

Jose Luis Garcia

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

163 papers	8,019 citations	49 h-index	85 g-index
169 ext. papers	9,072 ext. citations	6.1 avg, IF	5.6 L-index

#	Paper	IF	Citations
163	Genomic sequence of the pathogenic and allergenic filamentous fungus <i>Aspergillus fumigatus</i> . <i>Nature</i> , 2005 , 438, 1151-6	50.4	1114
162	Genomic analysis of the aromatic catabolic pathways from <i>Pseudomonas putida</i> KT2440. <i>Environmental Microbiology</i> , 2002 , 4, 824-41	5.2	380
161	Anaerobic catabolism of aromatic compounds: a genetic and genomic view. <i>Microbiology and Molecular Biology Reviews</i> , 2009 , 73, 71-133	13.2	312
160	The homogentisate pathway: a central catabolic pathway involved in the degradation of L-phenylalanine, L-tyrosine, and 3-hydroxyphenylacetate in <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 2004 , 186, 5062-77	3.5	190
159	Taking aim on bacterial pathogens: from phage therapy to enzybiotics. <i>Current Opinion in Microbiology</i> , 2007 , 10, 461-72	7.9	175
158	Microalgae, old sustainable food and fashion nutraceuticals. <i>Microbial Biotechnology</i> , 2017 , 10, 1017-1024	4.3	171
157	Structural basis for selective recognition of pneumococcal cell wall by modular endolysin from phage Cp-1. <i>Structure</i> , 2003 , 11, 1239-49	5.2	135
156	The phenylacetyl-CoA catabolon: a complex catabolic unit with broad biotechnological applications. <i>Molecular Microbiology</i> , 2001 , 39, 1434-42	4.1	132
155	Genome sequence of the olive tree, <i>Olea europaea</i> . <i>GigaScience</i> , 2016 , 5, 29	7.6	130
154	LytB, a novel pneumococcal murein hydrolase essential for cell separation. <i>Molecular Microbiology</i> , 1999 , 31, 1275-81	4.1	126
153	Genome and transcriptome analysis of the Mesoamerican common bean and the role of gene duplications in establishing tissue and temporal specialization of genes. <i>Genome Biology</i> , 2016 , 17, 32	18.3	124
152	Searching for autolysin functions. Characterization of a pneumococcal mutant deleted in the <i>lytA</i> gene. <i>FEBS Journal</i> , 1986 , 158, 289-93		115
151	Deciphering the genetic determinants for aerobic nicotinic acid degradation: the <i>nic</i> cluster from <i>Pseudomonas putida</i> KT2440. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 11329-34	11.5	112
150	The molecular characterization of the first autolytic lysozyme of <i>Streptococcus pneumoniae</i> reveals evolutionary mobile domains. <i>Molecular Microbiology</i> , 1999 , 33, 128-38	4.1	109
149	Whole genome sequencing of turbot (<i>Scophthalmus maximus</i> ; Pleuronectiformes): a fish adapted to demersal life. <i>DNA Research</i> , 2016 , 23, 181-92	4.5	103
148	Catabolism and biotechnological applications of cholesterol degrading bacteria. <i>Microbial Biotechnology</i> , 2012 , 5, 679-99	6.3	99
147	Purification and polar localization of pneumococcal LytB, a putative endo-beta-N-acetylglucosaminidase: the chain-dispersing murein hydrolase. <i>Journal of Bacteriology</i> , 2002 , 184, 4988-5000	3.5	97

146	Stabilization of penicillin G acylase from <i>Escherichia coli</i> : site-directed mutagenesis of the protein surface to increase multipoint covalent attachment. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 1249-51	4.8	93
145	The bzd gene cluster, coding for anaerobic benzoate catabolism, in <i>Azoarcus</i> sp. strain CIB. <i>Journal of Bacteriology</i> , 2004 , 186, 5762-74	3.5	92
144	One-step purification, covalent immobilization, and additional stabilization of poly-His-tagged proteins using novel heterofunctional chelate-epoxy supports. <i>Biotechnology and Bioengineering</i> , 2001 , 76, 269-76	4.9	91
143	The turnover of medium-chain-length polyhydroxyalkanoates in <i>Pseudomonas putida</i> KT2442 and the fundamental role of PhaZ depolymerase for the metabolic balance. <i>Environmental Microbiology</i> , 2010 , 12, 207-21	5.2	87
142	Extreme genomic erosion after recurrent demographic bottlenecks in the highly endangered Iberian lynx. <i>Genome Biology</i> , 2016 , 17, 251	18.3	85
141	Functionalization of Gold Surfaces for Specific and Reversible Attachment of a Fused β -Galactosidase and Choline-Receptor Protein. <i>Journal of the American Chemical Society</i> , 1997 , 119, 1043-1051	16.4	84
140	Molecular characterization of the safracin biosynthetic pathway from <i>Pseudomonas fluorescens</i> A2-2: designing new cytotoxic compounds. <i>Molecular Microbiology</i> , 2005 , 56, 144-54	4.1	84
139	In vivo immobilization of fusion proteins on bioplastics by the novel tag BioF. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 3205-12	4.8	82
138	Biological role of the pneumococcal amidase. Cloning of the <i>lytA</i> gene in <i>Streptococcus pneumoniae</i> . <i>FEBS Journal</i> , 1987 , 164, 621-4		80
137	Immobilization and single-step purification of fusion proteins using DEAE-cellulose. <i>FEBS Journal</i> , 1992 , 203, 153-9		79
136	Conformational selection of glycomimetics at enzyme catalytic sites: experimental demonstration of the binding of distinct high-energy distorted conformations of C-, S-, and O-glycosides by <i>E. Coli</i> beta-galactosidases. <i>Journal of the American Chemical Society</i> , 2002 , 124, 4804-10	16.4	78
135	One-step purification, covalent immobilization, and additional stabilization of a thermophilic poly-His-tagged beta-galactosidase from <i>Thermus</i> sp. strain T2 by using novel heterofunctional chelate-epoxy Sepabeads. <i>Biomacromolecules</i> , 2003 , 4, 107-13	6.9	76
134	Insights into pneumococcal pathogenesis from the crystal structure of the modular teichoic acid phosphorylcholine esterase Pce. <i>Nature Structural and Molecular Biology</i> , 2005 , 12, 533-8	17.6	75
133	Biotransformations catalyzed by multimeric enzymes: stabilization of tetrameric ampicillin acylase permits the optimization of ampicillin synthesis under dissociation conditions. <i>Biomacromolecules</i> , 2001 , 2, 95-104	6.9	73
132	Enhancing desulphurization by engineering a flavin reductase-encoding gene cassette in recombinant biocatalysts. <i>Environmental Microbiology</i> , 2000 , 2, 687-94	5.2	72
131	Disruption of β -oxidation pathway in <i>Pseudomonas putida</i> KT2442 to produce new functionalized PHAs with thioester groups. <i>Applied Microbiology and Biotechnology</i> , 2011 , 89, 1583-98	5.7	71
130	Structural requirements of choline derivatives for conversion of pneumococcal amidase. A new single-step procedure for purification of this autolysin. <i>FEBS Letters</i> , 1988 , 232, 308-12	3.8	71
129	Affinity chromatography of polyhistidine tagged enzymes. New dextran-coated immobilized metal ion affinity chromatography matrices for prevention of undesired multipoint adsorptions. <i>Journal of Chromatography A</i> , 2001 , 915, 97-106	4.5	68

128	Stabilization of heterodimeric enzyme by multipoint covalent immobilization: Penicillin G acylase from <i>Kluyvera citrophila</i> . <i>Biotechnology and Bioengineering</i> , 1993 , 42, 455-64	4.9	68
127	Bacterial degradation of benzoate: cross-regulation between aerobic and anaerobic pathways. <i>Journal of Biological Chemistry</i> , 2012 , 287, 10494-10508	5.4	66
126	Genetically engineered <i>Pseudomonas</i> : a factory of new bioplastics with broad applications. <i>Environmental Microbiology</i> , 2001 , 3, 612-8	5.2	66
125	Cholesterol metabolism in <i>Mycobacterium smegmatis</i> . <i>Environmental Microbiology Reports</i> , 2012 , 4, 168-82	5.7	64
124	Controlled autolysis facilitates the polyhydroxyalkanoate recovery in <i>Pseudomonas putida</i> KT2440. <i>Microbial Biotechnology</i> , 2011 , 4, 533-47	6.3	64
123	Engineering synthetic bacterial consortia for enhanced desulfurization and revalorization of oil sulfur compounds. <i>Metabolic Engineering</i> , 2016 , 35, 46-54	9.7	58
122	Unravelling the gallic acid degradation pathway in bacteria: the gal cluster from <i>Pseudomonas putida</i> . <i>Molecular Microbiology</i> , 2011 , 79, 359-74	4.1	58
121	BzdR, a repressor that controls the anaerobic catabolism of benzoate in <i>Azoarcus</i> sp. CIB, is the first member of a new subfamily of transcriptional regulators. <i>Journal of Biological Chemistry</i> , 2005 , 280, 10683-94	5.4	58
120	New Insights on Steroid Biotechnology. <i>Frontiers in Microbiology</i> , 2018 , 9, 958	5.7	54
119	Insights into pneumococcal fratricide from the crystal structures of the modular killing factor LytC. <i>Nature Structural and Molecular Biology</i> , 2010 , 17, 576-81	17.6	53
118	Elucidation of the molecular recognition of bacterial cell wall by modular pneumococcal phage endolysin CPL-1. <i>Journal of Biological Chemistry</i> , 2007 , 282, 24990-9	5.4	53
117	PHACOS, a functionalized bacterial polyester with bactericidal activity against methicillin-resistant <i>Staphylococcus aureus</i> . <i>Biomaterials</i> , 2014 , 35, 14-24	15.6	50
116	Promotion of multipoint covalent immobilization through different regions of genetically modified penicillin G acylase from <i>E. coli</i> . <i>Process Biochemistry</i> , 2010 , 45, 390-398	4.8	50
115	Whole-genome analysis of <i>Azoarcus</i> sp. strain CIB provides genetic insights to its different lifestyles and predicts novel metabolic features. <i>Systematic and Applied Microbiology</i> , 2015 , 38, 462-71	4.2	49
114	The PhaD regulator controls the simultaneous expression of the pha genes involved in polyhydroxyalkanoate metabolism and turnover in <i>Pseudomonas putida</i> KT2442. <i>Environmental Microbiology</i> , 2010 , 12, 1591-603	5.2	49
113	Identification and biochemical evidence of a medium-chain-length polyhydroxyalkanoate depolymerase in the <i>Bdellovibrio bacteriovorus</i> predatory hydrolytic arsenal. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 6017-26	4.8	49
112	Molecular characterization of the gallate dioxygenase from <i>Pseudomonas putida</i> KT2440. The prototype of a new subgroup of extradiol dioxygenases. <i>Journal of Biological Chemistry</i> , 2005 , 280, 35382-90	5.4	48
111	New challenges for syngas fermentation: towards production of biopolymers. <i>Journal of Chemical Technology and Biotechnology</i> , 2015 , 90, 1735-1751	3.5	47

110	Structural organization of the major autolysin from <i>Streptococcus pneumoniae</i> . <i>Journal of Biological Chemistry</i> , 1996 , 271, 6832-8	5.4	47
109	Molecular determinants of the hpa regulatory system of <i>Escherichia coli</i> : the HpaR repressor. <i>Nucleic Acids Research</i> , 2003 , 31, 6598-609	20.1	45
108	Genome organization and molecular analysis of the temperate bacteriophage MM1 of <i>Streptococcus pneumoniae</i> . <i>Journal of Bacteriology</i> , 2003 , 185, 2362-8	3.5	44
107	Molecular peculiarities of the <i>lytA</i> gene isolated from clinical pneumococcal strains that are bile insoluble. <i>Journal of Clinical Microbiology</i> , 2002 , 40, 2545-54	9.7	44
106	Crystal structure of CbpF, a bifunctional choline-binding protein and autolysis regulator from <i>Streptococcus pneumoniae</i> . <i>EMBO Reports</i> , 2009 , 10, 246-51	6.5	43
105	Stabilization of a multimeric beta-galactosidase from <i>Thermus</i> sp. strain T2 by immobilization on novel heterofunctional epoxy supports plus aldehyde-dextran cross-linking. <i>Biotechnology Progress</i> , 2004 , 20, 388-92	2.8	43
104	NMR investigations of protein-carbohydrate interactions binding studies and refined three-dimensional solution structure of the complex between the B domain of wheat germ agglutinin and N,NQN"-triacylchitotriose. <i>FEBS Journal</i> , 2000 , 267, 3965-78		43
103	Microbial synthesis of poly(beta-hydroxyalkanoates) bearing phenyl groups from <i>Pseudomonas putida</i> : chemical structure and characterization. <i>Biomacromolecules</i> , 2001 , 2, 562-7	6.9	41
102	A highly conserved mycobacterial cholesterol catabolic pathway. <i>Environmental Microbiology</i> , 2013 , 15, 2342-59	5.2	40
101	Characterization of the last step of the aerobic phenylacetic acid degradation pathway. <i>Microbiology (United Kingdom)</i> , 2007 , 153, 357-365	2.9	39
100	Genetic modification of the penicillin G acylase surface to improve its reversible immobilization on ionic exchangers. <i>Applied and Environmental Microbiology</i> , 2007 , 73, 312-9	4.8	37
99	Overproduction of <i>Thermus</i> sp. Strain T2 beta-galactosidase in <i>Escherichia coli</i> and preparation by using tailor-made metal chelate supports. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 1967-72	4.8	37
98	Characterization of genetic transformation in <i>Streptococcus oralis</i> NCTC 11427: expression of the pneumococcal amidase in <i>S. oralis</i> using a new shuttle vector. <i>Molecular Genetics and Genomics</i> , 1988 , 215, 53-7		37
97	Disposable amperometric magnetoimmunosensors for the specific detection of <i>Streptococcus pneumoniae</i> . <i>Biosensors and Bioelectronics</i> , 2010 , 26, 1225-30	11.8	35
96	Regulation of the <i>mhp</i> cluster responsible for 3-(3-hydroxyphenyl)propionic acid degradation in <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2003 , 278, 27575-85	5.4	33
95	Structural characterization of the unligated and choline-bound forms of the major pneumococcal autolysin <i>LytA</i> amidase. Conformational transitions induced by temperature. <i>Journal of Biological Chemistry</i> , 1996 , 271, 29152-61	5.4	33
94	Purification and characterization of the autolytic glycosidase of <i>Streptococcus pneumoniae</i> . <i>Biochemical and Biophysical Research Communications</i> , 1989 , 158, 251-6	3.4	32
93	Analysis of dibenzothiophene desulfurization in a recombinant <i>Pseudomonas putida</i> strain. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 875-7	4.8	31

92	Engineering the D-amino acid oxidase from <i>Trigonopsis variabilis</i> to facilitate its overproduction in <i>Escherichia coli</i> and its downstream processing by tailor-made metal chelate supports. <i>Enzyme and Microbial Technology</i> , 1999 , 25, 88-95	3.8	31
91	Structural and thermodynamic characterization of Pal, a phage natural chimeric lysin active against pneumococci. <i>Journal of Biological Chemistry</i> , 2004 , 279, 43697-707	5.4	30
90	<i>Mycobacterium smegmatis</i> is a suitable cell factory for the production of steroidal synthons. <i>Microbial Biotechnology</i> , 2017 , 10, 138-150	6.3	28
89	Development of amperometric magnetogenosensors coupled to asymmetric PCR for the specific detection of <i>Streptococcus pneumoniae</i> . <i>Analytical and Bioanalytical Chemistry</i> , 2011 , 399, 2413-20	4.4	28
88	Genetic characterization of the phenylacetyl-coenzyme A oxygenase from the aerobic phenylacetic acid degradation pathway of <i>Escherichia coli</i> . <i>Applied and Environmental Microbiology</i> , 2006 , 72, 7422-6	4.8	28
87	Genetic characterization of the styrene lower catabolic pathway of <i>Pseudomonas</i> sp. strain Y2. <i>Gene</i> , 2003 , 319, 71-83	3.8	28
86	Engineering <i>Mycobacterium smegmatis</i> for testosterone production. <i>Microbial Biotechnology</i> , 2017 , 10, 151-161	6.3	27
85	Structural studies of the lysozyme coded by the pneumococcal phage Cp-1. Conformational changes induced by choline. <i>FEBS Journal</i> , 1990 , 187, 409-16		27
84	Deciphering the transcriptional regulation of cholesterol catabolic pathway in mycobacteria: identification of the inducer of KstR repressor. <i>Journal of Biological Chemistry</i> , 2014 , 289, 17576-88	5.4	24
83	Characterization of the KstR-dependent promoter of the gene for the first step of the cholesterol degradative pathway in <i>Mycobacterium smegmatis</i> . <i>Microbiology (United Kingdom)</i> , 2011 , 157, 2670-2680	2.9	24
82	3-Hydroxyphenylpropionate and phenylpropionate are synergistic activators of the MhpR transcriptional regulator from <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2009 , 284, 21218-28	5.4	24
81	Single-step purification on DEAE-sephacel of recombinant polypeptides produced in <i>Escherichia coli</i> . <i>Nature Biotechnology</i> , 1992 , 10, 795-8	44.5	24
80	Reward for <i>Bdellovibrio bacteriovorus</i> for preying on a polyhydroxyalkanoate producer. <i>Environmental Microbiology</i> , 2013 , 15, 1204-15	5.2	23
79	Biochemical characterization of the transcriptional regulator BzdR from <i>Azoarcus</i> sp. CIB. <i>Journal of Biological Chemistry</i> , 2010 , 285, 35694-705	5.4	23
78	Genome sequence of the methanotrophic poly- β -hydroxybutyrate producer <i>Methylocystis parvus</i> OBBP. <i>Journal of Bacteriology</i> , 2012 , 194, 5709-10	3.5	23
77	Coregulation by phenylacetyl-coenzyme A-responsive PaaX integrates control of the upper and lower pathways for catabolism of styrene by <i>Pseudomonas</i> sp. strain Y2. <i>Journal of Bacteriology</i> , 2006 , 188, 4812-21	3.5	23
76	Selective and mild adsorption of large proteins on lowly activated immobilized metal ion affinity chromatography matrices. Purification of multimeric thermophilic enzymes overexpressed in <i>Escherichia coli</i> . <i>Journal of Chromatography A</i> , 2004 , 1055, 93-8	4.5	23
75	A gene containment strategy based on a restriction-modification system. <i>Environmental Microbiology</i> , 2000 , 2, 555-63	5.2	22

74	Isolation, characterization and physiological properties of an autolytic-deficient mutant of <i>Streptococcus pneumoniae</i> . <i>Molecular Genetics and Genomics</i> , 1986 , 204, 237-42		22
73	New tool for spreading proteins to the environment: Cry1Ab toxin immobilized to bioplastics. <i>Applied Microbiology and Biotechnology</i> , 2006 , 72, 88-93	5.7	21
72	The PaaX repressor, a link between penicillin G acylase and the phenylacetyl-coenzyme A catabolon of <i>Escherichia coli</i> W. <i>Journal of Bacteriology</i> , 2004 , 186, 2215-20	3.5	21
71	FLYCOP: metabolic modeling-based analysis and engineering microbial communities. <i>Bioinformatics</i> , 2018 , 34, i954-i963	7.2	21
70	Newly discovered penicillin acylase activity of aculeacin A acylase from <i>Actinoplanes utahensis</i> . <i>Applied and Environmental Microbiology</i> , 2007 , 73, 5378-81	4.8	20
69	VO1, a temperate bacteriophage of the type 19A multiresistant epidemic 8249 strain of <i>Streptococcus pneumoniae</i> : analysis of variability of lytic and putative C5 methyltransferase genes. <i>Microbial Drug Resistance</i> , 2003 , 9, 7-15	2.9	20
68	Aromatic metabolism versus carbon availability: the regulatory network that controls catabolism of less-preferred carbon sources in <i>Escherichia coli</i> . <i>FEMS Microbiology Reviews</i> , 2004 , 28, 503-18	15.1	20
67	Role of Asp-9 and Glu-36 in the active site of the pneumococcal CPL1 lysozyme: an evolutionary perspective of lysozyme mechanism. <i>Biochemistry</i> , 1992 , 31, 8495-9	3.2	20
66	Insights on the regulation of the phenylacetate degradation pathway from <i>Escherichia coli</i> . <i>Environmental Microbiology Reports</i> , 2014 , 6, 239-50	3.7	19
65	A finely tuned regulatory circuit of the nicotinic acid degradation pathway in <i>Pseudomonas putida</i> . <i>Environmental Microbiology</i> , 2011 , 13, 1718-32	5.2	19
64	Engineering alternative isobutanol production platforms. <i>AMB Express</i> , 2015 , 5, 119	4.1	18
63	Characterization of Ejl, the cell-wall amidase coded by the pneumococcal bacteriophage Ej-1. <i>Protein Science</i> , 2002 , 11, 1788-99	6.3	18
62	Insights into the structure-function relationships of pneumococcal cell wall lysozymes, LytC and Cpl-1. <i>Journal of Biological Chemistry</i> , 2008 , 283, 28618-28	5.4	18
61	Stabilization of a tetrameric enzyme (Amino acid ester hydrolase from <i>Acetobacter turbidans</i>) enables a very improved performance of ampicillin synthesis. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2001 , 11, 633-638		17
60	Testosterone Degradative Pathway of. <i>Genes</i> , 2019 , 10,	4.2	17
59	Identification of the <i>Geobacter metallireducens</i> bamVW two-component system, involved in transcriptional regulation of aromatic degradation. <i>Applied and Environmental Microbiology</i> , 2010 , 76, 383-5	4.8	16
58	Identification and analysis of a glutaryl-CoA dehydrogenase-encoding gene and its cognate transcriptional regulator from <i>Azoarcus</i> sp. CIB. <i>Environmental Microbiology</i> , 2008 , 10, 474-82	5.2	16
57	Oxygen-dependent regulation of the central pathway for the anaerobic catabolism of aromatic compounds in <i>Azoarcus</i> sp. strain CIB. <i>Journal of Bacteriology</i> , 2006 , 188, 2343-54	3.5	16

56	Overexpression of penicillin V acylase from <i>Streptomyces lavendulae</i> and elucidation of its catalytic residues. <i>Applied and Environmental Microbiology</i> , 2015 , 81, 1225-33	4.8	15
55	Cloning in <i>Escherichia coli</i> and molecular analysis of the sucrose system of the <i>Salmonella</i> plasmid SCR-53. <i>Molecular Genetics and Genomics</i> , 1985 , 201, 575-7		15
54	1,3-Propanediol production by NRRL-B199 from glycerol. Medium composition and operational conditions. <i>Biotechnology Reports (Amsterdam, Netherlands)</i> , 2015 , 6, 100-107	5.3	14
53	Clinical evaluation of a disposable amperometric magneto-genosensor for the detection and identification of <i>Streptococcus pneumoniae</i> . <i>Journal of Microbiological Methods</i> , 2014 , 103, 25-8	2.8	14
52	Design of catabolic cassettes for styrene biodegradation. <i>Antonie Van Leeuwenhoek</i> , 2003 , 84, 17-24	2.1	14
51	Molecular and functional analysis of the mce4 operon in <i>Mycobacterium smegmatis</i> . <i>Environmental Microbiology</i> , 2017 , 19, 3689-3699	5.2	13
50	The role of cofactor binding in tryptophan accessibility and conformational stability of His-tagged D-amino acid oxidase from <i>Trigonopsis variabilis</i> . <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2007 , 1774, 556-65	4	13
49	Unravelling the structure of the pneumococcal autolytic lysozyme. <i>Biochemical Journal</i> , 2005 , 391, 41-9	3.8	13
48	Allelic variation of polymorphic locus lytB, encoding a choline-binding protein, from streptococci of the mitis group. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 8706-13	4.8	13
47	Molecular characterization of a new gene cluster for steroid degradation in <i>Mycobacterium smegmatis</i> . <i>Environmental Microbiology</i> , 2017 , 19, 2546-2563	5.2	12
46	Unravelling the pleiotropic role of the MceG ATPase in <i>Mycobacterium smegmatis</i> . <i>Environmental Microbiology</i> , 2017 , 19, 2564-2576	5.2	12
45	Genome of sp. PHM005 Reveals a Complete and Active -AT PKS Gene Cluster for the Biosynthesis of Labrenzin. <i>Frontiers in Microbiology</i> , 2019 , 10, 2561	5.7	12
44	Pipelines for New Chemicals: a strategy to create new value chains and stimulate innovation-based economic revival in Southern European countries. <i>Environmental Microbiology</i> , 2014 , 16, 9-18	5.2	11
43	Identification of a missing link in the evolution of an enzyme into a transcriptional regulator. <i>PLoS ONE</i> , 2013 , 8, e57518	3.7	11
42	New insights into the BzdR-mediated transcriptional regulation of the anaerobic catabolism of benzoate in <i>Azoarcus</i> sp. CIB. <i>Microbiology (United Kingdom)</i> , 2008 , 154, 306-316	2.9	11
41	Production of a Thermoresistant Alpha-galactosidase from <i>Thermus</i> sp. Strain T2 for Food Processing. <i>Food Biotechnology</i> , 2007 , 21, 91-103	2.2	11
40	Unraveling the 17 β -Estradiol Degradation Pathway in NBRC 16725. <i>Frontiers in Microbiology</i> , 2020 , 11, 588300	5.7	11
39	Cloning, expression, and characterization of a peculiar choline-binding beta-galactosidase from <i>Streptococcus mitis</i> . <i>Applied and Environmental Microbiology</i> , 2009 , 75, 5972-80	4.8	10

38	Pneumococcal phosphorylcholine esterase, Pce, contains a metal binuclear center that is essential for substrate binding and catalysis. <i>Protein Science</i> , 2005 , 14, 3013-24	6.3	10
37	Identification and expression of the 11 β -steroid hydroxylase from <i>Cochliobolus lunatus</i> in <i>Corynebacterium glutamicum</i> . <i>Microbial Biotechnology</i> , 2019 , 12, 856-868	6.3	8
36	Engineering the l-Arabinose Isomerase from <i>Enterococcus Faecium</i> for d-Tagatose Synthesis. <i>Molecules</i> , 2017 , 22,	4.8	8
35	Characterization of the KstR2 regulator responsible of the lower cholesterol degradative pathway in <i>Mycobacterium smegmatis</i> . <i>Environmental Microbiology Reports</i> , 2015 , 7, 155-63	3.7	8
34	Cloning of the authentic bovine gene encoding pepsinogen a and its expression in microbial cells. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 2588-95	4.8	8
33	Construction of a chimeric thermostable pyrophosphatase to facilitate its purification and immobilization by using the choline-binding tag. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 4642-7	4.8	8
32	The subunit I of the respiratory-chain NADH dehydrogenase from <i>Cephalosporium acremonium</i> : the evolution of a mitochondrial gene. <i>Current Genetics</i> , 1986 , 10, 797-801	2.9	8
31	One-Step Immobilization and Stabilization of a Recombinant <i>Enterococcus faecium</i> DBFIQ E36 L-Arabinose Isomerase for D-Tagatose Synthesis. <i>Applied Biochemistry and Biotechnology</i> , 2019 , 188, 310-325	3.2	8
30	Quantifying dynamic mechanisms of auto-regulation in <i>Escherichia coli</i> with synthetic promoter in response to varying external phosphate levels. <i>Scientific Reports</i> , 2019 , 9, 2076	4.9	7
29	Unravelling a new catabolic pathway of C-19 steroids in <i>Mycobacterium smegmatis</i> . <i>Environmental Microbiology</i> , 2018 , 20, 1815-1827	5.2	7
28	Plasmids as Tools for Containment. <i>Microbiology Spectrum</i> , 2014 , 2,	8.9	7
27	The role of FIS protein in the physiological control of the expression of the <i>Escherichia coli</i> meta-hpa operon. <i>Microbiology (United Kingdom)</i> , 2008 , 154, 2151-2160	2.9	7
26	Genome Sequence of <i>Pseudomonas azelaica</i> Strain Aramco J. <i>Genome Announcements</i> , 2015 , 3,		6
25	Monitoring <i>Escherichia coli</i> growth in M63 media by ultrasonic noninvasive methods and correlation with spectrophotometric and HPLC techniques. <i>Applied Microbiology and Biotechnology</i> , 2010 , 85, 813-21	5.7	6
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