

Manuel Becerra

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7618907/publications.pdf>

Version: 2024-02-01

64
papers

1,354
citations

331259

21
h-index

360668

35
g-index

64
all docs

64
docs citations

64
times ranked

1701
citing authors

#	ARTICLE	IF	CITATIONS
1	Bioprospecting for Thermozyms and Characterization of a Novel Lipolytic Thermozyne Belonging to the SGNH/GDSL Family of Hydrolases. International Journal of Molecular Sciences, 2022, 23, 5733.	1.8	1
2	Characterization of a novel thermophilic metagenomic GH5 endoglucanase heterologously expressed in Escherichia coli and Saccharomyces cerevisiae. , 2022, 15, .		4
3	Biochemical and Structural Characterization of a novel thermophilic esterase EstD11 provide catalytic insights for the HSL family. Computational and Structural Biotechnology Journal, 2021, 19, 1214-1232.	1.9	17
4	Hot Springs Thermophilic Microbiomes. , 2021, , 87-105.		0
5	The HMGB Protein Klxr1, a DNA Binding Regulator of Kluyveromyces lactis Gene Expression Involved in Oxidative Metabolism, Growth, and dNTP Synthesis. Biomolecules, 2021, 11, 1392.	1.8	2
6	Biovalorization of cheese whey and molasses wastes to galactosidases by recombinant yeasts. , 2020, , 149-161.		6
7	Microbial diversity analysis and screening for novel xylanase enzymes from the sediment of the Lobios Hot Spring in Spain. Scientific Reports, 2019, 9, 11195.	1.6	37
8	Optimization of Saccharomyces cerevisiae β -galactosidase production and application in the degradation of raffinose family oligosaccharides. Microbial Cell Factories, 2019, 18, 172.	1.9	20
9	Bioconversion of Beet Molasses to Alpha-Galactosidase and Ethanol. Frontiers in Microbiology, 2019, 10, 405.	1.5	22
10	Heat-Loving β -Galactosidases from Cultured and Uncultured Microorganisms. Current Protein and Peptide Science, 2018, 19, 1224-1234.	0.7	4
11	Valuation of agro-industrial wastes as substrates for heterologous production of β -galactosidase. Microbial Cell Factories, 2018, 17, 137.	1.9	16
12	Advances of Functional Metagenomics in Harnessing Thermozyms. , 2018, , 289-307.		3
13	Cellulases from Thermophiles Found by Metagenomics. Microorganisms, 2018, 6, 66.	1.6	46
14	Structural features of <i>Aspergillus niger</i> β -galactosidase define its activity against glycoside linkages. FEBS Journal, 2017, 284, 1815-1829.	2.2	25
15	Rational mutagenesis by engineering disulphide bonds improves Kluyveromyces lactis beta-galactosidase for high-temperature industrial applications. Scientific Reports, 2017, 7, 45535.	1.6	24
16	Archaeal Biocommunication in Hot Springs Revealed by Metagenomics. , 2017, , 85-101.		0
17	Dual function of lxr1 in transcriptional regulation and recognition of cisplatin-DNA adducts is caused by differential binding through its two HMG-boxes. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2017, 1860, 256-269.	0.9	9
18	HMGB Proteins from Yeast to Human. Gene Regulation, DNA Repair and Beyond. , 2017, , .		2

#	ARTICLE	IF	CITATIONS
19	Extremophilic Esterases for Bioprocessing of Lignocellulosic Feedstocks. , 2017, , 205-223.		0
20	Improved bioethanol production in an engineered <i>Kluyveromyces lactis</i> strain shifted from respiratory to fermentative metabolism by deletion of NDI 1. <i>Microbial Biotechnology</i> , 2015, 8, 319-330.	2.0	15
21	Biobutanol from cheese whey. <i>Microbial Cell Factories</i> , 2015, 14, 27.	1.9	35
22	KlGcr1 controls glucose-6-phosphate dehydrogenase activity and responses to H ₂ O ₂ , cadmium and arsenate in <i>Kluyveromyces lactis</i> . <i>Fungal Genetics and Biology</i> , 2015, 82, 95-103.	0.9	7
23	Sky1 regulates the expression of sulfur metabolism genes in response to cisplatin. <i>Microbiology (United Kingdom)</i> , 2014, 160, 1357-1368.	0.7	6
24	Crystallization and preliminary X-ray diffraction data of Î ² -galactosidase from <i>Aspergillus niger</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2014, 70, 1529-1531.	0.4	4
25	The yeast hypoxic responses, resources for new biotechnological opportunities. <i>Biotechnology Letters</i> , 2012, 34, 2161-2173.	1.1	15
26	Structural basis of specificity in tetrameric <i>Kluyveromyces lactis</i> Î ² -galactosidase. <i>Journal of Structural Biology</i> , 2012, 177, 392-401.	1.3	88
27	lxr1p and the control of the <i>Saccharomyces cerevisiae</i> hypoxic response. <i>Applied Microbiology and Biotechnology</i> , 2012, 94, 173-184.	1.7	22
28	Comparative transcriptome analysis of yeast strains carrying slt2, rlm1, and pop2 deletions. <i>Genome</i> , 2011, 54, 99-109.	0.9	3
29	Two Proteins with Different Functions Are Derived from the <i>KlHEM13</i> Gene. <i>Eukaryotic Cell</i> , 2011, 10, 1331-1339.	3.4	1
30	Crystallization and preliminary X-ray diffraction data of Î [±] -galactosidase from <i>Saccharomyces cerevisiae</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010, 66, 44-47.	0.7	2
31	Crystallization and preliminary X-ray crystallographic analysis of Î ² -galactosidase from <i>Kluyveromyces lactis</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010, 66, 297-300.	0.7	9
32	lxr1p regulates oxygen-dependent <i>KlHEM13</i> transcription. <i>FEMS Yeast Research</i> , 2010, 10, 309-321.	1.1	13
33	Structural Analysis of <i>Saccharomyces cerevisiae</i> Î [±] -Galactosidase and Its Complexes with Natural Substrates Reveals New Insights into Substrate Specificity of GH27 Glycosidases. <i>Journal of Biological Chemistry</i> , 2010, 285, 28020-28033.	1.6	36
34	Growth phase-dependent expression of <i>Kluyveromyces lactis</i> genes and involvement of 3' UTR elements. <i>Process Biochemistry</i> , 2008, 43, 1153-1157.	1.8	2
35	<i>Kluyveromyces lactis</i> Î ² -galactosidase crystallization using full-factorial experimental design. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2008, 52-53, 178-182.	1.8	8
36	Functional motifs outside the kinase domain of yeast Srb10p. Their role in transcriptional regulation and protein-interactions with Tup1p and Srb11p. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2007, 1774, 1227-1235.	1.1	7

#	ARTICLE	IF	CITATIONS
37	An approach to the hypoxic and oxidative stress responses in <i>Kluyveromyces lactis</i> by analysis of mRNA levels. <i>FEMS Yeast Research</i> , 2007, 7, 702-714.	1.1	17
38	Secretion of a hybrid <i>K. lactis</i> - <i>A. niger</i> β -galactosidase. <i>Microbial Cell Factories</i> , 2006, 5, P66.	1.9	0
39	Secretion and properties of a hybrid <i>Kluyveromyces lactis</i> - <i>Aspergillus niger</i> beta-galactosidase. <i>Microbial Cell Factories</i> , 2006, 5, 41.	1.9	33
40	A transcriptome analysis of <i>Kluyveromyces lactis</i> growing in cheese whey. <i>International Dairy Journal</i> , 2006, 16, 207-214.	1.5	11
41	Reoxidation of cytosolic NADPH in <i>Kluyveromyces lactis</i> . <i>FEMS Yeast Research</i> , 2006, 6, 371-380.	1.1	43
42	Functional characterization of KIHEM13, a hypoxic gene of <i>Kluyveromyces lactis</i> . <i>Canadian Journal of Microbiology</i> , 2005, 51, 431-431.	0.8	1
43	Functional characterization of KIHEM13, a hypoxic gene of <i>Kluyveromyces lactis</i> . <i>Canadian Journal of Microbiology</i> , 2005, 51, 241-249.	0.8	11
44	Functional characterisation and transcriptional regulation of the KIHEM12 gene from <i>Kluyveromyces lactis</i> . <i>Current Genetics</i> , 2004, 46, 147-57.	0.8	6
45	Genome-wide analysis of <i>Kluyveromyces lactis</i> in wild-type and <i>rag2</i> mutant strains. <i>Genome</i> , 2004, 47, 970-978.	0.9	21
46	Engineered autolytic yeast strains secreting <i>Kluyveromyces lactis</i> β -galactosidase for production of heterologous proteins in lactose media. <i>Journal of Biotechnology</i> , 2004, 109, 131-137.	1.9	27
47	Genome-Wide Analysis of the Yeast Transcriptome Upon Heat and Cold Shock. <i>Comparative and Functional Genomics</i> , 2003, 4, 366-375.	2.0	18
48	Influence of substituting milk powder for whey powder on yoghurt quality. <i>Trends in Food Science and Technology</i> , 2002, 13, 334-340.	7.8	102
49	Genome-Wide analysis of yeast transcription upon calcium shortage. <i>Cell Calcium</i> , 2002, 32, 83-91.	1.1	9
50	Crystal Structure of the MuSK Tyrosine Kinase. <i>Structure</i> , 2002, 10, 1187-1196.	1.6	122
51	The yeast transcriptome in aerobic and hypoxic conditions: effects of <i>hap1</i> , <i>rox1</i> , <i>rox3</i> and <i>srb10</i> deletions. <i>Molecular Microbiology</i> , 2002, 43, 545-555.	1.2	77
52	The yeast transcriptome in aerobic and hypoxic conditions: effects of <i>hap1</i> , <i>rox1</i> , <i>rox3</i> and <i>srb10</i> deletions. <i>Molecular Microbiology</i> , 2002, 45, 265-265.	1.2	2
53	Metabolic engineering for direct lactose utilization by <i>Saccharomyces cerevisiae</i> . <i>Biotechnology Letters</i> , 2002, 24, 1391-1396.	1.1	10
54	Lactose bioconversion by calcium-alginate immobilization of <i>Kluyveromyces lactis</i> cells. <i>Enzyme and Microbial Technology</i> , 2001, 29, 506-512.	1.6	51

#	ARTICLE	IF	CITATIONS
55	Title is missing!. Biotechnology Letters, 2001, 23, 33-40.	1.1	13
56	New secretory strategies for <i>Kluyveromyces lactis</i> β -galactosidase. Protein Engineering, Design and Selection, 2001, 14, 379-386.	1.0	39
57	Title is missing!. Biotechnology Letters, 1998, 12, 253-256.	0.5	40
58	Dealing with different methods for <i>Kluyveromyces lactis</i> β -galactosidase purification. Biological Procedures Online, 1998, 1, 48-58.	1.4	19
59	Heterologous <i>Kluyveromyces lactis</i> β -galactosidase production and release by <i>Saccharomyces cerevisiae</i> osmotic-remedial thermosensitive autolytic mutants. Biochimica Et Biophysica Acta - General Subjects, 1997, 1335, 235-241.	1.1	27
60	Enzyme encapsulation on chitosan microbeads. Process Biochemistry, 1997, 32, 211-216.	1.8	69
61	Expression of a low-molecular-weight (10 kDa) calcium binding protein in glial cells of the brain of the trout (Teleostei). Anatomy and Embryology, 1997, 196, 403-416.	1.5	39
62	Sequence analysis of a 10 kb DNA fragment from yeast chromosome VII reveals a novel member of the dnaJ family. , 1996, 12, 145-148.		4
63	Yeast β -galactosidase in solid-state fermentations. Enzyme and Microbial Technology, 1996, 19, 39-44.	1.6	32
64	Utilizaci3n de screencasts para un aprendizaje activo. , 0, , 15-24.		0