## Jose Luis Garcia

## List of Publications by Year in Descending Order

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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.

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

| #   | Paper  | IF                        | Citations |
|-----|--|---------------------------|-----------|
| 163 | production of pederin by labrenzin pathway expansion <i>Metabolic Engineering Communications</i> , <b>2022</b> , 14, e00198  | 6.5                       | O         |
| 162 | Identification of trans-AT polyketide clusters in two marine bacteria reveals cryptic similarities between distinct symbiosis factors. <i>Environmental Microbiology</i> , <b>2021</b> , 23, 2509-2521                 | 5.2                       | 2         |
| 161 | Engineering the Steroid Hydroxylating System from in. <i>Microorganisms</i> , <b>2021</b> , 9,   | 4.9                       | 2         |
| 160 | Production of 11Ehydroxysteroids from sterols in a single fermentation step by Mycolicibacterium smegmatis. <i>Microbial Biotechnology</i> , <b>2021</b> , 14, 2514-2524   | 6.3                       | 4         |
| 159 | Unraveling the 17Estradiol Degradation Pathway in NBRC 16725. Frontiers in Microbiology, <b>2020</b> , 11, 588300  | 5.7                       | 11        |
| 158 | Heterologous production and biochemical characterization of a new highly glucose tolerant GH1 Eglucosidase from Anoxybacillus thermarum. <i>Process Biochemistry</i> , <b>2020</b> , 99, 1-8                           | 4.8                       | 5         |
| 157 | Degradation of Aromatic Compounds in Pseudomonas: A Systems Biology View <b>2019</b> , 639-687   |                           | О         |
| 156 | Bacterial Metabolism of Steroids <b>2019</b> , 315-336   |                           | 1         |
| 155 | Effect of Arthrospira supplementation on Oreochromis niloticus gut microbiota and flesh quality. <i>Aquaculture Research</i> , <b>2019</b> , 50, 1448-1458   | 1.9                       | 3         |
| 154 | Quantifying dynamic mechanisms of auto-regulation in Escherichia coli with synthetic promoter in response to varying external phosphate levels. <i>Scientific Reports</i> , <b>2019</b> , 9, 2076                      | 4.9                       | 7         |
| 153 | Identification and expression of the 11Esteroid hydroxylase from Cochliobolus lunatus in Corynebacterium glutamicum. <i>Microbial Biotechnology</i> , <b>2019</b> , 12, 856-868  | 6.3                       | 8         |
| 152 | Testosterone Degradative Pathway of. <i>Genes</i> , <b>2019</b> , 10,  | 4.2                       | 17        |
| 151 | Genome of sp. PHM005 Reveals a Complete and Active -AT PKS Gene Cluster for the Biosynthesis of Labrenzin. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 2561   | 5.7                       | 12        |
| 150 | One-Step Immobilization and Stabilization of a Recombinant Enterococcus faecium DBFIQ E36 L-Arabinose Isomerase for D-Tagatose Synthesis. <i>Applied Biochemistry and Biotechnology</i> , <b>2019</b> , 188, 310-325   | 3.2                       | 8         |
| 149 | Unravelling a new catabolic pathway of C-19 steroids in Mycobacterium smegmatis. <i>Environmental Microbiology</i> , <b>2018</b> , 20, 1815-1827   | 5.2                       | 7         |
| 148 | New Insights on Steroid Biotechnology. Frontiers in Microbiology, 2018, 9, 958   | 5.7                       | 54        |
| 147 | Effect of spirulina (Arthrospira platensis) supplementation on tilapia (Oreochromis niloticus) growth and stress responsiveness under hypoxia. <i>Spanish Journal of Agricultural Research</i> , <b>2018</b> , 16, e06 | 5 <b>đ</b> 6 <sup>1</sup> | 2         |

## (2015-2018)

| 146                      | FLYCOP: metabolic modeling-based analysis and engineering microbial communities. <i>Bioinformatics</i> , <b>2018</b> , 34, i954-i963  | 7.2            | 21                         |
|--------------------------|---|----------------|----------------------------|
| 145                      | Molecular characterization of a new gene cluster for steroid degradation in Mycobacterium smegmatis. <i>Environmental Microbiology</i> , <b>2017</b> , 19, 2546-2563  | 5.2            | 12                         |
| 144                      | Unravelling the pleiotropic role of the MceG ATPase in Mycobacterium smegmatis. <i>Environmental Microbiology</i> , <b>2017</b> , 19, 2564-2576   | 5.2            | 12                         |
| 143                      | Microalgae, old sustainable food and fashion nutraceuticals. <i>Microbial Biotechnology</i> , <b>2017</b> , 10, 1017-10   | ) <b>24</b> .3 | 171                        |
| 142                      | Engineering a bzd cassette for the anaerobic bioconversion of aromatic compounds. <i>Microbial Biotechnology</i> , <b>2017</b> , 10, 1418-1425  | 6.3            | 5                          |
| 141                      | Molecular and functional analysis of the mce4 operon in Mycobacterium smegmatis. <i>Environmental Microbiology</i> , <b>2017</b> , 19, 3689-3699  | 5.2            | 13                         |
| 140                      | Mycobacterium smegmatis is a suitable cell factory for the production of steroidic synthons. <i>Microbial Biotechnology</i> , <b>2017</b> , 10, 138-150   | 6.3            | 28                         |
| 139                      | Engineering Mycobacterium smegmatis for testosterone production. <i>Microbial Biotechnology</i> , <b>2017</b> , 10, 151-161   | 6.3            | 27                         |
| 138                      | Engineering the l-Arabinose Isomerase from Enterococcus Faecium for d-Tagatose Synthesis. <i>Molecules</i> , <b>2017</b> , 22,  | 4.8            | 8                          |
|                          |   |                |                            |
| 137                      | Production of 4-Ene-3-ketosteroids in Corynebacterium glutamicum. <i>Catalysts</i> , <b>2017</b> , 7, 316   | 4              | 4                          |
| 137                      | Production of 4-Ene-3-ketosteroids in Corynebacterium glutamicum. <i>Catalysts</i> , <b>2017</b> , 7, 316  Bacterial Metabolism of Steroids <b>2017</b> , 1-22  | 4              | 3                          |
|                          |   | 4              |                            |
| 136                      | Bacterial Metabolism of Steroids <b>2017</b> , 1-22  Complete mitochondrial genome of Polymastia littoralis (Demospongiae, Polymastiidae).  | 4.5            |                            |
| 136<br>135               | Bacterial Metabolism of Steroids <b>2017</b> , 1-22  Complete mitochondrial genome of Polymastia littoralis (Demospongiae, Polymastiidae).  Mitochondrial DNA, <b>2016</b> , 27, 312-3  Whole genome sequencing of turbot (Scophthalmus maximus; Pleuronectiformes): a fish adapted   | 4.5            | 3                          |
| 136<br>135<br>134        | Bacterial Metabolism of Steroids <b>2017</b> , 1-22  Complete mitochondrial genome of Polymastia littoralis (Demospongiae, Polymastiidae).  Mitochondrial DNA, <b>2016</b> , 27, 312-3  Whole genome sequencing of turbot (Scophthalmus maximus; Pleuronectiformes): a fish adapted to demersal life. DNA Research, <b>2016</b> , 23, 181-92  |                | 3<br>1<br>103              |
| 136<br>135<br>134        | Bacterial Metabolism of Steroids 2017, 1-22  Complete mitochondrial genome of Polymastia littoralis (Demospongiae, Polymastiidae).  Mitochondrial DNA, 2016, 27, 312-3  Whole genome sequencing of turbot (Scophthalmus maximus; Pleuronectiformes): a fish adapted to demersal life. DNA Research, 2016, 23, 181-92  Genome sequence of the olive tree, Olea europaea. GigaScience, 2016, 5, 29  Engineering synthetic bacterial consortia for enhanced desulfurization and revalorization of oil  | 7.6            | 3<br>1<br>103<br>130       |
| 136<br>135<br>134<br>133 | Bacterial Metabolism of Steroids 2017, 1-22  Complete mitochondrial genome of Polymastia littoralis (Demospongiae, Polymastiidae).  Mitochondrial DNA, 2016, 27, 312-3  Whole genome sequencing of turbot (Scophthalmus maximus; Pleuronectiformes): a fish adapted to demersal life. DNA Research, 2016, 23, 181-92  Genome sequence of the olive tree, Olea europaea. GigaScience, 2016, 5, 29  Engineering synthetic bacterial consortia for enhanced desulfurization and revalorization of oil sulfur compounds. Metabolic Engineering, 2016, 35, 46-54  Extreme genomic erosion after recurrent demographic bottlenecks in the highly endangered | 7.6<br>9.7     | 3<br>1<br>103<br>130<br>58 |

| 128 | 1,3-Propanediol production by NRRL-B199 from glycerol. Medium composition and operational conditions. <i>Biotechnology Reports (Amsterdam, Netherlands)</i> , <b>2015</b> , 6, 100-107                             | 5.3  | 14 |
|-----|--|------|----|
| 127 | Engineering alternative isobutanol production platforms. <i>AMB Express</i> , <b>2015</b> , 5, 119   | 4.1  | 18 |
| 126 | New challenges for syngas fermentation: towards production of biopolymers. <i>Journal of Chemical Technology and Biotechnology</i> , <b>2015</b> , 90, 1735-1751   | 3.5  | 47 |
| 125 | Genome Sequence of Pseudomonas azelaica Strain Aramco J. <i>Genome Announcements</i> , <b>2015</b> , 3,  |      | 6  |
| 124 | Overexpression of penicillin V acylase from Streptomyces lavendulae and elucidation of its catalytic residues. <i>Applied and Environmental Microbiology</i> , <b>2015</b> , 81, 1225-33                           | 4.8  | 15 |
| 123 | Draft Genome Sequence of Actinoplanes utahensis NRRL 12052, a Microorganism Involved in Industrial Production of Pharmaceutical Intermediates. <i>Genome Announcements</i> , <b>2015</b> , 3,                      |      | 4  |
| 122 | Characterization of the KstR2 regulator responsible of the lower cholesterol degradative pathway in Mycobacterium smegmatis. <i>Environmental Microbiology Reports</i> , <b>2015</b> , 7, 155-63                   | 3.7  | 8  |
| 121 | PHACOS, a functionalized bacterial polyester with bactericidal activity against methicillin-resistant Staphylococcus aureus. <i>Biomaterials</i> , <b>2014</b> , 35, 14-24   | 15.6 | 50 |
| 120 | Insights on the regulation of the phenylacetate degradation pathway from Escherichia coli. <i>Environmental Microbiology Reports</i> , <b>2014</b> , 6, 239-50   | 3.7  | 19 |
| 119 | Clinical evaluation of a disposable amperometric magneto-genosensor for the detection and identification of Streptococcus pneumoniae. <i>Journal of Microbiological Methods</i> , <b>2014</b> , 103, 25-8          | 2.8  | 14 |
| 118 | Plasmids as Tools for Containment. <i>Microbiology Spectrum</i> , <b>2014</b> , 2,   | 8.9  | 7  |
| 117 | Deciphering the transcriptional regulation of cholesterol catabolic pathway in mycobacteria: identification of the inducer of KstR repressor. <i>Journal of Biological Chemistry</i> , <b>2014</b> , 289, 17576-88 | 5.4  | 24 |
| 116 | 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> , <b>2014</b> , 16, 9-18      | 5.2  | 11 |
| 115 | A highly conserved mycobacterial cholesterol catabolic pathway. <i>Environmental Microbiology</i> , <b>2013</b> , 15, 2342-59  | 5.2  | 40 |
| 114 | Genome of the Psychrophilic Bacterium Bacillus psychrosaccharolyticus, a Potential Source of 2@Deoxyribosyltransferase for Industrial Nucleoside Synthesis. <i>Genome Announcements</i> , <b>2013</b> , 1,         |      | 5  |
| 113 | Reward for Bdellovibrio bacteriovorus for preying on a polyhydroxyalkanoate producer. <i>Environmental Microbiology</i> , <b>2013</b> , 15, 1204-15  | 5.2  | 23 |
| 112 | Identification of a missing link in the evolution of an enzyme into a transcriptional regulator. <i>PLoS ONE</i> , <b>2013</b> , 8, e57518   | 3.7  | 11 |
| 111 | Catabolism and biotechnological applications of cholesterol degrading bacteria. <i>Microbial Biotechnology</i> , <b>2012</b> , 5, 679-99   | 6.3  | 99 |

| 110 | Cholesterol metabolism in Mycobacterium smegmatis. Environmental Microbiology Reports, 2012, 4, 168  | - <b>3</b> 7 | 64 |
|-----|--|--------------|----|
| 109 | Genome sequence of the methanotrophic poly-Ehydroxybutyrate producer Methylocystis parvus OBBP. <i>Journal of Bacteriology</i> , <b>2012</b> , 194, 5709-10  | 3.5          | 23 |
| 108 | Identification and biochemical evidence of a medium-chain-length polyhydroxyalkanoate depolymerase in the Bdellovibrio bacteriovorus predatory hydrolytic arsenal. <i>Applied and Environmental Microbiology</i> , <b>2012</b> , 78, 6017-26 | 4.8          | 49 |
| 107 | Bacterial degradation of benzoate: cross-regulation between aerobic and anaerobic pathways.<br>Journal of Biological Chemistry, <b>2012</b> , 287, 10494-10508   | 5.4          | 66 |
| 106 | Selection of Ceratitis capitata (Diptera: Tephritidae) specific recombinant monoclonal phage display antibodies for prey detection analysis. <i>PLoS ONE</i> , <b>2012</b> , 7, e51440   | 3.7          | 1  |
| 105 | A finely tuned regulatory circuit of the nicotinic acid degradation pathway in Pseudomonas putida. <i>Environmental Microbiology</i> , <b>2011</b> , 13, 1718-32   | 5.2          | 19 |
| 104 | Unravelling the gallic acid degradation pathway in bacteria: the gal cluster from Pseudomonas putida. <i>Molecular Microbiology</i> , <b>2011</b> , 79, 359-74   | 4.1          | 58 |
| 103 | Controlled autolysis facilitates the polyhydroxyalkanoate recovery in Pseudomonas putida KT2440. <i>Microbial Biotechnology</i> , <b>2011</b> , 4, 533-47  | 6.3          | 64 |
| 102 | Characterization of the KstR-dependent promoter of the gene for the first step of the cholesterol degradative pathway in Mycobacterium smegmatis. <i>Microbiology (United Kingdom)</i> , <b>2011</b> , 157, 2670-268                         | <b>∂</b> ·9  | 24 |
| 101 | Development of amperometric magnetogenosensors coupled to asymmetric PCR for the specific detection of Streptococcus pneumoniae. <i>Analytical and Bioanalytical Chemistry</i> , <b>2011</b> , 399, 2413-20                                  | 4.4          | 28 |
| 100 | Disruption of Ebxidation pathway in Pseudomonas putida KT2442 to produce new functionalized PHAs with thioester groups. <i>Applied Microbiology and Biotechnology</i> , <b>2011</b> , 89, 1583-98  | 5.7          | 71 |
| 99  | Inhibition of Recombinant D-Amino Acid Oxidase from Trigonopsis variabilis by Salts. <i>Enzyme Research</i> , <b>2011</b> , 2011, 158541   | 2.4          |    |
| 98  | Insights into pneumococcal fratricide from the crystal structures of the modular killing factor LytC. <i>Nature Structural and Molecular Biology</i> , <b>2010</b> , 17, 576-81  | 17.6         | 53 |
| 97  | The turnover of medium-chain-length polyhydroxyalkanoates in Pseudomonas putida KT2442 and the fundamental role of PhaZ depolymerase for the metabolic balance. <i>Environmental Microbiology</i> , <b>2010</b> , 12, 207-21                 | 5.2          | 87 |
| 96  | The PhaD regulator controls the simultaneous expression of the pha genes involved in polyhydroxyalkanoate metabolism and turnover in Pseudomonas putida KT2442. <i>Environmental Microbiology</i> , <b>2010</b> , 12, 1591-603               | 5.2          | 49 |
| 95  | Identification of the Geobacter metallireducens bamVW two-component system, involved in transcriptional regulation of aromatic degradation. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 383-5                          | 4.8          | 16 |
| 94  | Biochemical characterization of the transcriptional regulator BzdR from Azoarcus sp. CIB. <i>Journal of Biological Chemistry</i> , <b>2010</b> , 285, 35694-705  | 5.4          | 23 |
| 93  | Monitoring Escherichia coli growth in M63 media by ultrasonic noninvasive methods and correlation with spectrophotometric and HPLC techniques. <i>Applied Microbiology and Biotechnology</i> , <b>2010</b> , 85, 813-21                      | 5.7          | 6  |

| 92 | Disposable amperometric magnetoimmunosensors for the specific detection of Streptococcus pneumoniae. <i>Biosensors and Bioelectronics</i> , <b>2010</b> , 26, 1225-30   | 11.8 | 35  |
|----|---|------|-----|
| 91 | Promotion of multipoint covalent immobilization through different regions of genetically modified penicillin G acylase from E. coli. <i>Process Biochemistry</i> , <b>2010</b> , 45, 390-398  | 4.8  | 50  |
| 90 | Crystallization of the pneumococcal autolysin LytC: in-house phasing using novel lanthanide complexes. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , <b>2010</b> , 66, 448-51  |      | 3   |
| 89 | A preliminary crystallographic study of recombinant NicX, an Fe(2+)-dependent 2,5-dihydroxypyridine dioxygenase from Pseudomonas putida KT2440. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , <b>2010</b> , 66, 549-53 |      | 4   |
| 88 | 3-Hydroxyphenylpropionate and phenylpropionate are synergistic activators of the MhpR transcriptional regulator from Escherichia coli. <i>Journal of Biological Chemistry</i> , <b>2009</b> , 284, 21218-28   | 5.4  | 24  |
| 87 | Analysis of dibenzothiophene desulfurization in a recombinant Pseudomonas putida strain. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 875-7  | 4.8  | 31  |
| 86 | Cloning, expression, and characterization of a peculiar choline-binding beta-galactosidase from Streptococcus mitis. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 5972-80  | 4.8  | 10  |
| 85 | Crystal structure of CbpF, a bifunctional choline-binding protein and autolysis regulator from Streptococcus pneumoniae. <i>EMBO Reports</i> , <b>2009</b> , 10, 246-51   | 6.5  | 43  |
| 84 | Crystal structure of CbpF, a bifunctional choline-binding protein and autolysis regulator from Streptococcus pneumoniae. <i>EMBO Reports</i> , <b>2009</b> , 10, 413-413  | 6.5  | 2   |
| 83 | Anaerobic catabolism of aromatic compounds: a genetic and genomic view. <i>Microbiology and Molecular Biology Reviews</i> , <b>2009</b> , 73, 71-133  | 13.2 | 312 |
| 82 | Identification and analysis of a glutaryl-CoA dehydrogenase-encoding gene and its cognate transcriptional regulator from Azoarcus sp. CIB. <i>Environmental Microbiology</i> , <b>2008</b> , 10, 474-82   | 5.2  | 16  |
| 81 | Deciphering the genetic determinants for aerobic nicotinic acid degradation: the nic cluster from Pseudomonas putida KT2440. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 11329-34     | 11.5 | 112 |
| 80 | The role of FIS protein in the physiological control of the expression of the Escherichia coli meta-hpa operon. <i>Microbiology (United Kingdom)</i> , <b>2008</b> , 154, 2151-2160   | 2.9  | 7   |
| 79 | Insights into the structure-function relationships of pneumococcal cell wall lysozymes, LytC and Cpl-1. <i>Journal of Biological Chemistry</i> , <b>2008</b> , 283, 28618-28  | 5.4  | 18  |
| 78 | New insights into the BzdR-mediated transcriptional regulation of the anaerobic catabolism of benzoate in Azoarcus sp. CIB. <i>Microbiology (United Kingdom)</i> , <b>2008</b> , 154, 306-316   | 2.9  | 11  |
| 77 | The role of cofactor binding in tryptophan accessibility and conformational stability of His-tagged D-amino acid oxidase from Trigonopsis variabilis. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2007</b> , 1774, 556-65     | 4    | 13  |
| 76 | Elucidation of the molecular recognition of bacterial cell wall by modular pneumococcal phage endolysin CPL-1. <i>Journal of Biological Chemistry</i> , <b>2007</b> , 282, 24990-9  | 5.4  | 53  |
| 75 | Production of a Thermoresistant Alpha-galactosidase from Thermus sp. Strain T2 for Food Processing. <i>Food Biotechnology</i> , <b>2007</b> , 21, 91-103  | 2.2  | 11  |

| 74 | Genetic modification of the penicillin G acylase surface to improve its reversible immobilization on ionic exchangers. <i>Applied and Environmental Microbiology</i> , <b>2007</b> , 73, 312-9                                       | 4.8                 | 37   |
|----|--|---------------------|------|
| 73 | Newly discovered penicillin acylase activity of aculeacin A acylase from Actinoplanes utahensis. <i>Applied and Environmental Microbiology</i> , <b>2007</b> , 73, 5378-81   | 4.8                 | 20   |
| 72 | Characterization of the last step of the aerobic phenylacetic acid degradation pathway. <i>Microbiology (United Kingdom)</i> , <b>2007</b> , 153, 357-365  | 2.9                 | 39   |
| 71 | Taking aim on bacterial pathogens: from phage therapy to enzybiotics. <i>Current Opinion in Microbiology</i> , <b>2007</b> , 10, 461-72  | 7.9                 | 175  |
| 70 | New tool for spreading proteins to the environment: Cry1Ab toxin immobilized to bioplastics. <i>Applied Microbiology and Biotechnology</i> , <b>2006</b> , 72, 88-93   | 5.7                 | 21   |
| 69 | Coregulation by phenylacetyl-coenzyme A-responsive PaaX integrates control of the upper and lower pathways for catabolism of styrene by Pseudomonas sp. strain Y2. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 4812-21       | 3.5                 | 23   |
| 68 | Genetic characterization of the phenylacetyl-coenzyme A oxygenase from the aerobic phenylacetic acid degradation pathway of Escherichia coli. <i>Applied and Environmental Microbiology</i> , <b>2006</b> , 72, 7422-6               | 4.8                 | 28   |
| 67 | Oxygen-dependent regulation of the central pathway for the anaerobic catabolism of aromatic compounds in Azoarcus sp. strain CIB. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 2343-54  | 3.5                 | 16   |
| 66 | Unravelling the structure of the pneumococcal autolytic lysozyme. <i>Biochemical Journal</i> , <b>2005</b> , 391, 41-9   | 3.8                 | 13   |
| 65 | Pneumococcal phosphorylcholine esterase, Pce, contains a metal binuclear center that is essential for substrate binding and catalysis. <i>Protein Science</i> , <b>2005</b> , 14, 3013-24  | 6.3                 | 10   |
| 64 | Molecular characterization of the safracin biosynthetic pathway from Pseudomonas fluorescens A2-2: designing new cytotoxic compounds. <i>Molecular Microbiology</i> , <b>2005</b> , 56, 144-54                                       | 4.1                 | 84   |
| 63 | Insights into pneumococcal pathogenesis from the crystal structure of the modular teichoic acid phosphorylcholine esterase Pce. <i>Nature Structural and Molecular Biology</i> , <b>2005</b> , 12, 533-8                             | 17.6                | 75   |
| 62 | Genomic sequence of the pathogenic and allergenic filamentous fungus Aspergillus fumigatus. <i>Nature</i> , <b>2005</b> , 438, 1151-6  | 50.4                | 1114 |
| 61 | Molecular characterization of the gallate dioxygenase from Pseudomonas putida KT2440. The prototype of a new subgroup of extradiol dioxygenases. <i>Journal of Biological Chemistry</i> , <b>2005</b> , 280, 3538                    | 3 <del>2:∕9</del> 0 | 48   |
| 60 | BzdR, a repressor that controls the anaerobic catabolism of benzoate in Azoarcus sp. CIB, is the first member of a new subfamily of transcriptional regulators. <i>Journal of Biological Chemistry</i> , <b>2005</b> , 280, 10683-94 | 5.4                 | 58   |
| 59 | Allelic variation of polymorphic locus lytB, encoding a choline-binding protein, from streptococci of the mitis group. <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 8706-13                                     | 4.8                 | 13   |
| 58 | Stabilization of a multimeric beta-galactosidase from Thermus sp. strain T2 by immobilization on novel heterofunctional epoxy supports plus aldehyde-dextran cross-linking. <i>Biotechnology Progress</i> , <b>2004</b> , 20, 388-92 | 2.8                 | 43   |
| 57 | Stabilization of penicillin G acylase from Escherichia coli: site-directed mutagenesis of the protein surface to increase multipoint covalent attachment. <i>Applied and Environmental Microbiology</i> , <b>2004</b> , 70, 1249-51  | 4.8                 | 93   |

| 56 | Cloning of the authentic bovine gene encoding pepsinogen a and its expression in microbial cells. <i>Applied and Environmental Microbiology</i> , <b>2004</b> , 70, 2588-95   | 4.8           | 8   |
|----|---|---------------|-----|
| 55 | The PaaX repressor, a link between penicillin G acylase and the phenylacetyl-coenzyme A catabolon of Escherichia coli W. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 2215-20  | 3.5           | 21  |
| 54 | The bzd gene cluster, coding for anaerobic benzoate catabolism, in Azoarcus sp. strain CIB. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 5762-74   | 3.5           | 92  |
| 53 | Structural and thermodynamic characterization of Pal, a phage natural chimeric lysin active against pneumococci. <i>Journal of Biological Chemistry</i> , <b>2004</b> , 279, 43697-707  | 5.4           | 30  |
| 52 | Aromatic metabolism versus carbon availability: the regulatory network that controls catabolism of less-preferred carbon sources in Escherichia coli. <i>FEMS Microbiology Reviews</i> , <b>2004</b> , 28, 503-18   | 15.1          | 20  |
| 51 | Selective and mild adsorption of large proteins on lowly activated immobilized metal ion affinity chromatography matrices. Purification of multimeric thermophilic enzymes overexpressed in Escherichia coli. <i>Journal of Chromatography A</i> , <b>2004</b> , 1055, 93-8 | 4.5           | 23  |
| 50 | In vivo immobilization of fusion proteins on bioplastics by the novel tag BioF. <i>Applied and Environmental Microbiology</i> , <b>2004</b> , 70, 3205-12   | 4.8           | 82  |
| 49 | The homogentisate pathway: a central catabolic pathway involved in the degradation of L-phenylalanine, L-tyrosine, and 3-hydroxyphenylacetate in Pseudomonas putida. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 5062-77  | 3.5           | 190 |
| 48 | Construction of a chimeric thermostable pyrophosphatase to facilitate its purification and immobilization by using the choline-binding tag. <i>Applied and Environmental Microbiology</i> , <b>2004</b> , 70, 4642  | 2 <b>-4</b> 8 | 8   |
| 47 | VO1, a temperate bacteriophage of the type 19A multiresistant epidemic 8249 strain of Streptococcus pneumoniae: analysis of variability of lytic and putative C5 methyltransferase genes. <i>Microbial Drug Resistance</i> , <b>2003</b> , 9, 7-15                          | 2.9           | 20  |
| 46 | Regulation of the mhp cluster responsible for 3-(3-hydroxyphenyl)propionic acid degradation in Escherichia coli. <i>Journal of Biological Chemistry</i> , <b>2003</b> , 278, 27575-85   | 5.4           | 33  |
| 45 | Design of catabolic cassettes for styrene biodegradation. <i>Antonie Van Leeuwenhoek</i> , <b>2003</b> , 84, 17-24  | 2.1           | 14  |
| 44 | Structural basis for selective recognition of pneumococcal cell wall by modular endolysin from phage Cp-1. <i>Structure</i> , <b>2003</b> , 11, 1239-49   | 5.2           | 135 |
| 43 | Molecular and biochemical analysis of the system regulating the lytic/lysogenic cycle in the pneumococcal temperate phage MM1. <i>FEMS Microbiology Letters</i> , <b>2003</b> , 222, 193-7  | 2.9           | 6   |
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| 40 | Molecular determinants of the hpa regulatory system of Escherichia coli: the HpaR repressor. <i>Nucleic Acids Research</i> , <b>2003</b> , 31, 6598-609   | 20.1          | 45  |
| 39 | Genome organization and molecular analysis of the temperate bacteriophage MM1 of Streptococcus pneumoniae. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 2362-8   | 3.5           | 44  |

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| 36 | Crystallization and preliminary X-ray diffraction studies of the complete modular endolysin from Cp-1, a phage infecting Streptococcus pneumoniae. <i>Acta Crystallographica Section D: Biological Crystallography</i> , <b>2002</b> , 58, 1487-9   |      | 2   |
| 35 | Genomic analysis of the aromatic catabolic pathways from Pseudomonas putida KT2440. <i>Environmental Microbiology</i> , <b>2002</b> , 4, 824-41   | 5.2  | 380 |
| 34 | Molecular peculiarities of the lytA gene isolated from clinical pneumococcal strains that are bile insoluble. <i>Journal of Clinical Microbiology</i> , <b>2002</b> , 40, 2545-54   | 9.7  | 44  |
| 33 | Purification and polar localization of pneumococcal LytB, a putative endo-beta-N-acetylglucosaminidase: the chain-dispersing murein hydrolase. <i>Journal of Bacteriology</i> , <b>2002</b> , 184, 4988-5000  | 3.5  | 97  |
| 32 | 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 E. Coli beta-galactosidases. <i>Journal of the American Chemical Society</i> , <b>2002</b> , 124, 4804-10 | 16.4 | 78  |
| 31 | Genetically engineered Pseudomonas: a factory of new bioplastics with broad applications. <i>Environmental Microbiology</i> , <b>2001</b> , 3, 612-8  | 5.2  | 66  |
| 30 | The phenylacetyl-CoA catabolon: a complex catabolic unit with broad biotechnological applications. <i>Molecular Microbiology</i> , <b>2001</b> , 39, 1434-42  | 4.1  | 132 |
| 29 | One-step purification, covalent immobilization, and additional stabilization of poly-His-tagged proteins using novel heterofunctional chelate-epoxy supports. <i>Biotechnology and Bioengineering</i> , <b>2001</b> , 76, 269-76  | 4.9  | 91  |
| 28 | 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> , <b>2001</b> , 915, 97-106   | 4.5  | 68  |
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