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List of Publications by Year in descending order

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567281 501196 38 825 15 28 citations h-index g-index papers 38 38 38 1099 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Lignocellulosic ethanol: Technology design and its impact on process efficiency. Biotechnology Advances, 2015, 33, 1091-1107.	11.7	151
2	Electrodialysis as a useful technique for lactic acid separation from a model solution and a fermentation broth. Desalination, 2004, 162, 361-372.	8.2	115
3	Use of a mixture of glucose and methanol as substrates for the production of recombinant trypsinogen in continuous cultures with Pichia pastoris Mut+. Journal of Biotechnology, 2012, 157, 180-188.	3.8	52
4	Transformation of raw feather waste into digestible peptides and amino acids. Journal of Chemical Technology and Biotechnology, 2016, 91, 1629-1637.	3.2	50
5	Acidogenesis, solventogenesis, metabolic stress response and life cycle changes in Clostridium beijerinckii NRRL B-598 at the transcriptomic level. Scientific Reports, 2019, 9, 1371.	3.3	48
6	Comparative analysis of high butanol tolerance and production in clostridia. Biotechnology Advances, 2018, 36, 721-738.	11.7	46
7	Effect of initial pH, different nitrogen sources, and cultivation time on the production of yellow or orange <i>Monascus purpureus</i> pigments and the mycotoxin citrinin. Food Science and Nutrition, 2019, 7, 3494-3500.	3.4	39
8	Evaluation of viability, metabolic activity and spore quantity in clostridial cultures during ABE fermentation. FEMS Microbiology Letters, 2016, 363, fnw031.	1.8	33
9	Continuous production of n-butanol by Clostridium pasteurianum DSM 525 using suspended and surface-immobilized cells. Journal of Biotechnology, 2015, 216, 29-35.	3.8	32
10	Chicken feather and wheat straw hydrolysate for direct utilization in biobutanol production. Renewable Energy, 2020, 145, 1941-1948.	8.9	31
11	Flow cytometry analysis of Clostridium beijerinckii NRRL B-598 populations exhibiting different phenotypes induced by changes in cultivation conditions. Biotechnology for Biofuels, 2018, 11, 99.	6.2	29
12	Complete genome sequence of Clostridium pasteurianum NRRL B-598, a non-type strain producing butanol. Journal of Biotechnology, 2015, 214, 113-114.	3.8	24
13	Transcriptional analysis of amino acid, metal ion, vitamin and carbohydrate uptake in butanol-producing Clostridium beijerinckii NRRL B-598. PLoS ONE, 2019, 14, e0224560.	2.5	19
14	A transcriptional response of Clostridium beijerinckii NRRL B-598 to a butanol shock. Biotechnology for Biofuels, 2019, 12, 243.	6.2	18
15	Transcription profiling of butanol producer Clostridium beijerinckii NRRL B-598 using RNA-Seq. BMC Genomics, 2018, 19, 415.	2.8	17
16	Rapid flow cytometric method for viability determination of solventogenic clostridia. Folia Microbiologica, 2012, 57, 307-311.	2.3	16
17	Use of fluorescent staining and flow cytometry for monitoring physiological changes in solventogenic clostridia. Anaerobe, 2014, 29, 113-117.	2.1	13
18	Comparison of expression of key sporulation, solventogenic and acetogenic genes in C. beijerinckii NRRL B-598 and its mutant strain overexpressing spoOA. Applied Microbiology and Biotechnology, 2017, 101, 8279-8291.	3.6	12

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19	Use of wheat straw and chicken feather hydrolysates as a complete medium for lactic acid production. Czech Journal of Food Sciences, 2018, 36, 146-153.	1.2	11
20	Effective continuous acetone–butanol–ethanol production with full utilization of cassava by immobilized symbiotic TSH06. Biotechnology for Biofuels, 2019, 12, 219.	6.2	10
21	Phenotypic and genomic analysis of isopropanol and 1,3-propanediol producer Clostridium diolis DSM 15410. Genomics, 2021, 113, 1109-1119.	2.9	9
22	Comparison of Lactic Acid Production by L. casei in Batch, Fed-batch and Continuous Cultivation, Testing the use of Feather Hydrolysate as a Complex Nitrogen Source. Brazilian Archives of Biology and Technology, 0, 63, .	0.5	9
23	Phenotypic and Genomic Analysis of Clostridium beijerinckii NRRL B-598 Mutants With Increased Butanol Tolerance. Frontiers in Bioengineering and Biotechnology, 2020, 8, 598392.	4.1	6
24	Transcriptomic studies of solventogenic clostridia, Clostridium acetobutylicum and Clostridium beijerinckii. Biotechnology Advances, 2022, 58, 107889.	11.7	6
25	Diversity and Evolution of Clostridium beijerinckii and Complete Genome of the Type Strain DSM 791T. Processes, 2021, 9, 1196.	2.8	5
26	Deeper below the surfaceâ€"transcriptional changes in selected genes of Clostridium beijerinckii in response to butanol shock. MicrobiologyOpen, 2021, 10, e1146.	3.0	5
27	Application of Flow Cytometry to Saccharomyces cerevisiae Population Analysis. Chimia, 2005, 59, 745-748.	0.6	4
28	Production and cleavage of a fusion protein of porcine trypsinogen and enhanced green fluorescent protein (EGFP) in Pichia pastoris. Folia Microbiologica, 2018, 63, 773-787.	2.3	4
29	Identification and Validation of Reference Genes in Clostridium beijerinckii NRRL B-598 for RT-qPCR Using RNA-Seq Data. Frontiers in Microbiology, 2021, 12, 640054.	3.5	4
30	Changes in efflux pump activity of Clostridium beijerinckii throughout ABE fermentation. Applied Microbiology and Biotechnology, 2021, 105, 877-889.	3.6	3
31	Effect of a Monascus sp. Red Yeast Rice Extract on Germination of Bacterial Spores. Frontiers in Microbiology, 2021, 12, 686100.	3.5	2
32	Thermophilic bacteria colony growth and its consequences in the food industry. Czech Journal of Food Sciences, 2004, 22, 1-8.	1.2	1
33	Microbial production of butanol from food industry waste. , 2020, , 163-180.		1
34	Monascus Secondary Metabolites. , 2015, , 1-31.		0
35	Title is missing!. , 2019, 14, e0224560.		0
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38	8 Title is missing!. , 2019, 14, e0224560.		0