MarÃ-a Isabel GonzÃ;lez-Siso

List of Publications by Year in descending order

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105 papers

3,111 citations

30 h-index 52 g-index

106 all docs

106 docs citations

106 times ranked 3219 citing authors

#	Article	IF	CITATIONS
1	The biotechnological utilization of cheese whey: A review. Bioresource Technology, 1996, 57, 1-11.	9.6	587
2	New Extremophilic Lipases and Esterases from Metagenomics. Current Protein and Peptide Science, 2014, 15, 445-455.	1.4	144
3	In vitro effects of mangiferin on superoxide concentrations and expression of the inducible nitric oxide synthase, tumour necrosis factor- \hat{l} ± and transforming growth factor- \hat{l} 2 genes. Biochemical Pharmacology, 2003, 65, 1361-1371.	4.4	140
4	Metagenomics of Thermophiles with a Focus on Discovery of Novel Thermozymes. Frontiers in Microbiology, 2016, 7, 1521.	3 . 5	98
5	Structural basis of specificity in tetrameric Kluyveromyces lactis \hat{l}^2 -galactosidase. Journal of Structural Biology, 2012, 177, 392-401.	2.8	88
6	Respirofermentative metabolism in Kluyveromyces lactis:. Enzyme and Microbial Technology, 2000, 26, 699-705.	3.2	81
7	Sugar metabolism, redox balance and oxidative stress response in the respiratory yeast Kluyveromyces lactis. Microbial Cell Factories, 2009, 8, 46.	4.0	75
8	Codon usage in Kluyveromyces lactis and in yeast cytochrome c-encoding genes. Gene, 1994, 139, 43-49.	2.2	71
9	Enzyme encapsulation on chitosan microbeads. Process Biochemistry, 1997, 32, 211-216.	3.7	69
10	Hot Spring Metagenomics. Life, 2013, 3, 308-320.	2.4	69
10	Hot Spring Metagenomics. Life, 2013, 3, 308-320. Respirofermentative metabolism in Kluyveromyces lactis: Ethanol production and the Crabtree effect. Enzyme and Microbial Technology, 1996, 18, 585-591.	3.2	59
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11	Respirofermentative metabolism in Kluyveromyces lactis: Ethanol production and the Crabtree effect. Enzyme and Microbial Technology, 1996, 18, 585-591. Transcriptome analysis of the thermotolerant yeast Kluyveromyces marxianus CCT 7735 under ethanol	3.2	59
11 12	Respirofermentative metabolism in Kluyveromyces lactis: Ethanol production and the Crabtree effect. Enzyme and Microbial Technology, 1996, 18, 585-591. Transcriptome analysis of the thermotolerant yeast Kluyveromyces marxianus CCT 7735 under ethanol stress. Applied Microbiology and Biotechnology, 2017, 101, 6969-6980. Metagenomics of an Alkaline Hot Spring in Galicia (Spain): Microbial Diversity Analysis and Screening	3.2	59 57
11 12 13	Respirofermentative metabolism in Kluyveromyces lactis: Ethanol production and the Crabtree effect. Enzyme and Microbial Technology, 1996, 18, 585-591. Transcriptome analysis of the thermotolerant yeast Kluyveromyces marxianus CCT 7735 under ethanol stress. Applied Microbiology and Biotechnology, 2017, 101, 6969-6980. Metagenomics of an Alkaline Hot Spring in Galicia (Spain): Microbial Diversity Analysis and Screening for Novel Lipolytic Enzymes. Frontiers in Microbiology, 2015, 6, 1291. Lactose bioconversion by calcium-alginate immobilization of Kluyveromyces lactis cells. Enzyme and	3.2 3.6 3.5	59 57 54
11 12 13	Respirofermentative metabolism in Kluyveromyces lactis: Ethanol production and the Crabtree effect. Enzyme and Microbial Technology, 1996, 18, 585-591. Transcriptome analysis of the thermotolerant yeast Kluyveromyces marxianus CCT 7735 under ethanol stress. Applied Microbiology and Biotechnology, 2017, 101, 6969-6980. Metagenomics of an Alkaline Hot Spring in Galicia (Spain): Microbial Diversity Analysis and Screening for Novel Lipolytic Enzymes. Frontiers in Microbiology, 2015, 6, 1291. Lactose bioconversion by calcium-alginate immobilization of Kluyveromyces lactis cells. Enzyme and Microbial Technology, 2001, 29, 506-512. Kluyveromyces marxianus as a host for heterologous protein synthesis. Applied Microbiology and	3.2 3.6 3.5	59 57 54 51
11 12 13 14	Respirofermentative metabolism in Kluyveromyces lactis: Ethanol production and the Crabtree effect. Enzyme and Microbial Technology, 1996, 18, 585-591. Transcriptome analysis of the thermotolerant yeast Kluyveromyces marxianus CCT 7735 under ethanol stress. Applied Microbiology and Biotechnology, 2017, 101, 6969-6980. Metagenomics of an Alkaline Hot Spring in Galicia (Spain): Microbial Diversity Analysis and Screening for Novel Lipolytic Enzymes. Frontiers in Microbiology, 2015, 6, 1291. Lactose bioconversion by calcium-alginate immobilization of Kluyveromyces lactis cells. Enzyme and Microbial Technology, 2001, 29, 506-512. Kluyveromyces marxianus as a host for heterologous protein synthesis. Applied Microbiology and Biotechnology, 2016, 100, 6193-6208.	3.2 3.6 3.2 3.6	59 57 54 51 49

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19	Reoxidation of the NADPH produced by the pentose phosphate pathway is necessary for the utilization of glucose byKluyveromyces lactis rag2mutants. FEBS Letters, 1996, 387, 7-10.	2.8	41
20	Title is missing!. Biotechnology Letters, 1998, 12, 253-256.	0.5	40
21	Heterologous expression of glucose oxidase in the yeast Kluyveromyces marxianus. Microbial Cell Factories, 2010, 9, 4.	4.0	40
22	New secretory strategies for Kluyveromyces lactis \hat{l}^2 -galactosidase. Protein Engineering, Design and Selection, 2001, 14, 379-386.	2.1	39
23	Microbial diversity analysis and screening for novel xylanase enzymes from the sediment of the Lobios Hot Spring in Spain. Scientific Reports, 2019, 9, 11195.	3.3	37
24	Depuration and valuation of mussel-processing wastes. Characterization of amylolytic postincubates from different species grown on an effluent. Bioresource Technology, 1992, 42, 133-140.	9.6	35
25	Kluyveromyces lactis immobilization on corn grits for milk whey lactose hydrolysis. Enzyme and Microbial Technology, 1994, 16, 303-310.	3.2	35
26	Biobutanol from cheese whey. Microbial Cell Factories, 2015, 14, 27.	4.0	35
27	Heterologous expression of a thermophilic esterase in Kluyveromyces yeasts. Applied Microbiology and Biotechnology, 2011, 89, 375-385.	3. 6	34
28	Secretion and properties of a hybrid Kluyveromyces lactis-Aspergillus niger beta-galactosidase. Microbial Cell Factories, 2006, 5, 41.	4.0	33
29	Yeast \hat{l}^2 -galactosidase in solid-state fermentations. Enzyme and Microbial Technology, 1996, 19, 39-44.	3.2	32
30	The nuclear genes encoding the internal (KINDI1) and external (KINDE1) alternative NAD(P)H:ubiquinone oxidoreductases of mitochondria from Kluyveromyces lactis. Biochimica Et Biophysica Acta - Bioenergetics, 2005, 1707, 199-210.	1.0	31
31	Permeabilization of Kluyveromyces lactis cells for milk whey saccharification: A comparison of different treatments. Biotechnology Letters, 1992, 6, 289-292.	0.5	29
32	Covalent immobilization of \hat{l}^2 -galactosidase on corn grits. A system for lactose hydrolysis without diffusional resistance. Process Biochemistry, 1994, 29, 7-12.	3.7	27
33	Heterologous Kluyveromyces lactis \hat{l}^2 -galactosidase production and release by Saccharomyces cerevisiae osmotic-remedial thermosensitive autolytic mutants. Biochimica Et Biophysica Acta - General Subjects, 1997, 1335, 235-241.	2.4	27
34	Engineered autolytic yeast strains secreting Kluyveromyces lactis \hat{l}^2 -galactosidase for production of heterologous proteins in lactose media. Journal of Biotechnology, 2004, 109, 131-137.	3.8	27
35	Heterologous expression of an esterase from Thermus thermophilus HB27 in Saccharomyces cerevisiae. Journal of Biotechnology, 2010, 145, 226-232.	3.8	25
36	Characterization of the second external alternative dehydrogenase from mitochondria of the respiratory yeast Kluyveromyces lactis. Biochimica Et Biophysica Acta - Bioenergetics, 2006, 1757, 1476-1484.	1.0	24

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37	Rational mutagenesis by engineering disulphide bonds improves Kluyveromyces lactis beta-galactosidase for high-temperature industrial applications. Scientific Reports, 2017, 7, 45535.	3.3	24
38	\hat{l}^2 -Galactosidase production by Kluyveromyces lactis on milk whey: batch versus fed-batch cultures. Process Biochemistry, 1994, 29, 565-568.	3.7	22
39	lxr1p and the control of the Saccharomyces cerevisiae hypoxic response. Applied Microbiology and Biotechnology, 2012, 94, 173-184.	3.6	22
40	Bioconversion of Beet Molasses to Alpha-Galactosidase and Ethanol. Frontiers in Microbiology, 2019, 10, 405.	3.5	22
41	Enhancement of the bioproduction potential of an amylaceous effluent. Bioresource Technology, 1993, 44, 155-163.	9.6	21
42	Genome-wide analysis of Kluyveromyces lactis in wild-type and rag2 mutant strains. Genome, 2004, 47, 970-978.	2.0	21
43	Regulatory factors controlling transcription of $\langle i \rangle$ Saccharomyces cerevisiae IXR1 $\langle i \rangle$ by oxygen levels: a model of transcriptional adaptation from aerobiosis to hypoxia implicating $\langle i \rangle$ ROX1 $\langle i \rangle$ and $\langle i \rangle$ IXR1 $\langle i \rangle$ cross-regulation. Biochemical Journal, 2010, 425, 235-243.	3.7	20
44	Optimization of Saccharomyces cerevisiae \hat{l} ±-galactosidase production and application in the degradation of raffinose family oligosaccharides. Microbial Cell Factories, 2019, 18, 172.	4.0	20
45	Dealing with different methods for Kluyveromyces lactis \hat{l}^2 -galactosidase purification. Biological Procedures Online, 1998, 1, 48-58.	2.9	19
46	Genome-Wide Analysis of the Yeast Transcriptome Upon Heat and Cold Shock. Comparative and Functional Genomics, 2003, 4, 366-375.	2.0	18
47	Functional characterization of KlHAP1: A model to foresee different mechanisms of transcriptional regulation by Hap1p in yeasts. Gene, 2007, 405, 96-107.	2.2	18
48	The role of glutathione reductase in the interplay between oxidative stress response and turnover of cytosolic NADPH in Kluyveromyces lactis. FEMS Yeast Research, 2008, 8, 597-606.	2.3	18
49	An approach to the hypoxic and oxidative stress responses inKluyveromyces lactisby analysis of mRNA levels. FEMS Yeast Research, 2007, 7, 702-714.	2.3	17
50	Production and characterization of two N-terminal truncated esterases from Thermus thermophilus HB27 in a mesophilic yeast: Effect of N-terminus in thermal activity and stability. Protein Expression and Purification, 2011, 78, 120-130.	1.3	17
51	Cloning, expression, purification and characterization of an oligomeric His-tagged thermophilic esterase from Thermus thermophilus HB27. Process Biochemistry, 2014, 49, 927-935.	3.7	17
52	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.	4.1	17
53	Valuation of agro-industrial wastes as substrates for heterologous production of α-galactosidase. Microbial Cell Factories, 2018, 17, 137.	4.0	16
54	RFLP analysis of PCR-amplified small subunit ribosomal DNA of three fish microsporidian species. Parasitology, 2000, 120, 113-119.	1.5	15

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55	The yeast hypoxic responses, resources for new biotechnological opportunities. Biotechnology Letters, 2012, 34, 2161-2173.	2,2	15
56	<i>Kluyveromyces lactis</i> : A Suitable Yeast Model to Study Cellular Defense Mechanisms against Hypoxia-Induced Oxidative Stress. Oxidative Medicine and Cellular Longevity, 2012, 2012, 1-14.	4.0	15
57	Improved bioethanol production in an engineered K luyveromyces lactis strain shifted from respiratory to fermentative metabolism by deletion of NDI 1. Microbial Biotechnology, 2015, 8, 319-330.	4.2	15
58	Isolation and characterization of two nuclear genes encoding glutathione and thioredoxin reductases from the yeast Kluyveromyces lactis. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2004, 1678, 170-175.	2.4	14
59	Title is missing!. Biotechnology Letters, 2001, 23, 33-40.	2.2	13
60	lxr1p regulates oxygen-dependent â€fHEM13â€ftranscription. FEMS Yeast Research, 2010, 10, 309-321.	2.3	13
61	Thermus thermophilus as a Source of Thermostable Lipolytic Enzymes. Microorganisms, 2015, 3, 792-808.	3 . 6	13
62	Production and Characterization of an Extracellular Acid Protease from Thermophilic Brevibacillus sp. OA30 Isolated from an Algerian Hot Spring. Microorganisms, 2018, 6, 31.	3.6	13
63	Proteomic Analysis of the Oxidative Stress Response inKluyveromyces lactisand Effect of Glutathione Reductase Depletion. Journal of Proteome Research, 2010, 9, 2358-2376.	3.7	12
64	Functional characterization of KlHEM13, a hypoxic gene of Kluyveromyces lactis. Canadian Journal of Microbiology, 2005, 51, 241-249.	1.7	11
65	A transcriptome analysis of Kluyveromyces lactis growing in cheese whey. International Dairy Journal, 2006, 16, 207-214.	3.0	11
66	lxr1 Regulates Ribosomal Gene Transcription and Yeast Response to Cisplatin. Scientific Reports, 2018, 8, 3090.	3.3	11
67	Exploring the taxonomical and functional profile of As Burgas hot spring focusing on thermostable \hat{l}^2 -galactosidases. Scientific Reports, 2021, 11, 101.	3.3	11
68	Amylolysis in systems with ?-amylase and glucoamylase. A comparative study of six procedures of evaluation. Biotechnology Letters, 1987, 1, 195.	0.5	10
69	A HPLC method for specific determination ofl±-amylase and glucoamylase in complex enzymatic preparations. Chromatographia, 1989, 27, 328-332.	1.3	10
70	Metabolic engineering for direct lactose utilization by Saccharomyces cerevisiae. Biotechnology Letters, 2002, 24, 1391-1396.	2.2	10
71	Effects of chitinolytic and proteolytic enzymes on in vitro phagocytosis of microsporidians by spleen macrophages of turbot, Scophthalmus maximus L Veterinary Immunology and Immunopathology, 1997, 59, 171-180.	1.2	9
72	Crystallization and preliminary X-ray crystallographic analysis of \hat{l}^2 -galactosidase from <i>Kluyveromyces lactis </i> . Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 297-300.	0.7	9

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73	Microfungus-yeast mixed cultures in the degradation of amylaceous wastes. I: Interactions affecting amylolytic activity. Biotechnology Letters, 1988, 10, 431-436.	2.2	8
74	DNA probes for detection of the fish microsporidians Microgemma caulleryi and Tetramicra brevifilum. Parasitology, 1999, 119, 267-272.	1.5	8
75	Cloning Genes From a Library Using a Clustering Strategy and PCR. Molecular Biotechnology, 2004, 26, 35-38.	2.4	8
76	Kluyveromyces lactis \hat{l}^2 -galactosidase crystallization using full-factorial experimental design. Journal of Molecular Catalysis B: Enzymatic, 2008, 52-53, 178-182.	1.8	8
77	PICDI, a simple program for codon bias calculation. Molecular Biotechnology, 1996, 5, 191-195.	2.4	7
78	Functional motifs outside the kinase domain of yeast Srb10p. Their role in transcriptional regulation and protein-interactions with $Tup1p$ and $Srb11p$. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2007, 1774, 1227-1235.	2.3	7
79	KlGcr1 controls glucose-6-phosphate dehydrogenase activity and responses to H2O2, cadmium and arsenate in Kluyveromyces lactis. Fungal Genetics and Biology, 2015, 82, 95-103.	2.1	7
80	A factorial experimental design for investigation of the effects of temperature, incubation time, and pathogen-to-phagocyte ratio on in vitro phagocytosis by turbot adherent cells. Comparative Biochemistry and Physiology C, Comparative Pharmacology and Toxicology, 1995, 112, 215-220.	0.5	6
81	Functional characterisation and transcriptional regulation of the KIHEM12 gene from Kluyveromyces lactis. Current Genetics, 2004, 46, 147-57.	1.7	6
82	A functional analysis of <i>Kluyveromyces lactis</i> glutathione reductase. Yeast, 2010, 27, 431-441.	1.7	6
83	Sky1 regulates the expression of sulfur metabolism genes in response to cisplatin. Microbiology (United Kingdom), 2014, 160, 1357-1368.	1.8	6
84	Isolation and transcriptional regulation of the Kluyveromyces lactis FBA1 (fructose-1,6-bisphosphate) Tj ETQq0 0 () rgBJ /Ove	erlgck 10 Tf 5
85	SKY1 and IXR1 interactions, their effects on cisplatin and spermine resistance in Saccharomyces cerevisiae. Canadian Journal of Microbiology, 2012, 58, 184-188.	1.7	5
86	Editorial for the Special Issue: Thermophiles and Thermozymes. Microorganisms, 2019, 7, 62.	3.6	5
87	Microfungus-yeast mixed cultures in the degradation of amylaceous wastes. II: An experimental design for optimization of yeast production. Biotechnology Letters, 1988, 2, 171-176.	0.5	4
88	Chromosomal mapping of the KICYC1 gene from Kluyveromyces lactis. Genome, 1994, 37, 515-517.	2.0	4
89	A simple form of immobilisation and its effects on morphologic trends and metabolic activity of pellet-forming microfungi. Bioresource Technology, 1994, 48, 237-243.	9.6	4
90	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

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91	Heat-Loving \hat{l}^2 -Galactosidases from Cultured and Uncultured Microorganisms. Current Protein and Peptide Science, 2018, 19, 1224-1234.	1.4	4
92	Characterization of a novel thermophilic metagenomic GH5 endoglucanase heterologously expressed in Escherichia coli and Saccharomyces cerevisiae. , 2022, 15 , .		4
93	Proteomic Analyses Reveal that Sky1 Modulates Apoptosis and Mitophagy in Saccharomyces cerevisiae Cells Exposed to Cisplatin. International Journal of Molecular Sciences, 2014, 15, 12573-12590.	4.1	3
94	Advances of Functional Metagenomics in Harnessing Thermozymes., 2018,, 289-307.		3
95	Structural determination of Enzyme-Graphene Nanocomposite Sensor Material. Scientific Reports, 2019, 9, 15519.	3.3	3
96	The Kluyveromyces lactis gene KlGSK-3 combines functions which in Saccharomyces cerevisiae are performed by MCK1 and MSD1. Current Genetics, 1998, 33, 262-267.	1.7	2
97	Mouse antibody response to a microsporidian parasite following inoculation with a gene coding for parasite ribosomal RNA. Vaccine, 2002, 20, 2648-2655.	3.8	2
98	A functional analysis of <i>KlSRB10</i> : implications in <i>Kluyveromyces lactis</i> transcriptional regulation. Yeast, 2007, 24, 1061-1073.	1.7	2
99	Functional characterization of KIHEM13, a hypoxic gene of Kluyveromyces lactis. Canadian Journal of Microbiology, 2005, 51, 431-431.	1.7	1
100	Two Proteins with Different Functions Are Derived from the <i>KlHEM13</i> Gene. Eukaryotic Cell, 2011, 10, 1331-1339.	3.4	1
101	Reactivity of a Recombinant Esterase from Thermus thermophilus HB27 in Aqueous and Organic Media. Microorganisms, 2022, 10, 915.	3.6	1
102	Bioprospecting for Thermozymes and Characterization of a Novel Lipolytic Thermozyme Belonging to the SGNH/GDSL Family of Hydrolases. International Journal of Molecular Sciences, 2022, 23, 5733.	4.1	1
103	Heterologous Aspergillus niger \hat{l}^2 -galactosidase secretion by Saccharomyces cerevisiae. Journal of Biotechnology, 2007, 131, S199-S200.	3.8	O
104	Archaeal Biocommunication in Hot Springs Revealed by Metagenomics., 2017,, 85-101.		0
105	Extremophilic Esterases for Bioprocessing of Lignocellulosic Feedstocks., 2017,, 205-223.		0