

JosÃ© MuÃ±oz-Dorado

List of Publications by Year in descending order

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3,534

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304743

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47

docs citations

47

times ranked

4144

citing authors

#	ARTICLE	IF	CITATIONS
1	Biodegradation and biological treatments of cellulose, hemicellulose and lignin: an overview. International Microbiology, 2002, 5, 53-63.	2.4	1,195
2	Complete genome sequence of the myxobacterium Sorangium cellulosum. Nature Biotechnology, 2007, 25, 1281-1289.	17.5	354
3	Myxobacteria: Moving, Killing, Feeding, and Surviving Together. Frontiers in Microbiology, 2016, 7, 781.	3.5	274
4	A gene encoding a protein serine/threonine kinase is required for normal development of <i>M. xanthus</i> , a gram-negative bacterium. Cell, 1991, 67, 995-1006.	28.9	264
5	Bacterial predation: 75 years and counting!. Environmental Microbiology, 2016, 18, 766-779.	3.8	190
6	Crystal Structure of <i>Myxococcus xanthus</i> Nucleoside Diphosphate Kinase and its Interaction with a Nucleotide Substrate at 2.0 Å... Resolution. Journal of Molecular Biology, 1993, 234, 1230-1247.	4.2	122
7	Eukaryotic-like protein kinases in the prokaryotes and the myxobacterial kinome. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 15950-15955.	7.1	105
8	<scp>PomZ</scp>, a <scp>ParA</scp>â€¢like protein, regulates <scp>Z</scp>â€¢ring formation and cell division in <i><scp>M</scp>yxococcus xanthus</i>. Molecular Microbiology, 2013, 87, 235-253.	2.5	103
9	<i>Myxococcus xanthus</i> induces actinorhodin overproduction and aerial mycelium formation by <i>Streptomyces coelicolor</i>. Microbial Biotechnology, 2011, 4, 175-183.	4.2	86
10	Identification of a putative eukaryotic-like protein kinase family in the developmental bacterium <i>Myxococcus xanthus</i> . Journal of Bacteriology, 1992, 174, 5450-5453.	2.2	66
11	Autophosphorylation of nucleoside diphosphate kinase from <i>Myxococcus xanthus</i> . Journal of Bacteriology, 1993, 175, 1176-1181.	2.2	65
12	<i>Myxococcus xanthus</i> , a gram-negative bacterium, contains a transmembrane protein serine/threonine kinase that blocks the secretion of beta-lactamase by phosphorylation.. Genes and Development, 1995, 9, 972-983.	5.9	58
13	Rhizobial galactogluconan determines the predatory pattern of <scp><i>M</i></scp><i>yxococcus xanthus</i> and protects <scp><i>S</i></scp><i>inorhizobium meliloti</i> from predation. Environmental Microbiology, 2014, 16, 2341-2350.	3.8	56
14	CorE from <i>Myxococcus xanthus</i> Is a Copper-Dependent RNA Polymerase Sigma Factor. PLoS Genetics, 2011, 7, e1002106.	3.5	49
15	The antibiotic crisis: How bacterial predators can help. Computational and Structural Biotechnology Journal, 2020, 18, 2547-2555.	4.1	45
16	A Large Family of Eukaryotic-Like Protein Ser/Thr Kinases of<i>Myxococcus xanthus</i>, a Developmental Bacterium. Microbial & Comparative Genomics, 2000, 5, 103-120.	0.4	41
17	Copper induction of carotenoid synthesis in the bacterium <i>Myxococcus xanthus</i> . Molecular Microbiology, 2005, 56, 1159-1168.	2.5	34
18	Eukaryotic-like protein serine/threonine kinases in <i>Myxococcus xanthus</i> , a developmental bacterium exhibiting social behavior. Journal of Cellular Biochemistry, 1993, 51, 29-33.	2.6	33

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19	Differential Expression of the Three Multicopper Oxidases from <i>Myxococcus xanthus</i> . <i>Journal of Bacteriology</i> , 2007, 189, 4887-4898.	2.2	31
20	Differential Regulation of Six Heavy Metal Efflux Systems in the Response of <i>< i> Myxococcus xanthus </i></i> to Copper. <i>Applied and Environmental Microbiology</i> , 2010, 76, 6069-6076.	3.1	31
21	Transcriptome dynamics of the <i>Myxococcus xanthus</i> multicellular developmental program. <i>ELife</i> , 2019, 8, .	6.0	31
22	Comprehensive Set of Integrative Plasmid Vectors for Copper-Inducible Gene Expression in <i>Myxococcus xanthus</i> . <i>Applied and Environmental Microbiology</i> , 2012, 78, 2515-2521.	3.1	29
23	In depth analysis of the mechanism of action of metal-dependent sigma factors: characterization of CorE2 from <i>< i> Myxococcus xanthus </i></i> . <i>Nucleic Acids Research</i> , 2016, 44, 5571-5584.	14.5	28
24	Characterization of manganese-dependent peroxidase isoenzymes from the ligninolytic fungus <i>Phanerochaete flavido-alba</i> . <i>Research in Microbiology</i> , 2002, 153, 547-554.	2.1	23
25	Oar, a 115-kilodalton membrane protein required for development of <i>Myxococcus xanthus</i> . <i>Journal of Bacteriology</i> , 1993, 175, 4756-4763.	2.2	22
26	Metalâ€“responsive RNA polymerase extracytoplasmic function (ECF) sigma factors. <i>Molecular Microbiology</i> , 2019, 112, 385-398.	2.5	21
27	Expression and Physiological Role of Three <i>< i> Myxococcus xanthus </i></i> Copper-Dependent P _{1B} -Type ATPases during Bacterial Growth and Development. <i>Applied and Environmental Microbiology</i> , 2010, 76, 6077-6084.	3.1	19
28	Copper and Melanin Play a Role in <i>Myxococcus xanthus</i> Predation on <i>Sinorhizobium meliloti</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 94.	3.5	18
29	The complex global response to copper in the multicellular bacterium <i>< i> Myxococcus xanthus </i></i> . <i>Metallomics</i> , 2018, 10, 876-886.	2.4	16
30	Role of Two Novel Two-Component Regulatory Systems in Development and Phosphatase Expression in <i>Myxococcus xanthus</i> . <i>Journal of Bacteriology</i> , 2003, 185, 1376-1383.	2.2	15
31	Identification of cis- and trans-acting elements involved in the expression of cold shock-inducible TIP1 gene of yeast <i>Saccharomyces cerevisiae</i> . <i>Nucleic Acids Research</i> , 1994, 22, 560-568.	14.5	13
32	The <i>Myxococcus xanthus</i> Two-Component System CorSR Regulates Expression of a Gene Cluster Involved in Maintaining Copper Tolerance during Growth and Development. <i>PLoS ONE</i> , 2013, 8, e68240.	2.5	13
33	Dissection of the sensor domain of the copper-responsive histidine kinase CorS from <i>Myxococcus xanthus</i> . <i>Environmental Microbiology Reports</i> , 2016, 8, 363-370.	2.4	12
34	A novel mechanism of bacterial adaptation mediated by copper-dependent RNA polymerase I f factors. <i>Transcription</i> , 2012, 3, 63-67.	3.1	9
35	Glycerol 3-Phosphate Inhibits Swarming and Aggregation of <i>Myxococcus xanthus</i> . <i>Journal of Bacteriology</i> , 2001, 183, 6135-6139.	2.2	8
36	Crystallization and preliminary X-ray diffraction analysis of nucleoside diphosphate kinase from <i>Myxococcus xanthus</i> . <i>Journal of Molecular Biology</i> , 1991, 220, 5-7.	4.2	7

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37	PhoR1-PhoP1, a Third Two-Component System of the Family PhoRP from <i>Myxococcus xanthus</i> : Role in Development. <i>Journal of Bacteriology</i> , 2005, 187, 4976-4983.	2.2	7
38	Mechanisms of Action of Non-Canonical ECF Sigma Factors. <i>International Journal of Molecular Sciences</i> , 2022, 23, 3601.	4.1	6
39	mlpB, a gene encoding a new lipoprotein in <i>Myxococcus xanthus</i> . <i>Journal of Applied Microbiology</i> , 2002, 92, 134-139.	3.1	4
40	phoR1, a gene encoding a new histidine protein kinase <i>Myxococcus xanthus</i> . <i>Antonie Van Leeuwenhoek</i> , 2003, 83, 361-368.	1.7	4
41	<i>Myxococcus xanthus</i> Pph2 Is a Manganese-dependent Protein Phosphatase Involved in Energy Metabolism. <i>Journal of Biological Chemistry</i> , 2009, 284, 28720-28728.	3.4	3
42	Protein Ser/Thr Kinases and Phosphatases in <i>Myxococcus xanthus</i> . , 2014, , 191-210.		3
43	Identification of the <i>Myxococcus xanthus</i> 59-kDa membrane-associated GTP-binding protein as a proton-translocating ATPase. <i>Gene</i> , 1994, 138, 133-137.	2.2	1
44	Pkn1., 1995, , 356-357.		0