CITATION REPORT List of articles citing

A comparative view of metabolite and substrate stress and tolerance in microbial bioprocessing: From biofuels and chemicals, to biocatalysis and bioremediation

DOI: 10.1016/j.ymben.2010.03.004 Metabolic Engineering, 2010, 12, 307-31.

Source: https://exaly.com/paper-pdf/48343147/citation-report.pdf

Version: 2024-04-28

This report has been generated based on the citations recorded by exaly.com for the above article. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

#	Paper	IF	Citations
457	Microbial degradation of tetrachloromethane: mechanisms and perspectives for bioremediation. 2010 , 74, 257-75		38
456	Transcriptional analysis of Lactobacillus brevis to N-butanol and ferulic acid stress responses. <i>PLoS ONE</i> , 2011 , 6, e21438	3.7	42
455	Switching Clostridium acetobutylicum to an ethanol producer by disruption of the butyrate/butanol fermentative pathway. <i>Metabolic Engineering</i> , 2011 , 13, 464-73	9.7	66
454	Styrene biosynthesis from glucose by engineered E. coli. <i>Metabolic Engineering</i> , 2011 , 13, 544-54	9.7	192
453	A stable yeast strain efficiently producing cholesterol instead of ergosterol is functional for tryptophan uptake, but not weak organic acid resistance. <i>Metabolic Engineering</i> , 2011 , 13, 555-69	9.7	72
452	Metabolic engineering of Thermobifida fusca for direct aerobic bioconversion of untreated lignocellulosic biomass to 1-propanol. <i>Metabolic Engineering</i> , 2011 , 13, 570-7	9.7	66
451	Engineering topology and kinetics of sucrose metabolism in Saccharomyces cerevisiae for improved ethanol yield. <i>Metabolic Engineering</i> , 2011 , 13, 694-703	9.7	74
450	Metabolic engineering of Clostridium acetobutylicum: recent advances to improve butanol production. 2011 , 22, 634-47		293
449	Dehalogenase gene detection and microbial diversity of a chlorinated hydrocarbon contaminated site. <i>World Journal of Microbiology and Biotechnology</i> , 2011 , 27, 2407-2414	4.4	6
448	Fermentative production of butanolthe industrial perspective. 2011 , 22, 337-43		560
447	Evolution combined with genomic study elucidates genetic bases of isobutanol tolerance in Escherichia coli. <i>Microbial Cell Factories</i> , 2011 , 10, 18	6.4	139
446	Engineering microbes for tolerance to next-generation biofuels. 2011 , 4, 32		216
445	Elucidating acetate tolerance in E. coli using a genome-wide approach. <i>Metabolic Engineering</i> , 2011 , 13, 214-24	9.7	50
444	L-ascorbic acid producing yeasts learn from plants how to recycle it. <i>Metabolic Engineering</i> , 2011 , 13, 177-85	9.7	16
443	Photosynthesis driven conversion of carbon dioxide to fatty alcohols and hydrocarbons in cyanobacteria. <i>Metabolic Engineering</i> , 2011 , 13, 169-76	9.7	184
442	Computation of metabolic fluxes and efficiencies for biological carbon dioxide fixation. <i>Metabolic Engineering</i> , 2011 , 13, 150-8	9.7	52
441	Engineering the robustness of Clostridium acetobutylicum by introducing glutathione biosynthetic capability. <i>Metabolic Engineering</i> , 2011 , 13, 426-34	9.7	64

440	Linking genes to microbial growth kinetics: an integrated biochemical systems engineering approach. <i>Metabolic Engineering</i> , 2011 , 13, 401-13	9.7	22	
439	Bioconversion of d-xylose to d-xylonate with Kluyveromyces lactis. <i>Metabolic Engineering</i> , 2011 , 13, 383	- 9 . 1 7	248	
438	Metabolic engineering of Clostridium tyrobutyricum for n-butanol production. <i>Metabolic Engineering</i> , 2011 , 13, 373-82	9.7	177	
437	Metabolic engineering of cyanobacteria for 1-butanol production from carbon dioxide. <i>Metabolic Engineering</i> , 2011 , 13, 353-63	9.7	314	
436	Coexisting/Coexpressing Genomic Libraries (CoGeL) identify interactions among distantly located genetic loci for developing complex microbial phenotypes. 2011 , 39, e152		41	
435	From Pathways to Genomes and Beyond: The Metabolic Engineering Toolbox and Its Place in Biofuels Production. 2011 , 1,		3	
434	Laboratory-evolved mutants of an exogenous global regulator, IrrE from Deinococcus radiodurans, enhance stress tolerances of Escherichia coli. <i>PLoS ONE</i> , 2011 , 6, e16228	3.7	55	
433	Membrane stresses induced by overproduction of free fatty acids in Escherichia coli. 2011 , 77, 8114-28		109	
432	Global transcriptome response to ionic liquid by a tropical rain forest soil bacterium, Enterobacter lignolyticus. 2012 , 109, E2173-82		81	
431	Significant rewiring of the transcriptome and proteome of an Escherichia coli strain harboring a tailored exogenous global regulator IrrE. <i>PLoS ONE</i> , 2012 , 7, e37126	3.7	18	
430	Characterization of acetonitrile-tolerant marine bacterium Exiguobacterium sp. SBH81 and its tolerance mechanism. 2012 , 27, 30-5		8	
429	Effects of nutritional enrichment on the production of acetone-butanol-ethanol (ABE) by Clostridium acetobutylicum. 2012 , 50, 1063-6		17	
428	Improving butanol fermentation to enter the advanced biofuel market. 2012, 3,		33	
427	Over-expression of stress protein-encoding genes helps Clostridium acetobutylicum to rapidly adapt to butanol stress. 2012 , 34, 1643-9		50	
426	Synthetic Escherichia coli consortia engineered for syntrophy demonstrate enhanced biomass productivity. 2012 , 157, 159-66		96	
425	A transcriptional study of acidogenic chemostat cells of Clostridium acetobutylicumcellular behavior in adaptation to n-butanol. 2012 , 161, 366-77		41	
424	Multi-copy genes that enhance the yield of mammalian G protein-coupled receptors in Escherichia coli. <i>Metabolic Engineering</i> , 2012 , 14, 591-602	9.7	22	
423	Visualizing evolution in real time to determine the molecular mechanisms of n-butanol tolerance in Escherichia coli. <i>Metabolic Engineering</i> , 2012 , 14, 579-90	9.7	91	

422	Metabolic engineering of D-xylose pathway in Clostridium beijerinckii to optimize solvent production from xylose mother liquid. <i>Metabolic Engineering</i> , 2012 , 14, 569-78	9.7	86
421	A selection platform for carbon chain elongation using the CoA-dependent pathway to produce linear higher alcohols. <i>Metabolic Engineering</i> , 2012 , 14, 504-11	9.7	112
420	ATP drives direct photosynthetic production of 1-butanol in cyanobacteria. 2012, 109, 6018-23		293
419	Quantitative iTRAQ LC-MS/MS proteomics reveals metabolic responses to biofuel ethanol in cyanobacterial Synechocystis sp. PCC 6803. 2012 , 11, 5286-300		114
418	Development of microorganisms for cellulose-biofuel consolidated bioprocessings: metabolic engineers' tricks. 2012 , 3, e201210007		33
417	Physiological predisposition of various Clostridium species to synthetize 1,3-propanediol from glycerol. 2012 , 47, 1308-1319		51
416	Engineering the robustness of industrial microbes through synthetic biology. 2012 , 20, 94-101		57
415	Enzymatic dynamics of microbial acid tolerance response (ATR) during the enhanced biohydrogen production process via anaerobic digestion. 2012 , 37, 10655-10662		9
414	Exploring the combinatorial genomic space in Escherichia coli for ethanol tolerance. 2012 , 7, 1337-45		26
413	Proteomic analysis reveals resistance mechanism against biofuel hexane in Synechocystis sp. PCC 6803. 2012 , 5, 68		67
412	Towards a Synthetic Biology of the Stress-Response and the Tolerance Phenotype: Systems Understanding and Engineering of the Clostridium acetobutylicum Stress-Response and Tolerance to Toxic Metabolites. 2012 , 193-219		1
411	Metabolic uncouplers in environmental research: a critical review. 2012 , 28,		6
410	Bioprocessing for biofuels. 2012 , 23, 390-5		65
409	Clostridia: the importance of their exceptional substrate and metabolite diversity for biofuel and biorefinery applications. 2012 , 23, 364-81		310
408	Escherichia coli for biofuel production: bridging the gap from promise to practice. 2012 , 30, 538-45		69
407	Physiological heterogeneity of Pseudomonas taetrolens during lactobionic acid production. 2012 , 96, 1465-77		24
406	Stoichiometric and energetic analyses of non-photosynthetic CO2-fixation pathways to support synthetic biology strategies for production of fuels and chemicals. 2012 , 1, 380-395		165
405	Toward a semisynthetic stress response system to engineer microbial solvent tolerance. 2012 , 3,		67

404	Metabolic engineering of biocatalysts for carboxylic acids production. 2012 , 3, e201210011		26
403	RNA-seq based identification and mutant validation of gene targets related to ethanol resistance in cyanobacterial Synechocystis sp. PCC 6803. 2012 , 5, 89		62
402	Enhancing E. coli tolerance towards oxidative stress via engineering its global regulator cAMP receptor protein (CRP). <i>PLoS ONE</i> , 2012 , 7, e51179	3.7	38
401	Application of meta-transcriptomics and -proteomics to analysis of in situ physiological state. <i>Frontiers in Microbiology</i> , 2012 , 3, 184	5.7	16
400	Synthetic feedback loop model for increasing microbial biofuel production using a biosensor. <i>Frontiers in Microbiology</i> , 2012 , 3, 360	5.7	39
399	Butanol production from renewable biomass: rediscovery of metabolic pathways and metabolic engineering. 2012 , 7, 186-98		116
398	High-titer n-butanol production by clostridium acetobutylicum JB200 in fed-batch fermentation with intermittent gas stripping. 2012 , 109, 2746-56		176
397	Improvements in Biobutanol Fermentation and Their Impacts on Distillation Energy Consumption and Wastewater Generation. 2012 , 5, 504-514		57
396	From network models to network responses: integration of thermodynamic and kinetic properties of yeast genome-scale metabolic networks. 2012 , 12, 129-43		60
395	Application of phenotypic microarrays to environmental microbiology. 2012 , 23, 41-8		33
395 394	Application of phenotypic microarrays to environmental microbiology. 2012, 23, 41-8 Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as an expression vector. 2012, 113, 596-603		33
	Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as	9.7	
394	Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as an expression vector. 2012 , 113, 596-603 Engineering of vesicle trafficking improves heterologous protein secretion in Saccharomyces	9.7	6
394	Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as an expression vector. 2012 , 113, 596-603 Engineering of vesicle trafficking improves heterologous protein secretion in Saccharomyces cerevisiae. <i>Metabolic Engineering</i> , 2012 , 14, 120-7	9.7	6
394 393 392	Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as an expression vector. 2012 , 113, 596-603 Engineering of vesicle trafficking improves heterologous protein secretion in Saccharomyces cerevisiae. <i>Metabolic Engineering</i> , 2012 , 14, 120-7 Construction of microbial cell factories for industrial bioprocesses. 2012 , 87, 445-450 Integrated OMICS guided engineering of biofuel butanol-tolerance in photosynthetic Synechocystis	9-7	6 81 28
394 393 392 391	Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as an expression vector. 2012 , 113, 596-603 Engineering of vesicle trafficking improves heterologous protein secretion in Saccharomyces cerevisiae. <i>Metabolic Engineering</i> , 2012 , 14, 120-7 Construction of microbial cell factories for industrial bioprocesses. 2012 , 87, 445-450 Integrated OMICS guided engineering of biofuel butanol-tolerance in photosynthetic Synechocystis sp. PCC 6803. 2013 , 6, 106	9-7	6 81 28
394 393 392 391 390	Identification of a methanol-inducible promoter from Rhodococcus erythropolis PR4 and its use as an expression vector. 2012, 113, 596-603 Engineering of vesicle trafficking improves heterologous protein secretion in Saccharomyces cerevisiae. <i>Metabolic Engineering</i> , 2012, 14, 120-7 Construction of microbial cell factories for industrial bioprocesses. 2012, 87, 445-450 Integrated OMICS guided engineering of biofuel butanol-tolerance in photosynthetic Synechocystis sp. PCC 6803. 2013, 6, 106 RecA-mediated SOS response provides a geraniol tolerance in Escherichia coli. 2013, 167, 357-64 Cell morphology variations of Klebsiella pneumoniae induced by acetate stress using biomimetic	9-7	6 81 28 62

386	Overexpression of fetA (ybbL) and fetB (ybbM), Encoding an Iron Exporter, Enhances Resistance to Oxidative Stress in Escherichia coli. 2013 , 79, 7210-9	28
385	Comparative proteomic and metabolomic analysis of Staphylococcus warneri SG1 cultured in the presence and absence of butanol. 2013 , 12, 4478-89	26
384	Global metabolomic and network analysis of Escherichia coli responses to exogenous biofuels. 2013 , 12, 5302-12	44
383	Microbial engineering strategies to improve cell viability for biochemical production. <i>Biotechnology Advances</i> , 2013 , 31, 903-14	3 48
382	Dissecting the assays to assess microbial tolerance to toxic chemicals in bioprocessing. 2013 , 31, 643-53	35
381	Engineering a synthetic pathway in cyanobacteria for isopropanol production directly from carbon dioxide and light. <i>Metabolic Engineering</i> , 2013 , 20, 101-8	115
380	Two-stage in situ gas stripping for enhanced butanol fermentation and energy-saving product recovery. 2013 , 135, 396-402	138
379	Engineering of transcriptional regulators enhances microbial stress tolerance. <i>Biotechnology Advances</i> , 2013 , 31, 986-91	3 53
378	Exploring the Capabilities of the Geobiosphere's Microbial Genome. 2013 , 59, 688-698	5
377	Microevolution from shock to adaptation revealed strategies improving ethanol tolerance and production in Thermoanaerobacter. 2013 , 6, 103	11
376	Pulsed addition of HMF and furfural to batch-grown xylose-utilizing Saccharomyces cerevisiae results in different physiological responses in glucose and xylose consumption phase. 2013 , 6, 181	33
375	The Clostridium small RNome that responds to stress: the paradigm and importance of toxic metabolite stress in C. acetobutylicum. 2013 , 14, 849	43
374	Transcription factors and genetic circuits orchestrating the complex, multilayered response of Clostridium acetobutylicum to butanol and butyrate stress. 2013 , 7, 120	59
373	Biological Production of Butanol and Higher Alcohols. 2013 , 235-262	18
372	Using transcriptomics to improve butanol tolerance of Synechocystis sp. strain PCC 6803. 2013 , 79, 7419-27	73
371	Identification of transport proteins involved in free fatty acid efflux in Escherichia coli. 2013 , 195, 135-44	90
370	Identification and engineering of cholesterol oxidases involved in the initial step of sterols catabolism in Mycobacterium neoaurum. <i>Metabolic Engineering</i> , 2013 , 15, 75-87	61
369	Quantitative proteomics reveals dynamic responses of Synechocystis sp. PCC 6803 to next-generation biofuel butanol. 2013 , 78, 326-45	85

(2013-2013)

368	Metabolic engineering of Torulopsis glabrata for malate production. <i>Metabolic Engineering</i> , 2013 , 19, 10-6	9.7	66
367	Engineering improved ethanol production in Escherichia coli with a genome-wide approach. <i>Metabolic Engineering</i> , 2013 , 17, 1-11	9.7	40
366	GroESL overexpression imparts Escherichia coli tolerance to i-, n-, and 2-butanol, 1,2,4-butanetriol and ethanol with complex and unpredictable patterns. <i>Metabolic Engineering</i> , 2013 , 15, 196-205	9.7	92
365	Quantitative trait analysis of yeast biodiversity yields novel gene tools for metabolic engineering. <i>Metabolic Engineering</i> , 2013 , 17, 68-81	9.7	43
364	A modified pathway for the production of acetone in Escherichia coli. <i>Metabolic Engineering</i> , 2013 , 15, 218-25	9.7	21
363	Metabolic engineering of Clostridium acetobutylicum ATCC 824 for the high-yield production of a biofuel composed of an isopropanol/butanol/ethanol mixture. <i>Metabolic Engineering</i> , 2013 , 18, 1-8	9.7	114
362	Activating transhydrogenase and NAD kinase in combination for improving isobutanol production. <i>Metabolic Engineering</i> , 2013 , 16, 1-10	9.7	90
361	Detoxification of Lignocellulose Hydrolysates: Biochemical and Metabolic Engineering Toward White Biotechnology. 2013 , 6, 388-401		144
360	Engineering Clostridium acetobutylicum for alcohol production. 2013 , 166, 25-33		36
359	n-Butanol Production from Acid-Pretreated Jatropha Seed Cake by Clostridium acetobutylicum. 2013 , 6, 991-999		9
358	Synthetic tolerance: three noncoding small RNAs, DsrA, ArcZ and RprA, acting supra-additively against acid stress. 2013 , 41, 8726-37		80
357	Physiological and transcriptional responses of Saccharomyces cerevisiae to d-limonene show changes to the cell wall but not to the plasma membrane. 2013 , 79, 3590-600		62
356	Genetic determinants for n-butanol tolerance in evolved Escherichia coli mutants: cross adaptation and antagonistic pleiotropy between n-butanol and other stressors. 2013 , 79, 5313-20		42
355	Cell-free Biosystems in the Production of Electricity and Bioenergy. 2013 , 137, 125-52		4
354	Selection and optimization of a salting-out extraction system for recovery of biobutanol from fermentation broth. 2013 , 13, 464-471		26
353	Molecular adaptation mechanisms employed by ethanologenic bacteria in response to lignocellulose-derived inhibitory compounds. 2013 , 9, 598-612		76
352	Industrial robustness: understanding the mechanism of tolerance for the Populus hydrolysate-tolerant mutant strain of Clostridium thermocellum. <i>PLoS ONE</i> , 2013 , 8, e78829	3.7	19
351	Comparative proteomics analysis of engineered Saccharomyces cerevisiae with enhanced biofuel precursor production. <i>PLoS ONE</i> , 2013 , 8, e84661	3.7	12

350	Design and development of synthetic microbial platform cells for bioenergy. <i>Frontiers in Microbiology</i> , 2013 , 4, 92	5.7	29
349	Exometabolomics approaches in studying the application of lignocellulosic biomass as fermentation feedstock. 2013 , 3, 119-43		10
348	Regulation Systems of Bacteria such as Escherichia coli in Response to Nutrient Limitation and Environmental Stresses. 2013 , 4, 1-35		126
347	Artificially constructed quorum-sensing circuits are used for subtle control of bacterial population density. <i>PLoS ONE</i> , 2014 , 9, e104578	3.7	10
346	Enhancing stress-resistance for efficient microbial biotransformations by synthetic biology. 2014 , 2, 44		11
345	Protein Network Signatures Associated with Exogenous Biofuels Treatments in Cyanobacterium Synechocystis sp. PCC 6803. 2014 , 2, 48		8
344	. 2014,		8
343	1,3-Propanediol production from crude glycerol by Clostridium butyricum DSP1 in repeated batch. 2014 , 17, 322-328		27
342	Near-real-time analysis of the phenotypic responses of Escherichia coli to 1-butanol exposure using Raman Spectroscopy. 2014 , 196, 3983-91		28
341	Microbiological Metabolism Under Chemical Stress. 2014 , 497-509		1
340	Evaluation of asymmetric polydimethylsiloxane-polyvinylidene fluoride composite membrane and incorporated with acetone-butanol-ethanol fermentation for butanol recovery. 2014 , 188, 158-65		35
339	The effect of ethanol on cell properties and steroid 1-en-dehydrogenation biotransformation of Arthrobacter simplex. <i>Biotechnology and Applied Biochemistry</i> , 2014 , 61, 555-64	2.8	9
338	Metabolomic basis of laboratory evolution of butanol tolerance in photosynthetic Synechocystis sp. PCC 6803. <i>Microbial Cell Factories</i> , 2014 , 13, 151	6.4	50
337	Pseudomonas 2.0: genetic upgrading of P. putida KT2440 as an enhanced host for heterologous gene expression. <i>Microbial Cell Factories</i> , 2014 , 13, 159	6.4	152
336	Quantifying the effects of the division of labor in metabolic pathways. 2014 , 360, 222-242		23
335	Microbial tolerance engineering toward biochemical production: from lignocellulose to products. 2014 , 29, 99-106		70
334	Robustness of Pseudomonas putida KT2440 as a host for ethanol biosynthesis. 2014 , 31, 562-71		48
333	Overexpression of sigma factor SigB improves temperature and butanol tolerance of Synechocystis sp. PCC6803. 2014 , 182-183, 54-60		50

332	Cloning, expression, purification, crystallization and X-ray crystallographic analysis of (S)-3-hydroxybutyryl-CoA dehydrogenase from Clostridium butyricum. 2014 , 70, 485-8		2	
331	Engineering biofuel tolerance in non-native producing microorganisms. <i>Biotechnology Advances</i> , 2014 , 32, 541-8	17.8	62	
330	Two non-exclusive strategies employed to protect Torulopsis glabrata against hyperosmotic stress. 2014 , 98, 3099-110		3	
329	Comparison of extraction phases for a two-phase culture of a recombinant E. coli producing retinoids. 2014 , 36, 497-505		4	
328	Inhibitory effects of substrate and soluble end products on biohydrogen production of the alkalithermophile Caloramator celer: Kinetic, metabolic and transcription analyses. 2014 , 39, 6391-6401		35	
327	Proteomic analyses of the phase transition from acidogenesis to solventogenesis using solventogenic and non-solventogenic Clostridium acetobutylicum strains. 2014 , 98, 5105-15		26	
326	Characterization and engineering of 3-ketosteroid-?1-dehydrogenase and 3-ketosteroid-9Hydroxylase in Mycobacterium neoaurum ATCC 25795 to produce 9Hydroxy-4-androstene-3,17-dione through the catabolism of sterols. <i>Metabolic Engineering</i> , 2014 , 24, 181-91	9.7	76	
325	Application of new metabolic engineering tools for Clostridium acetobutylicum. 2014 , 98, 5823-37		62	
324	Engineering stress tolerance of Escherichia coli by stress-induced mutagenesis (SIM)-based adaptive evolution. 2014 , 9, 120-7		18	
323	Characterization of gas stripping and its integration with acetoneButanolBthanol fermentation for high-efficient butanol production and recovery. 2014 , 83, 55-61		71	
322	High-yield production of manganese peroxidase, lignin peroxidase, and versatile peroxidase in Phanerochaete chrysosporium. 2014 , 98, 9283-94		42	
321	Microbial inhibitors: formation and effects on acetone-butanol-ethanol fermentation of lignocellulosic biomass. 2014 , 98, 9151-72		97	
320	Overexpression of the Lactobacillus plantarum peptidoglycan biosynthesis murA2 gene increases the tolerance of Escherichia coli to alcohols and enhances ethanol production. 2014 , 98, 8399-411		19	
319	Exploring the heterologous genomic space for building, stepwise, complex, multicomponent tolerance to toxic chemicals. 2014 , 3, 476-86		13	
318	Biofuel toxicity and mechanisms of biofuel tolerance in three model cyanobacteria. 2014 , 5, 121-132		23	
317	Towards lactic acid bacteria-based biorefineries. <i>Biotechnology Advances</i> , 2014 , 32, 1216-1236	17.8	124	
316	Assessment of bacterial resistance to organic solvents using a combined confocal laser scanning and atomic force microscopy (CLSM/AFM). 2014 , 107, 23-9		25	
315	Transcriptomic analysis of Clostridium thermocellum Populus hydrolysate-tolerant mutant strain shows increased cellular efficiency in response to Populus hydrolysate compared to the wild type strain. 2014 , 14, 215		8	

314	Engineering clostridia for butanol production from biorenewable resources: from cells to process integration. 2014 , 6, 43-54		57
313	Enhancing Corynebacterium glutamicum robustness by over-expressing a gene, mshA, for mycothiol glycosyltransferase. 2014 , 36, 1453-9		17
312	Isobutanol. 2014 , 327-352		2
311	Stable high-titer n-butanol production from sucrose and sugarcane juice by Clostridium acetobutylicum JB200 in repeated batch fermentations. 2014 , 163, 172-9		67
310	Cloning, expression, purification, crystallization and X-ray crystallographic analysis of the (S)-3-hydroxybutyryl-CoA dehydrogenase PaaH1 from Ralstonia eutropha H16. 2014 , 70, 955-8		1
309	Scale-up of anaerobic 1,3-propanediol production by Clostridium butyricum DSP1 from crude glycerol. 2014 , 14, 45		32
308	Butanol tolerance regulated by a two-component response regulator Slr1037 in photosynthetic Synechocystis sp. PCC 6803. 2014 , 7, 89		28
307	Structural insights into substrate specificity of crotonase from the n-butanol producing bacterium Clostridium acetobutylicum. 2014 , 451, 431-5		10
306	Co-expression of TAL1 and ADH1 in recombinant xylose-fermenting Saccharomyces cerevisiae improves ethanol production from lignocellulosic hydrolysates in the presence of furfural. 2014 , 117, 165-169		55
305	High temperature stimulates acetic acid accumulation and enhances the growth inhibition and ethanol production by Saccharomyces cerevisiae under fermenting conditions. 2014 , 98, 6085-94		37
304	Bioprocessing Technologies. 2014 , 133-166		1
303	Succinic Acid. 2014 , 435-472		13
302	Dynamic model for isopropanol production by Cupriavidus necator. 2014 , 47, 4388-4393		3
301	Exacerbation of substrate toxicity by IPTG in Escherichia coli BL21(DE3) carrying a synthetic metabolic pathway. <i>Microbial Cell Factories</i> , 2015 , 14, 201	6.4	88
300	Adaptation of the autotrophic acetogen Sporomusa ovata to methanol accelerates the conversion of CO2 to organic products. 2015 , 5, 16168		56
299	Dynamic interplay of multidrug transporters with TolC for isoprenol tolerance in Escherichia coli. 2015 , 5, 16505		15
298	Engineering improved bio-jet fuel tolerance in Escherichia coli using a transgenic library from the hydrocarbon-degrader Marinobacter aquaeolei. 2015 , 8, 165		18
297	Engineering protein folding and translocation improves heterologous protein secretion in Saccharomyces cerevisiae. 2015 , 112, 1872-82		44

296	Bioremediation of Polluted Waters Using Microorganisms. 2015,	43
295	Changes in membrane plasmalogens of Clostridium pasteurianum during butanol fermentation as determined by lipidomic analysis. <i>PLoS ONE</i> , 2015 , 10, e0122058	25
294	Mitochondrial fusion and fission are involved in stress tolerance of Candida glabrata. <i>Bioresources and Bioprocessing</i> , 2015 , 2,	4
293	Mechanisms of solvent resistance mediated by interplay of cellular factors in Pseudomonas putida. 2015 , 39, 555-66	112
292	Untargeted metabolomics analysis revealed changes in the composition of glycerolipids and phospholipids in Bacillus subtilis under 1-butanol stress. 2015 , 99, 5971-83	19
291	Tolerance engineering in bacteria for the production of advanced biofuels and chemicals. 2015 , 23, 498-508	155
29 0	Metabolic engineering of Clostridium tyrobutyricum for n-butanol production from maltose and soluble starch by overexpressing glucosidase. 2015 , 99, 6155-65	19
289	Complex and extensive post-transcriptional regulation revealed by integrative proteomic and transcriptomic analysis of metabolite stress response in Clostridium acetobutylicum. 2015 , 8, 81	27
288	Expression of heterologous sigma factors enables functional screening of metagenomic and heterologous genomic libraries. 2015 , 6, 7045	45
287	Studies on the influence of different metabolic uncouplers on the biodegradation of toluene in a differential biofilter reactor. 2015 , 20, 915-923	5
286	Production of Industrially Relevant Isoprenoid Compounds in Engineered Microbes. 2015, 303-334	14
285	Aromatic catabolic pathway selection for optimal production of pyruvate and lactate from lignin. Metabolic Engineering, 2015 , 28, 240-247 9.7	155
284	Engineering Escherichia coli Cell Factories for n-Butanol Production. 2016 , 155, 141-63	4
283	A novel ionic liquid-tolerant Fusarium oxysporum BN secreting ionic liquid-stable cellulase: consolidated bioprocessing of pretreated lignocellulose containing residual ionic liquid. 2015 , 181, 18-25	43
282	The effect of high concentrations of glycerol on the growth, metabolism and adaptation capacity of Clostridium butyricum DSP1. 2015 , 18, 128-133	29
281	A paradigm shift in biomass technology from complete to partial cellulose hydrolysis: lessons learned from nature. 2015 , 6, 69-72	17
280	Metabolomic analysis reveals functional overlapping of three signal transduction proteins in regulating ethanol tolerance in cyanobacterium Synechocystis sp. PCC 6803. 2015 , 11, 770-82	22
279	Effect of 1,3-propanediol, organic acids, and ethanol on growth and metabolism of Clostridium butyricum DSP1. 2015 , 99, 3179-89	11

278	Thermal and solvent stress cross-tolerance conferred to Corynebacterium glutamicum by adaptive laboratory evolution. 2015 , 81, 2284-98		57
277	Improvement of acetic acid tolerance of Saccharomyces cerevisiae using a zinc-finger-based artificial transcription factor and identification of novel genes involved in acetic acid tolerance. 2015 , 99, 2441-9		41
276	Global transcriptional analysis of Escherichia coli expressing IrrE, a regulator from Deinococcus radiodurans, in response to NaCl shock. 2015 , 11, 1165-71		20
275	Elucidating butanol tolerance mediated by a response regulator Sll0039 in Synechocystis sp. PCC 6803 using a metabolomic approach. 2015 , 99, 1845-57		21
274	Metabolic Regulation and Coordination of the Metabolism in Bacteria in Response to a Variety of Growth Conditions. 2016 , 155, 1-54		23
273	Genome reduction boosts heterologous gene expression in Pseudomonas putida. <i>Microbial Cell Factories</i> , 2015 , 14, 23	6.4	108
272	Evolution for exogenous octanoic acid tolerance improves carboxylic acid production and membrane integrity. <i>Metabolic Engineering</i> , 2015 , 29, 180-188	9.7	77
271	Metabolic engineering of Clostridium tyrobutyricum for n-butanol production: effects of CoA transferase. 2015 , 99, 4917-30		34
270	cAMP receptor protein (CRP)-mediated resistance/tolerance in bacteria: mechanism and utilization in biotechnology. 2015 , 99, 4533-43		19
269	Dynamics of membrane fatty acid composition of succinic acid-producing Anaerobiospirillum succiniciproducens. 2015 , 193, 130-3		8
268	Chaotropicity: a key factor in product tolerance of biofuel-producing microorganisms. 2015 , 33, 228-59		117
267	Membrane transporter engineering in industrial biotechnology and whole cell biocatalysis. 2015 , 33, 237-46		136
266	Toxicological challenges to microbial bioethanol production and strategies for improved tolerance. 2015 , 24, 2156-74		10
265	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. 2015 , 54, 10170-10182		22
264	Applying systems biology tools to studyn-butanol degradation inPseudomonas putidaKT2440. 2015 , 15, 760-771		20
263	Metabolic and process engineering of Clostridium cellulovorans for biofuel production from cellulose. <i>Metabolic Engineering</i> , 2015 , 32, 39-48	9.7	96
262	Synthetic scaffolds for pathway enhancement. 2015 , 36, 98-106		63
261	Building cellular pathways and programs enabled by the genetic diversity of allo-genomes and meta-genomes. 2015 , 36, 16-31		1

(2016-2015)

260	A metabolomics-based strategy for identification of gene targets for phenotype improvement and its application to 1-butanol tolerance in Saccharomyces cerevisiae. 2015 , 8, 144	24
259	Evaluation of hydrophobic micro-zeolite-mixed matrix membrane and integrated with acetone-butanol-ethanol fermentation for enhanced butanol production. 2015 , 8, 105	45
258	Current status and prospects of industrial bio-production of n-butanol in China. <i>Biotechnology Advances</i> , 2015 , 33, 1493-501	118
257	Trade-Offs in Improving Biofuel Tolerance Using Combinations of Efflux Pumps. 2015 , 4, 1056-63	47
256	Engineering Clostridium acetobutylicum with a histidine kinase knockout for enhanced n-butanol tolerance and production. 2015 , 99, 1011-22	99
255	Microorganisms in Biorefineries. 2015 ,	3
254	Engineering Propionibacterium freudenreichii subsp. shermanii for enhanced propionic acid fermentation: effects of overexpressing propionyl-CoA:Succinate CoA transferase. <i>Metabolic</i> 9.7 <i>Engineering</i> , 2015 , 27, 46-56	43
253	Can Microbially Derived Advanced Biofuels Ever Compete with Conventional Bioethanol? A Critical Review. 2016 , 11,	2
252	Pathway and Strain Design for Biofuels Production. 2016 , 97-116	2
251	The Role of GroE Chaperonins in Developing Biocatalysts for Biofuel and Chemical Production. 2016 , 05,	2
250	Evolutionary Methods for Improving the Production of Biorenewable Fuels and Chemicals. 2016 , 265-290	8
249	Rebalancing Redox to Improve Biobutanol Production by. 2015 , 3,	9
248	Characterizing the Phenotypic Responses of Escherichia coli to Multiple 4-Carbon Alcohols with Raman Spectroscopy. 2016 , 2, 3	9
247	Metabolic Fingerprinting of Pseudomonas putida DOT-T1E Strains: Understanding the Influence of Divalent Cations in Adaptation Mechanisms Following Exposure to Toluene. 2016 , 6,	
246	Regulatory mechanisms related to biofuel tolerance in producing microbes. 2016 , 121, 320-32	5
245	Inhibition of dark fermentative bio-hydrogen production: A review. 2016 , 41, 6713-6733	172
244	Transcriptional kinetics of the cross-talk between the ortho-cleavage and TOL pathways of toluene biodegradation in Pseudomonas putida mt-2. 2016 , 228, 112-123	10
243	Very high gravity ethanol and fatty acid production of Zymomonas mobilis without amino acid and vitamin. 2016 , 43, 861-71	23

242	Design of online off-gas analysis system for anaerobic ABE fermentation and the strategy for improving biobutanol production. 2016 , 51, 555-560	4
241	Impact of furfural on biohydrogen production from glucose and xylose in continuous-flow systems. 2016 , 93, 302-311	37
240	Mycothiol peroxidase MPx protects Corynebacterium glutamicum against acid stress by scavenging ROS. 2016 , 38, 1221-8	7
239	A re-look at the biochemical strategies to enhance butanol production. 2016 , 94, 187-200	41
238	Intracellular metabolic changes of Clostridium acetobutylicum and promotion to butanol tolerance during biobutanol fermentation. 2016 , 78, 297-306	17
237	Cell-Free Mixing of Escherichia coli Crude Extracts to Prototype and Rationally Engineer High-Titer Mevalonate Synthesis. 2016 , 5, 1578-1588	101
236	A Two-Layer Gene Circuit for Decoupling Cell Growth from Metabolite Production. 2016 , 3, 133-143	70
235	Enhance nisin yield via improving acid-tolerant capability of Lactococcus lactis F44. 2016 , 6, 27973	32
234	Anaerobes in Biotechnology. 2016 ,	6
233	Bioconversion of L-phenylalanine to 2-phenylethanol by the novel stress-tolerant yeast Candida glycerinogenes WL2002-5. 2016 , 7, 418-423	27
232	Engineering membrane and cell-wall programs for tolerance to toxic chemicals: Beyond solo genes. 2016 , 33, 56-66	51
231	Ionic liquid-tolerant microorganisms and microbial communities for lignocellulose conversion to bioproducts. 2016 , 100, 10237-10249	34
230	CO fixation by anaerobic non-photosynthetic mixotrophy for improved carbon conversion. 2016 , 7, 12800	86
229	The Saccharomyces cerevisiae poly(A) binding protein Pab1 as a target for eliciting stress tolerant phenotypes. 2015 , 5, 18318	9
228	Genome-wide Escherichia coli stress response and improved tolerance towards industrially relevant chemicals. <i>Microbial Cell Factories</i> , 2016 , 15, 176	32
227	Enhancing muconic acid production from glucose and lignin-derived aromatic compounds via increased protocatechuate decarboxylase activity. 2016 , 3, 111-119	149
226	Biological Processes for Hydrogen Production. 2016 , 156, 155-193	4
225	Simultaneous enhancement of phenolic compound degradations by Acinetobacter strain V2 via a step-wise continuous acclimation process. 2016 , 56, 627-34	8

224	Metabolic analysis of the response of DOT-T1E strains to toluene using Fourier transform infrared spectroscopy and gas chromatography mass spectrometry. 2016 , 12, 112	7
223	Genetic engineering of Bacillus megaterium for high-yield production of the major teleost progestogens 17段0団- and 17段0D1昰rihydroxy-4-pregnen-3-one. <i>Metabolic Engineering</i> , 9.7 2016 , 36, 19-27	9
222	Membrane engineering via trans unsaturated fatty acids production improves Escherichia coli robustness and production of biorenewables. <i>Metabolic Engineering</i> , 2016 , 35, 105-113	76
221	Comparative transcriptomics elucidates adaptive phenol tolerance and utilization in lipid-accumulating Rhodococcus opacus PD630. 2016 , 44, 2240-54	76
220	Anaerobic Fermentation for Production of Carboxylic Acids as Bulk Chemicals from Renewable Biomass. 2016 , 156, 323-361	18
219	Application of nanofiltration for production of 1,3-propanediol in membrane bioreactor. 2016 , 268, 164-170	9
218	Gene expression and molecular characterization of a chaperone protein HtpG from Bacillus licheniformis. 2016 , 85, 179-91	1
217	Balancing Photosynthesis and Respiration Increases Microalgal Biomass Productivity during Photoheterotrophy on Glycerol. 2016 , <i>4</i> , 1611-1618	33
216	Bioreactor Engineering Research and Industrial Applications I. 2016,	4
215	Enhancing acetone biosynthesis and acetone-butanol-ethanol fermentation performance by co-culturing Clostridium acetobutylicum/Saccharomyces cerevisiae integrated with exogenous acetate addition. 2016 , 200, 111-20	38
214	The ability of Achromobacter sp. KW1 strain to biodegrade isomers of chlorotoluene. 2017 , 92, 2134-2141	9
213	Metabolic engineering of Clostridium tyrobutyricum for n-butanol production from sugarcane juice. 2017 , 101, 4327-4337	27
212	Membrane engineering of S. cerevisiae targeting sphingolipid metabolism. 2017 , 7, 41868	6
211	Microbial response to environmental stresses: from fundamental mechanisms to practical applications. 2017 , 101, 3991-4008	54
210	A New Player in the Biorefineries Field: Phasin PhaP Enhances Tolerance to Solvents and Boosts Ethanol and 1,3-Propanediol Synthesis in Escherichia coli. 2017 , 83,	18
209	Metabolic engineering of Escherichia coli for higher alcohols production: An environmentally friendly alternative to fossil fuels. 2017 , 77, 580-589	15
208	Tuning and elucidation of the colony dimorphism in Rhodococcus ruber associated with cell flocculation in large scale fermentation. 2017 , 101, 6321-6332	10
207	Engineering tolerance to industrially relevant stress factors in yeast cell factories. 2017 , 17,	88

206	The Efficient Clade: Lactic Acid Bacteria for Industrial Chemical Production. 2017, 35, 756-769		70
205	Engineering microbial phenotypes through rewiring of genetic networks. 2017 , 45, 4984-4993		8
204	The impact of succinate trace on pWW0 and ortho-cleavage pathway transcription in Pseudomonas putida mt-2 during toluene biodegradation. 2017 , 234, 397-405		10
203	High-efficient n-butanol production by co-culturing Clostridium acetobutylicum and Saccharomyces cerevisiae integrated with butyrate fermentative supernatant addition. <i>World Journal of Microbiology and Biotechnology</i> , 2017 , 33, 76	4.4	29
202	Reassessing Escherichia coli as a cell factory for biofuel production. 2017 , 45, 92-103		39
201	Systematic and functional identification of small non-coding RNAs associated with exogenous biofuel stress in cyanobacterium sp. PCC 6803. 2017 , 10, 57		19
200	Expression of Heterologous Sigma Factor Expands the Searchable Space for Biofuel Tolerance Mechanisms. 2017 , 6, 1343-1350		8
199	Proteomic Analysis of Metabolic Responses to Biofuels and Chemicals in Photosynthetic Cyanobacteria. 2017 , 585, 355-376		1
198	Compatible solute addition to biological systems treating waste/wastewater to counteract osmotic and other environmental stresses: a review. 2017 , 37, 865-879		40
197	The Tolerome: A Database of Transcriptome-Level Contributions to Diverse Escherichia coli Resistance and Tolerance Phenotypes. 2017 , 6, 2302-2315		7
196	Comparative genomic analysis of Clostridium acetobutylicum for understanding the mutations contributing to enhanced butanol tolerance and production. 2017 , 263, 36-44		27
195	Chassis and key enzymes engineering for monoterpenes production. <i>Biotechnology Advances</i> , 2017 , 35, 1022-1031	17.8	17
194	Comparative phenotypic analysis of "Clostridium neonatale" and Clostridium butyricum isolates from neonates. 2017 , 48, 76-82		8
193	Inactivation of Ascaris Eggs in Human Fecal Material Through In Situ Production of Carboxylic Acids. 2017 , 51, 9729-9738		12
192	Efficacy of ionic liquids on the growth and simultaneous xylanase production by Sporotrichum thermophile: membrane integrity, composition and morphological investigation. 2017 , 7, 21114-21123		7
191	Bioremediation 3.0: Engineering pollutant-removing bacteria in the times of systemic biology. <i>Biotechnology Advances</i> , 2017 , 35, 845-866	17.8	165
190	Enhanced robustness in acetone-butanol-ethanol fermentation with engineered Clostridium beijerinckii overexpressing adhE2 and ctfAB. 2017 , 243, 1000-1008		25
189	Improving Escherichia coli membrane integrity and fatty acid production by expression tuning of FadL and OmpF. <i>Microbial Cell Factories</i> , 2017 , 16, 38	6.4	33

188	The significance of proline and glutamate on butanol chaotropic stress in 168. 2017 , 10, 122		15
187	Dynamics of two methanogenic microbiomes incubated in polycyclic aromatic hydrocarbons, naphthenic acids, and oil field produced water. 2017 , 10, 123		3
186	Comparative genomic analysis of Mycobacterium neoaurum MN2 and MN4 substrate and product tolerance. 2017 , 7, 181		2
185	Development of stress tolerant Saccharomyces cerevisiae strains by metabolic engineering: New aspects from cell flocculation and zinc supplementation. 2017 , 123, 141-146		19
184	Growth of Pseudomonas taiwanensis VLB120 biofilms in the presence of n-butanol. 2017, 10, 745-755		12
183	Microbial Bioprocesses. 2017 , 45-72		
182	Engineering microbial cell factories for the production of plant natural products: from design principles to industrial-scale production. <i>Microbial Cell Factories</i> , 2017 , 16, 125	6.4	64
181	Awakening sleeping beauty: production of propionic acid in Escherichia coli through the sbm operon requires the activity of a methylmalonyl-CoA epimerase. <i>Microbial Cell Factories</i> , 2017 , 16, 121	6.4	9
180	Improving phloroglucinol tolerance and production in Escherichia coli by GroESL overexpression. <i>Microbial Cell Factories</i> , 2017 , 16, 227	6.4	14
179	Comprehensive characterization of toxicity of fermentative metabolites on microbial growth. 2017 , 10, 262		41
178	Native efflux pumps of Escherichia coli responsible for short and medium chain alcohol. 2018 , 133, 149-	156	14
177	Enhanced butyric acid production in Clostridium tyrobutyricum by overexpression of rate-limiting enzymes in the Embden-Meyerhof-Parnas pathway. 2018 , 272-273, 14-21		15
176	Biotreatment optimization of rice straw hydrolyzates for ethanolic fermentation with Scheffersomyces stipitis. 2018 , 112, 19-28		13
175	Integrated analysis of isopentenyl pyrophosphate (IPP) toxicity in isoprenoid-producing Escherichia coli. <i>Metabolic Engineering</i> , 2018 , 47, 60-72	9.7	62
174	Enhanced Delignification of Lignocellulosic Biomass by Recombinant Fungus Phanerochaete chrysosporium Overexpressing Laccases and Peroxidases. 2018 , 28, 1-13		13
173	Effect of nitrogen gas sparging on dark fermentative biohydrogen production using suspended and immobilized cells of anaerobic mixed bacteria from potato waste. 2018 , 9, 595-604		7
172	Escherichia coli as a host for metabolic engineering. <i>Metabolic Engineering</i> , 2018 , 50, 16-46	9.7	153
171	Thermochemical wastewater valorization via enhanced microbial toxicity tolerance. <i>Energy and Environmental Science</i> , 2018 , 11, 1625-1638	35.4	51

170	Enhanced polymalic acid production from the glyoxylate shunt pathway under exogenous alcohol stress. 2018 , 275, 24-30		9
169	Biofuel production with a stress-resistant and growth phase-independent promoter: mechanism revealed by in vitro transcription assays. 2018 , 102, 2929-2940		7
168	Fungal synthesis of chiral phosphonic synthetic platform - Scope and limitations of the method. <i>Bioorganic Chemistry</i> , 2018 , 77, 402-410	5.1	6
167	Methods for enhancing cyanobacterial stress tolerance to enable improved production of biofuels and industrially relevant chemicals. 2018 , 102, 1617-1628		13
166	Comparative analysis of high butanol tolerance and production in clostridia. <i>Biotechnology Advances</i> , 2018 , 36, 721-738	17.8	37
165	Energy efficiency of acetone, butanol, and ethanol (ABE) recovery by heat-integrated distillation. 2018 , 41, 395-405		20
164	Prospects of Solvent Tolerance in Butanol Fermenting Bacteria. <i>Biofuel and Biorefinery Technologies</i> , 2018 , 249-264	1	3
163	Genome-wide identification of tolerance mechanisms toward p-coumaric acid in Pseudomonas putida. 2018 , 115, 762-774		40
162	. 2018,		0
161	Metabolic Engineering for Biocatalyst Robustness to Organic Inhibitors. 2018 , 239-265		1
160	Improved acid-stress tolerance of Lactococcus lactis NZ9000 and Escherichia coli BL21 by overexpression of the anti-acid component recT. 2018 , 45, 1091-1101		7
159	Synthesis and techno-economic assessment of microbial-based processes for terpenes production. 2018 , 11, 294		21
158	A framework for the identification of promising bio-based chemicals. 2018 , 115, 2328-2340		15
157	Modification of membrane lipid compositions in single-celled organisms - From basics to applications. 2018 , 147, 50-65		23
157 156		6.4	23 18
	applications. 2018, 147, 50-65 Engineered global regulator H-NS improves the acid tolerance of E. coli. <i>Microbial Cell Factories</i> ,	6.4	
156	applications. 2018, 147, 50-65 Engineered global regulator H-NS improves the acid tolerance of E. coli. <i>Microbial Cell Factories</i> , 2018, 17, 118 Spatial organization of enzymes to enhance synthetic pathways in microbial chassis: a systematic	·	18

152	The effects of disruption in membrane lipid biosynthetic genes on 1-butanol tolerance of Bacillus subtilis. 2018 , 102, 9279-9289		3	
151	Efficient yeast surface-display of novel complex synthetic cellulosomes. <i>Microbial Cell Factories</i> , 2018 , 17, 122	6.4	23	
150	Problems of Solventogenicity, Solvent Tolerance: An Introduction. 2018, 327-334		1	
149	Enhancement of antroquinonol production during batch fermentation using pH control coupled with an oxygen vector. 2019 , 99, 449-456		8	
148	Chasing bacterial chassis for metabolic engineering: a perspective review from classical to non-traditional microorganisms. 2019 , 12, 98-124		112	
147	Enhancement of antroquinonol and antrodin C productions via in situ extractive fermentation of Antrodia camphorata S-29. 2019 , 103, 8351-8361		4	
146	Bacterial Acid Resistance Toward Organic Weak Acid Revealed by RNA-Seq Transcriptomic Analysis in. <i>Frontiers in Microbiology</i> , 2019 , 10, 1616	5.7	13	
145	Enhancing butanol tolerance of reveals hydrophobic interaction of multi-tasking chaperone SecB. 2019 , 12, 164		6	
144	Quantitative proteomic analysis reveals the ethanologenic metabolism regulation of by exogenous ethanol addition. 2019 , 12, 166		11	
143	Perspectives, Scope, Advancements, and Challenges of Microbial Technologies Treating Textile Industry Effluents. 2019 , 237-260		3	
142	Effect of Sequential Acclimation to Various Carbon Sources on the Proteome of LMG 23690 and Its Tolerance to Downstream Process Stresses. <i>Frontiers in Microbiology</i> , 2019 , 10, 608	5.7	3	
141	A review on commercial-scale high-value products that can be produced alongside cellulosic ethanol. 2019 , 12, 240		213	
140	Evaluation of Relationship Between Crystallization Structure and Thermal-Mechanical Performance of PLA with MCC Addition. 2019 , 4, 10174-10180		3	
139	The fabrication of a 3D current collector with bitter melon-like TiO2NCNFs for highly stable lithiumBulfur batteries. 2019 , 1, 527-531		2	
138	Enhanced butanol-hydrogen coproduction by Clostridium beijerinckii with biochar as cell's carrier. 2019 , 294, 122141		29	
137	Effect of the Structural and Regulatory Heat Shock Proteins on Hydrocarbon Degradation by Rhodococcus pyridinivorans 5Ap. 2019 , 88, 573-579		3	
136	New technologies provide more metabolic engineering strategies for bioethanol production in Zymomonas mobilis. 2019 , 103, 2087-2099		29	
135	Improved -Butanol Production from Clostridium cellulovorans by Integrated Metabolic and Evolutionary Engineering. 2019 , 85,		53	

Efficient synthesis of enantiopure amines from alcohols using resting cells and ammonia. 2019, 21, 3846-3857 17 134 Tolerance against butanol stress by disrupting succinylglutamate desuccinylase in .. 2019, 9, 11683-11695 133 4 Overexpression of endogenous stress-tolerance related genes in Saccharomyces cerevisiae 132 11 improved strain robustness and production of heterologous cellobiohydrolase. 2019, 19, Phenotypic and genomic analysis of ZM4 mutants with enhanced ethanol tolerance. 2019, 23, e00328 131 On the use of oxygenic photosynthesis for the sustainable production of commodity chemicals. 130 7 2019. 166. 413-427 Co-production of solvents and organic acids in butanol fermentation by in the presence of 129 13 lignin-derived phenolics.. 2019, 9, 6919-6927 Exceptional solvent tolerance in Yarrowia lipolytica is enhanced by sterols. Metabolic Engineering, 128 9.7 25 **2019**, 54, 83-95 Improvement of stress tolerance and riboflavin production of Bacillus subtilis by introduction of 16 127 heat shock proteins from thermophilic bacillus strains. 2019, 103, 4455-4465 Mixed-species biofilms for high-cell-density application of Synechocystis sp. PCC 6803 in capillary 126 38 reactors for continuous cyclohexane oxidation to cyclohexanol. 2019, 282, 171-178 Expression of Metazoan Annexins in Yeast Provides Protection Against Deleterious Effects of the 125 Biofuel Isobutanol. 2019, 9, 18603 Clostridium sp. as Bio-Catalyst for Fuels and Chemicals Production in a Biorefinery Context. 2019, 9, 962 124 18 Enhancing Expression of 3-Ketosteroid-9Hydroxylase Oxygenase, an Enzyme with Broad Substrate Range and High Hydroxylation Ability, in Mycobacterium sp. LY-1. 2019, 187, 1238-1254 Cyanobacterial Secretion Systems: Understanding Fundamental Mechanisms Toward Technological 8 122 Applications. 2019, 359-381 The genome and transcriptome of Lactococcus lactis ssp. lactis F44 and G423: Insights into 121 6 adaptation to the acidic environment. 2019, 102, 1044-1058 Biosynthesis of Polyphenols in Recombinant Micro-organisms. 2019, 237-262 120 1 119 Cellulolytic thermophilic microorganisms in white biotechnology: a review. 2020, 65, 25-43 11 Trends in Systems Biology for the Analysis and Engineering of Clostridium acetobutylicum 118 14 Metabolism. 2020, 28, 118-140 Aspergillus niger upregulated glycerolipid metabolism and ethanol utilization pathway under 117 ethanol stress. **2020**, 9, e00948

(2020-2020)

116	Screening a genomic library for genes involved in propionate tolerance in Yarrowia lipolytica. 2020 , 37, 131-140		6
115	Consolidated bioprocessing for butanol production of cellulolytic Clostridia: development and optimization. 2020 , 13, 410-422		22
114	The biological mechanisms of butanol tolerance and the application of solvent-tolerant bacteria for environmental protection. 2020 , 95, 1290-1297		2
113	Microbial response to acid stress: mechanisms and applications. 2020 , 104, 51-65		104
112	Biofuels Production (S ustainability and Advances in Microbial Bioresources. <i>Biofuel and Biorefinery Technologies</i> , 2020 ,	1	11
111	Degeneration of industrial bacteria caused by genetic instability. <i>World Journal of Microbiology and Biotechnology</i> , 2020 , 36, 119	4.4	8
110	Optimization of n-butanol synthesis in Lactobacillus brevis via the functional expression of thl, hbd, crt and ter. 2020 , 47, 1099-1108		2
109	Automated Conditional Screening of Multiple Strains in Parallel Adaptive Fed-Batch Cultivations. 2020 , 7,		5
108	Production optimization of yellow laccase from Stropharia sp. ITCC 8422 and enzyme-mediated depolymerization and hydrolysis of lignocellulosic biomass for biorefinery application. 2020 , 1		24
107	Ink-Based Additive Nanomanufacturing of Functional Materials for Human-Integrated Smart Wearables. 2020 , 2, 2000117		9
106	Metabolic and Process Engineering of for Butyl Acetate Production in One Step. 2020 , 68, 9475-9487		7
105	Clostridium thermocellum: A microbial platform for high-value chemical production from lignocellulose. 2020 , 113, 111-161		8
104	Role of efflux in enhancing butanol tolerance of bacteria. 2020 , 320, 17-27		12
103	Pathway dissection, regulation, engineering and application: lessons learned from biobutanol production by solventogenic clostridia. 2020 , 13, 39		36
102	Enzyme Assembly for Compartmentalized Metabolic Flux Control. 2020, 10,		4
101	Half-Preparative Scale Synthesis of (S)-1-Phenylethane-1,2-Diol as a Result of 2-Phenylethanol Hydroxylation with Aspergillus niger (IAFB 2301) Assistance. 2020 , 12, 989		1
100	Degradation and transformation of furfural derivatives from hydrothermal pre-treated algae and lignocellulosic biomass during hydrogen fermentation. 2020 , 131, 109983		8
99	Microbial Diversity, Interventions and Scope. 2020 ,		1

98	Compatible solutes adaptive alterations in Arthrobacter simplex during exposure to ethanol, and the effect of trehalose on the stress resistance and biotransformation performance. 2020 , 43, 895-908		3
97	An increase in cell membrane permeability in the in situ extractive fermentation improves the production of antroquinonol from Antrodia camphorata S-29. 2020 , 47, 197-207		4
96	Improving dark fermentative hydrogen production through zero-valent iron/copper (Fe/Cu) micro-electrolysis. 2020 , 42, 445-451		3
95	Common problems associated with the microbial productions of aromatic compounds and corresponding metabolic engineering strategies. <i>Biotechnology Advances</i> , 2020 , 41, 107548	17.8	21
94	Enhanced Production of 2,3-Butanediol in Recombinant Escherichia coli Using Response Regulator DR1558 Derived from Deinococcus radiodurans. 2020 , 25, 45-52		8
93	How to outwit nature: Omics insight into butanol tolerance. <i>Biotechnology Advances</i> , 2021 , 46, 107658	17.8	6
92	Bioremediation of heavy metals from wastewater using nanomaterials. 2021 , 23, 9617-9640		16
91	Reducing agents assisted fed-batch fermentation to enhance ABE yields. 2021 , 227, 113627		6
90	Recent technological and strategical developments in the biomanufacturing of 1,3-propanediol from glycerol. 2021 , 18, 2467-2490		6
89	Strain dynamics of specific contaminant bacteria modulate the performance of ethanol biorefineries.		1
88	Real-Time Optogenetics System for Controlling Gene Expression Using a Model-Based Design. 2021 , 93, 3181-3188		1
87	Effect of conductive material for overcoming inhibitory conditions derived from red algae-based substrate on biohydrogen production. 2021 , 285, 119059		7
86	Quorum Sensing-Mediated and Growth Phase-Dependent Regulation of Metabolic Pathways in H4. <i>Frontiers in Microbiology</i> , 2021 , 12, 567942	5.7	1
85	Inhibition of the metabolism of mixed cultures of purple phototrophic bacteria by typical refinery and petrochemistry wastewater pollutants. 2021 , 96, 1893-1901		
84	Sulphate-Reducing Bacteria Response to Extreme pH Environments and the Effect of Their Activities on Microbial Corrosion. 2021 , 11, 2201		5
83	Product Export in Cyanobacteria. 2021 , 369-406		3
82	Adaptive Laboratory Evolution Restores Solvent Tolerance in Plasmid-Cured Pseudomonas putida S12: a Molecular Analysis. 2021 , 87,		4
81	Homology- and cross-resistance of Lactobacillus plantarum to acid and osmotic stress and the influence of induction conditions on its proliferation by RNA-Seq. 2021 , 61, 576-590		1

80	A high-efficient strategy for combinatorial engineering paralogous gene family: A case study on histidine kinases in Clostridium. 2021 , 118, 2770-2780		2
79	Exposure to 1-Butanol Exemplifies the Response of the Thermoacidophilic Archaeon Sulfolobus acidocaldarius to Solvent Stress. 2021 , 87,		3
78	Proteomic Responses to Butanol Stress. <i>Frontiers in Microbiology</i> , 2021 , 12, 674639	5.7	2
77	Dynamic regulation of membrane integrity to enhance l-malate stress tolerance in Candida glabrata. 2021 , 118, 4347-4359		2
76	The Role of Metabolic Engineering Technologies for the Production of Fatty Acids in Yeast. 2021 , 10,		2
75	The Plasma Membrane at the Cornerstone Between Flexibility and Adaptability: Implications for as a Cell Factory. <i>Frontiers in Microbiology</i> , 2021 , 12, 715891	5.7	2
74	Raman spectroscopy in cell biology and microbiology.		10
73	Physiological limitations and opportunities in microbial metabolic engineering. 2022 , 20, 35-48		8
72	Biocatalysis in Green and Blue: Cyanobacteria. 2021 , 39, 875-889		11
71	ComX improves acid tolerance by regulating the expression of late competence proteins in Lactococcus lactis F44. 2021 , 104, 9556-9569		O
70	Bisphenol A Removal by the Fungus IM 6482-Analysis of the Cellular and Subcellular Level. 2021 , 22,		2
69	Sweet sorghum juice as an alternative carbon source and adaptive evolution of Lactobacillus brevis NIE9.3.3 in sweet sorghum juice and biodiesel derived crude glycerol to improve 1, 3 propanediol production. <i>Journal of Environmental Chemical Engineering</i> , 2021 , 9, 106086	5.8	2
68	Challenges and opportunities in biological funneling of heterogeneous and toxic substrates beyond lignin. 2021 , 73, 1-13		9
67	Efflux identification and engineering for ansamitocin P-3 production in Actinosynnema pretiosum. 2021 , 105, 695-706		3
66	An Adaptive Laboratory Evolution Method to Accelerate Autotrophic Metabolism. 2018, 1671, 149-161		2
65	Insights into Organic-Solvent-Tolerant Bacteria and Their Biotechnological Potentials. 2015 , 129-149		1
64	Tolerance of Microbial Biocatalysts to Feedstocks, Products, and Environmental Conditions. 2016 , 73-100)	2
63	Evolutionary Engineering of Microorganisms to Overcome Toxicity During Lignocellulose Hydrolysates Utilization. 2017 , 181-200		1

62	Challenges in the Application of Synthetic Biology Toward Synthesis of Commodity Products by Cyanobacteria via "Direct Conversion". <i>Advances in Experimental Medicine and Biology</i> , 2018 , 1080, 3-26	3.6	9
61	Engineering transport systems for microbial production. 2020 , 111, 33-87		6
60	Building cell factories for the production of advanced fuels. 2019 , 47, 1701-1714		4
59	Exceptional Solvent Tolerance inYarrowia lipolyticals Enhanced by Sterols.		3
58	Synthetic biology: mapping the scientific landscape. <i>PLoS ONE</i> , 2012 , 7, e34368	3.7	63
57	Identification and characterization of two functionally unknown genes involved in butanol tolerance of Clostridium acetobutylicum. <i>PLoS ONE</i> , 2012 , 7, e38815	3.7	23
56	Improving fatty acid availability for bio-hydrocarbon production in Escherichia coli by metabolic engineering. <i>PLoS ONE</i> , 2013 , 8, e78595	3.7	21
55	Metabolomics Analysis Reveals the Participation of Efflux Pumps and Ornithine in the Response of Pseudomonas putida DOT-T1E Cells to Challenge with Propranolol. <i>PLoS ONE</i> , 2016 , 11, e0156509	3.7	8
54	Fumarate Production by Torulopsis glabrata: Engineering Heterologous Fumarase Expression and Improving Acid Tolerance. <i>PLoS ONE</i> , 2016 , 11, e0164141	3.7	6
53	Major progress in microbial ecology of hypoxia in the shallow eutrophic lakes. <i>Hupo Kexue/Journal of Lake Sciences</i> , 2015 , 27, 567-574	0.5	4
52	Beneficial biofilms. <i>AIMS Bioengineering</i> , 2015 , 2, 437-448	3.4	9
51	Industrial production, application, microbial biosynthesis and degradation of furanic compound, hydroxymethylfurfural (HMF). <i>AIMS Microbiology</i> , 2018 , 4, 261-273	4.5	20
50	Technological Approach of Bioremediation Using Microbial Tools. <i>Advances in Environmental Engineering and Green Technologies Book Series</i> , 2017 , 134-154	0.4	5
49	Characterizing metabolic stress-induced phenotypes of PCC6803 with Raman spectroscopy. <i>PeerJ</i> , 2020 , 8, e8535	3.1	7
48	Recent progress on n-butanol production by lactic acid bacteria. <i>World Journal of Microbiology and Biotechnology</i> , 2021 , 37, 205	4.4	O
47	Preliminary Study of Biotransformation of Aldehydes and Ketones by Clostridium saccharoperbutylacetonicum N1-4 (ATCC 13564). <i>Biotechnology</i> , 2013 , 12, 179-182	0.1	
46	Engineering Central Metabolism for Production of Higher Alcohol-based Biofuels. 2016 , 1-34		2
45	Problems of Solventogenicity, Solvent Tolerance: An Introduction. 2017 , 1-8		

Comprehensive Characterization of Toxicity of Fermentative Metabolites on Microbial Growth.

43	Transporters Related to Stress Responses and Their Potential Application in Synechocystis sp. PCC 6803. <i>Advances in Experimental Medicine and Biology</i> , 2018 , 1080, 27-53	3.6	
42	Bacteria for Butanol Production: Bottlenecks, Achievements and Prospects. <i>Journal of Pure and Applied Microbiology</i> , 2019 , 13, 1429-1440	0.9	
41	Cyanobacterial Biofuel Production: Current Development, Challenges and Future Needs. <i>Biofuel and Biorefinery Technologies</i> , 2020 , 35-62	1	1
40	Microbial Clean Up Strategy for Polluted Water. 2020 , 219-229		
39	Microbes: The Next-Generation Bioenergy Producers. 2020 , 29-60		
38	Factors affecting the bioremediation of industrial and domestic wastewaters. 2022, 461-472		О
37	Global Climate Change and Microbial Ecology: Current Scenario and Management. 2021 , 285-313		
36	Toward improved terpenoids biosynthesis: strategies to enhance the capabilities of cell factories. <i>Bioresources and Bioprocessing</i> , 2022 , 9,	5.2	О
35	Prospects of microbes in mitigations of environmental degradation in the liver ecosystem. 2022, 429-4	154	Ο
34	Molecular biology: Fantastic toolkits to improve knowledge and application of acetic acid bacteria <i>Biotechnology Advances</i> , 2022 , 107911	17.8	О
33	Toward low-cost biological and hybrid biological/catalytic conversion of cellulosic biomass to fuels. Energy and Environmental Science,	35.4	7
32	Enhancement of triterpene production via in situ extractive fermentation of Sanghuangporus vaninii YC-1 <i>Biotechnology and Applied Biochemistry</i> , 2021 ,	2.8	1
31	Exploring the limits of carbohydrate conversion and product formation in open mixed culture fermentation. <i>Journal of Environmental Chemical Engineering</i> , 2022 , 10, 107513	6.8	Ο
30	Engineering for life in toxicity: key to industrializing microbial synthesis of high energy density fuels. <i>Engineering Microbiology</i> , 2022 , 2, 100013		
29	Metabolic transformation of cyanobacteria for biofuel production Chemosphere, 2022, 134342	8.4	O
28	Consistent Value Creation from Bioprocess Data with Customized Algorithms: Opportunities Beyond Multivariate Analysis. 2022 , 231-264		1
27	Adult Worm Protective and Diagnostic Proteins in -Butanol Extracts Revealed by Proteomic Analysis <i>Pathogens</i> , 2021 , 11,	4.5	

- 26 Table_1.XLSX. **2019**,
- 25 Table_1.xlsx. **2019**,
- 24 Table_2.xlsx. **2019**,

18

12

- 23 Cross-Protection Response. 2022, 549-573
- 22 Sublethal Injury Adaptation in Foodborne Pathogens. 2022, 517-536
- An updated review on advancement in fermentative production strategies for biobutanol using

 Clostridium spp.. Environmental Science and Pollution Research, 2022,

 5.1 O
- Current knowledge on cyanobacterial biobutanol production: advances, challenges, and prospects.

 Reviews in Environmental Science and Biotechnology, 1
- 13.9 1
- Screening for amidoxime reductases in plant roots and Saccharomyces cerevisiae Development of biocatalytic method for chemoselective amidine synthesis.. *Bioorganic Chemistry*, **2022**, 124, 105815

Bioengineering in microbial production of biobutanol from renewable resources. 2022, 307-334

- - О
- Compartmentalization and transporter engineering strategies for terpenoid synthesis. *Microbial Cell Factories*, **2022**, 21,
- 6.4 2
- Enhanced Fermentative Hydrogen Production from Food Waste in Continuous Reactor after Butyric Acid Treatment. *Energies*, **2022**, 15, 4048
- 3.1 0
- Modification of substrate and product transport systems in Klebsiella pneumoniae to improve 1,3-propanediol production. *FEMS Microbiology Letters*,
- 2.9
- Editorial: Microorganisms for Consolidated 2nd Generation Biorefining. *Frontiers in Microbiology*, 13,
- 5.7
- Bioethanol production from biodegradable wastes using native yeast isolates from Ethiopian traditional alcoholic beverages. *Biocatalysis and Agricultural Biotechnology*, **2022**, 43, 102401
- 4.2 0
- Metabolic Engineering of Shikimic Acid Biosynthesis Pathway for the Production of Shikimic Acid and Its Branched Products in Microorganisms: Advances and Prospects. **2022**, 27, 4779
- О
- Listeria monocytogenes is a solvent tolerant organism secreting a solvent stable lipase: potential biotechnological applications.
- Co-expression of an isopropanol synthetic operon and eGFP to monitor the robustness of Cupriavidus necator during isopropanol production. **2022**, 110114
- Enhancing the prebiotic and antioxidant effects of exopolysaccharides derived from Cordyceps militaris by enzyme-digestion. **2022**, 167, 113830

CITATION REPORT

8	Bioremediation on the Crossroads of Technology for Environmental Clean-Up: An Overview. 2022 , 3-25	O
7	Molecular Tools- A Future Perspective Approach for Monitoring Landfill Leachates and Validating Bioremediation Process. 2022 , 95-113	O
6	Current progress on engineering microbial strains and consortia for production of cellulosic butanol through consolidated bioprocessing.	0
5	Novel approaches toward bio-butanol production from renewable feedstocks. 2023 , 105-138	O
4	Comparison of plasmid stabilization systems during heterologous isopropanol production in fed-batch bioreactor. 2023 , 366, 25-34	0
3	Adaptive laboratory evolution boosts Yarrowia lipolytica tolerance to vanillic acid. 2023, 367, 42-52	O
2	Antarctic heavy metal pollution and remediation efforts: state of the art of research and scientific publications.	0
1	Strain and process engineering toward continuous industrial fermentation.	0