017, 45, v-viii.	6.6	0
90	Production of Fatty Acids and Derivatives by Metabolic Engineering of Bacteria. , 2017, , 1-24.		0

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91	Production of Fatty Acids and Derivatives by Metabolic Engineering of Bacteria. , 2017, , 435-458.		0
92	Directed Evolution of an Adenylation Domain Specificity Code. FASEB Journal, 2018, 32, 530.6.	0.5	0