

# Santiago Gutiérrez Martínez

## List of Publications by Year in descending order

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124  
papers

4,969  
citations

66234

42  
h-index

110170

64  
g-index

125  
all docs

125  
docs citations

125  
times ranked

3220  
citing authors

#	ARTICLE	IF	CITATIONS
1	Use of the volatile trichodiene to reduce Fusarium head blight and trichothecene contamination in wheat. <i>Microbial Biotechnology</i> , 2022, 15, 513-527.	2.0	10
2	Volatile-mediated interactions between <i>Trichoderma harzianum</i> and <i>Acanthoscelides obtectus</i> : A novel in vitro methodology to evaluate the impact of microbial volatile compounds on dry grain storage pests. <i>Biological Control</i> , 2022, 169, 104868.	1.4	3
3	Novel culture chamber to evaluate in vitro plant-microbe volatile interactions: Effects of <i>Trichoderma harzianum</i> volatiles on wheat plantlets. <i>Plant Science</i> , 2022, 320, 111286.	1.7	4
4	Use of VOC Chambers to evaluate the impact of microbial volatile compounds on dry grain insect pests. <i>MethodsX</i> , 2022, 9, 101734.	0.7	1
5	Organic and Conventional Bean Pesticides in Development of Autochthonous <i>Trichoderma</i> Strains. <i>Journal of Fungi</i> (Basel, Switzerland), 2022, 8, 603.	1.5	4
6	Phylogenomic Analysis of a 55.1-kb 19-Gene Dataset Resolves a Monophyletic <i>Fusarium</i> that Includes the <i>Fusarium solani</i> Species Complex. <i>Phytopathology</i> , 2021, 111, 1064-1079.	1.1	107
7	Influence of Fungicide Application and Vine Age on <i>Trichoderma</i> Diversity as Source of Biological Control Agents. <i>Agronomy</i> , 2021, 11, 446.	1.3	6
8	Influence of Physicochemical Characteristics of Bean Crop Soil in <i>Trichoderma</i> spp. Development. <i>Agronomy</i> , 2021, 11, 274.	1.3	6
9	Volatile Organic Compound Chamber: A Novel Technology for Microbiological Volatile Interaction Assays. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 248.	1.5	10
10	The Influence of Temperature on the Growth, Sporulation, Colonization, and Survival of <i>Trichoderma</i> spp. in Grapevine Pruning Wounds. <i>Agronomy</i> , 2021, 11, 1771.	1.3	10
11	Fungal Secondary Metabolism. , 2021, , 54-63.		0
12	Germination and Agronomic Traits of <i>Phaseolus vulgaris</i> L. Beans Sprayed with <i>Trichoderma</i> Strains and Attacked by <i>Acanthoscelides obtectus</i> . <i>Agronomy</i> , 2021, 11, 2130.	1.3	2
13	Distribution, Function, and Evolution of a Gene Essential for Trichothecene Toxin Biosynthesis in <i>Trichoderma</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 791641.	1.5	10
14	Editorial: Designing Bio-Formulations Based on Organic Amendments, Beneficial Microbes and Their Metabolites. <i>Frontiers in Microbiology</i> , 2021, 12, 832149.	1.5	2
15	Identification of plant genes putatively involved in the perception of fungal ergosterol squalene. <i>Journal of Integrative Plant Biology</i> , 2020, 62, 927-947.	4.1	16
16	Self-Inhibitory Activity of <i>Trichoderma</i> Soluble Metabolites and Their Antifungal Effects on <i>Fusarium oxysporum</i> . <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 176.	1.5	5
17	<i>Trichoderma</i> trichothecenes. , 2020, , 281-301.		4
18	Spores of <i>Trichoderma</i> strains sprayed over <i>Acanthoscelides obtectus</i> and <i>Phaseolus vulgaris</i> L. beans: Effects in the biology of the bean weevil. <i>Journal of Stored Products Research</i> , 2020, 88, 101666.	1.2	10

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19	Influence of Substrates in the Development of Bean and in Pathogenicity of <i>Rhizoctonia solani</i> JG K&Afrac14hn. <i>Agronomy</i> , 2020, 10, 707.	1.3	4
20	Colonization of <i>Vitis vinifera</i> L. by the Endophyte <i>Trichoderma</i> sp. Strain T154: Biocontrol Activity Against <i>Phaeoacremonium minimum</i> . <i>Frontiers in Plant Science</i> , 2020, 11, 1170.	1.7	29
21	Evaluation of substrates and additives to <i>Trichoderma harzianum</i> development by qPCR. <i>Agronomy Journal</i> , 2020, 112, 3188-3194.	0.9	4
22	Genetic bases for variation in structure and biological activity of trichothecene toxins produced by diverse fungi. <i>Applied Microbiology and Biotechnology</i> , 2020, 104, 5185-5199.	1.7	21
23	Fungal Horizontal Gene Transfer: A History Beyond the Phylogenetic Kingdoms. , 2019, , 315-336.		5
24	A cytochrome P450 monooxygenase gene required for biosynthesis of the trichothecene toxin harzianum A in <i>Trichoderma</i> . <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 8087-8103.	1.7	13
25	Effect of trichodiene synthase encoding gene expression in <i>Trichoderma</i> strains on their effectiveness in the control of <i>Acanthoscelides obtectus</i> . <i>Journal of Stored Products Research</i> , 2019, 83, 275-280.	1.2	9
26	Synthesis of Trichodermin Derivatives and Their Antimicrobial and Cytotoxic Activities. <i>Molecules</i> , 2019, 24, 3811.	1.7	9
27	Effect of <i>Trichoderma velutinum</i> and <i>Rhizoctonia solani</i> on the Metabolome of Bean Plants ( <i>Phaseolus vulgaris</i> L.). <i>International Journal of Molecular Sciences</i> , 2019, 20, 549.	1.8	36
28	Requirement of Two Acyltransferases for 4-O-Acylation during Biosynthesis of Harzianum A, an Antifungal Trichothecene Produced by <i>Trichoderma arundinaceum</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 723-734.	2.4	12
29	Role of <i>Trichoderma arundinaceum</i> tri10 in regulation of terpene biosynthetic genes and in control of metabolic flux. <i>Fungal Genetics and Biology</i> , 2019, 122, 31-46.	0.9	16
30	Relevance of the deletion of the <i>Tatri4</i> gene in the secondary metabolome of <i>Trichoderma arundinaceum</i> . <i>Organic and Biomolecular Chemistry</i> , 2018, 16, 2955-2965.	1.5	18
31	Investigations of <i>Trichoderma</i> spp. and <i>Beauveria bassiana</i> as biological control agent for <i>Xylotrechus arvicola</i> , a major insect pest in Spanish vineyards. <i>Journal of Economic Entomology</i> , 2018, 111, 2585-2591.	0.8	20
32	Effect of trichodiene production by <i>Trichoderma harzianum</i> on <i>Acanthoscelides obtectus</i> . <i>Journal of Stored Products Research</i> , 2018, 77, 231-239.	1.2	23
33	Evolution of structural diversity of trichothecenes, a family of toxins produced by plant pathogenic and entomopathogenic fungi. <i>PLoS Pathogens</i> , 2018, 14, e1006946.	2.1	141
34	Effect of deletion of a trichothecene toxin regulatory gene on the secondary metabolism transcriptome of the saprotrophic fungus <i>Trichoderma arundinaceum</i> . <i>Fungal Genetics and Biology</i> , 2018, 119, 29-46.	0.9	27
35	Inhibitory activity of <i>Beauveria bassiana</i> and <i>Trichoderma</i> spp. on the insect pests <i>Xylotrechus arvicola</i> (Coleoptera: Cerambycidae) and <i>Acanthoscelides obtectus</i> (Coleoptera: Chrisomelidae:). <i>Tj ETQq1 1 0.7843134 rgBT /Overlock</i>		
36	Involvement of <i>Trichoderma harzianum</i> Epl-1 Protein in the Regulation of <i>Botrytis</i> Virulence- and Tomato Defense-Related Genes. <i>Frontiers in Plant Science</i> , 2017, 8, 880.	1.7	40

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37	Involvement of the Transcriptional Coactivator ThMBF1 in the Biocontrol Activity of <i>Trichoderma harzianum</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 2273.	1.5	20
38	Nitrogen Metabolism and Growth Enhancement in Tomato Plants Challenged with <i>Trichoderma harzianum</i> Expressing the <i>Aspergillus nidulans</i> Acetamidase <i>amdS</i> Gene. <i>Frontiers in Microbiology</i> , 2016, 7, 1182.	1.5	34
39	Development of a qPCR Strategy to Select Bean Genes Involved in Plant Defense Response and Regulated by the <i>Trichoderma velutinum</i> – <i>Rhizoctonia solani</i> Interaction. <i>Frontiers in Plant Science</i> , 2016, 7, 1109.	1.7	46
40	Trichothecenes and aspinolides produced by <i>Trichoderma arundinaceum</i> regulate expression of <i>Botrytis cinerea</i> genes involved in virulence and growth. <i>Environmental Microbiology</i> , 2016, 18, 3991-4004.	1.8	25
41	Botrydial and botcinins produced by <i>Botrytis cinerea</i> regulate the expression of <i>Trichoderma arundinaceum</i> genes involved in trichothecene biosynthesis. <i>Molecular Plant Pathology</i> , 2016, 17, 1017-1031.	2.0	14
42	The Cerato-Platanin protein Epl-1 from <i>Trichoderma harzianum</i> is involved in mycoparasitism, plant resistance induction and self cell wall protection. <i>Scientific Reports</i> , 2016, 5, 17998.	1.6	77
43	Insecticidal activity of <i>Trichoderma harzianum</i> against <i>Xylotrechus arvicola</i> and <i>Acanthoscelides obtectus</i> immature stages. <i>Planta Medica</i> , 2016, 81, S1-S381.	0.7	2
44	Effect of Farnesol, a compound produced by <i>Trichoderma</i> when growing on bean ( <i>Phaseolus vulgaris</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	0.7	1
45	Trichodiene Production in a <i>Trichoderma harzianum</i> <i>erg1</i> -Silenced Strain Provides Evidence of the Importance of the Sterol Biosynthetic Pathway in Inducing Plant Defense-Related Gene Expression. <i>Molecular Plant-Microbe Interactions</i> , 2015, 28, 1181-1197.	1.4	38
46	The importance of chorismate mutase in the biocontrol potential of <i>Trichoderma parareesei</i> . <i>Frontiers in Microbiology</i> , 2015, 6, 1181.	1.5	28
47	Influence of <i>Rhizoctonia solani</i> and <i>Trichoderma</i> spp. in growth of bean ( <i>Phaseolus vulgaris</i> L.) and in the induction of plant defense-related genes. <i>Frontiers in Plant Science</i> , 2015, 6, 685.	1.7	116
48	<i>Trichoderma</i> Transformation Methods. <i>Fungal Biology</i> , 2015, , 41-48.	0.3	4
49	Effects of Trichothecene Production on the Plant Defense Response and Fungal Physiology: Overexpression of the <i>Trichoderma arundinaceum</i> <i>tri4</i> Gene in <i>T. harzianum</i> . <i>Applied and Environmental Microbiology</i> , 2015, 81, 6355-6366.	1.4	37
50	Novel aspinolide production by <i>Trichoderma arundinaceum</i> with a potential role in <i>Botrytis cinerea</i> antagonistic activity and plant defence priming. <i>Environmental Microbiology</i> , 2015, 17, 1103-1118.	1.8	56
51	Production of trichodiene by <i>Trichoderma harzianum</i> alters the perception of this biocontrol strain by plants and antagonized fungi. <i>Environmental Microbiology</i> , 2015, 17, 2628-2646.	1.8	64
52	Overexpression of <i>erg1</i> gene in <i>Trichoderma harzianum</i> CECT 2413: effect on the induction of tomato defence-related genes. <i>Journal of Applied Microbiology</i> , 2014, 117, 812-823.	1.4	22
53	Secondary Metabolism and Antimicrobial Metabolites of <i>Trichoderma</i> . , 2014, , 125-137.		59
54	Relevance of trichothecenes in fungal physiology: Disruption of <i>tri5</i> in <i>Trichoderma arundinaceum</i> . <i>Fungal Genetics and Biology</i> , 2013, 53, 22-33.	0.9	89

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55	The contribution of <i>Trichoderma</i> to balancing the costs of plant growth and defense. <i>International Microbiology</i> , 2013, 16, 69-80.	1.1	173
56	Involvement of <i>Trichoderma</i> Trichothecenes in the Biocontrol Activity and Induction of Plant Defense-Related Genes. <i>Applied and Environmental Microbiology</i> , 2012, 78, 4856-4868.	1.4	143
57	Overexpression of the trichodiene synthase gene <i>tri5</i> increases trichodermin production and antimicrobial activity in <i>Trichoderma brevicompactum</i> . <i>Fungal Genetics and Biology</i> , 2011, 48, 285-296.	0.9	110
58	Identification of Loci and Functional Characterization of Trichothecene Biosynthesis Genes in Filamentous Fungi of the Genus <i>Trichoderma</i> . <i>Applied and Environmental Microbiology</i> , 2011, 77, 4867-4877.	1.4	124
59	Functional Analysis of the <i>Trichoderma harzianum</i> <i>nox1</i> Gene, Encoding an NADPH Oxidase, Relates Production of Reactive Oxygen Species to Specific Biocontrol Activity against <i>Pythium ultimum</i> . <i>Applied and Environmental Microbiology</i> , 2011, 77, 3009-3016.	1.4	92
60	Overexpression of the <i>Trichoderma brevicompactum</i> <i>tri5</i> Gene: Effect on the Expression of the Trichodermin Biosynthetic Genes and on Tomato Seedlings. <i>Toxins</i> , 2011, 3, 1220-1232.	1.5	45
61	TvDim1 of <i>Trichoderma virens</i> is involved in redox-processes and confers resistance to oxidative stresses. <i>Current Genetics</i> , 2010, 56, 63-73.	0.8	18
62	Transgenic expression of the <i>Trichoderma harzianum</i> <i>hsp70</i> gene increases <i>Arabidopsis</i> resistance to heat and other abiotic stresses. <i>Journal of Plant Physiology</i> , 2010, 167, 659-665.	1.6	161
63	The ThPG1 Endopolygalacturonase Is Required for the <i>Trichoderma harzianum</i> "Plant Beneficial Interaction. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 1021-1031.	1.4	173
64	Cloning and characterization of the <i>Thcut1</i> gene encoding a cutinase of <i>Trichoderma harzianum</i> T34. <i>Current Genetics</i> , 2008, 54, 301-312.	0.8	29
65	Overexpression of a <i>Trichoderma</i> HSP70 gene increases fungal resistance to heat and other abiotic stresses. <i>Fungal Genetics and Biology</i> , 2008, 45, 1506-1513.	0.9	68
66	Age-Related Clinical, Serological, and Histopathological Features of Celiac Disease. <i>American Journal of Gastroenterology</i> , 2008, 103, 2360-2365.	0.2	114
67	Partial silencing of a hydroxy-methylglutaryl-CoA reductase-encoding gene in <i>Trichoderma harzianum</i> CECT 2413 results in a lower level of resistance to lovastatin and lower antifungal activity. <i>Fungal Genetics and Biology</i> , 2007, 44, 269-283.	0.9	60
68	The heterologous overexpression of <i>hsp23</i> , a small heat-shock protein gene from <i>Trichoderma virens</i> , confers thermotolerance to <i>T. harzianum</i> . <i>Current Genetics</i> , 2007, 52, 45-53.	0.8	39
69	Cloning and characterization of the <i>erg1</i> gene of <i>Trichoderma harzianum</i> : Effect of the <i>erg1</i> silencing on ergosterol biosynthesis and resistance to terbinafine. <i>Fungal Genetics and Biology</i> , 2006, 43, 164-178.	0.9	77
70	ThPTR2, a di/tri-peptide transporter gene from <i>Trichoderma harzianum</i> . <i>Fungal Genetics and Biology</i> , 2006, 43, 234-246.	0.9	41
71	Genome-wide analysis of differentially expressed genes from <i>Penicillium chrysogenum</i> grown with a repressing or a non-repressing carbon source. <i>Current Genetics</i> , 2006, 49, 85-96.	0.8	15
72	Detection of peptaibols and partial cloning of a putative peptaibol synthetase gene from <i>T. harzianum</i> CECT 2413. <i>Folia Microbiologica</i> , 2006, 51, 114-120.	1.1	18

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73	Generation, annotation and analysis of ESTs from <i>Trichoderma harzianum</i> CECT 2413. <i>BMC Genomics</i> , 2006, 7, 193.	1.2	60
74	A comparison of the phenotypic and genetic stability of recombinant <i>Trichoderma</i> spp. generated by protoplast- and <i>Agrobacterium</i> -mediated transformation. <i>Journal of Microbiology</i> , 2006, 44, 383-95.	1.3	43
75	Detection of putative peptide synthetase genes in <i>Trichoderma</i> species: Application of this method to the cloning of a gene from <i>T. harzianum</i> CECT 2413. <i>FEMS Microbiology Letters</i> , 2005, 244, 139-148.	0.7	41
76	Screening of antimicrobial activities in <i>Trichoderma</i> isolates representing three <i>Trichoderma</i> sections. <i>Mycological Research</i> , 2005, 109, 1397-1406.	2.5	47
77	Stable transformants of the azaphilone pigment-producing <i>Monascus purpureus</i> obtained by protoplast transformation and <i>Agrobacterium</i> -mediated DNA transfer. <i>Current Genetics</i> , 2003, 43, 447-452.	0.8	42
78	Expression of a synthetic copy of the bovine chymosin gene in <i>Aspergillus awamori</i> from constitutive and pH-regulated promoters and secretion using two different pre-pro sequences. <i>Biotechnology and Bioengineering</i> , 2003, 83, 249-259.	1.7	29
79	The isopenicillin N acyltransferases of <i>Aspergillus nidulans</i> and <i>Penicillium chrysogenum</i> differ in their ability to maintain the 40-kDa alpha-beta heterodimer in an undissociated form. <i>FEBS Journal</i> , 2003, 270, 1958-1968.	0.2	19
80	Co-transformation with autonomous replicating and integrative plasmids in <i>Penicillium chrysogenum</i> is highly efficient and leads in some cases to rescue of the intact integrative plasmid. <i>Fungal Genetics and Biology</i> , 2003, 40, 83-92.	0.9	14
81	A Novel Epimerization System in Fungal Secondary Metabolism Involved in the Conversion of Isopenicillin N into Penicillin N in <i>Acremonium chrysogenum</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 46216-46225.	1.6	71
82	Silencing of the <i>Aspergillopepsin B</i> ( <i>pepB</i> ) Gene of <i>Aspergillus awamori</i> by Antisense RNA Expression or Protease Removal by Gene Disruption Results in a Large Increase in Thaumatin Production. <i>Applied and Environmental Microbiology</i> , 2002, 68, 3550-3559.	1.4	49
83	Subcellular localization of the homocitrate synthase in <i>Penicillium chrysogenum</i> . <i>Molecular Genetics and Genomics</i> , 2002, 266, 711-719.	1.0	13
84	The <i>cefT</i> gene of <i>Acremonium chrysogenum</i> C10 encodes a putative multidrug efflux pump protein that significantly increases cephalosporin C production. <i>Molecular Genetics and Genomics</i> , 2002, 267, 673-683.	1.0	73
85	Characterization of the <i>gdhA</i> Gene from the Phytopathogen <i>Botrytis cinerea</i> . <i>Fungal Genetics and Biology</i> , 2001, 34, 193-206.	0.9	4
86	Targeted Inactivation of the <i>mecB</i> Gene, Encoding Cystathionine- $\beta$ -Lyase, Shows that the Reverse Transsulfuration Pathway Is Required for High-Level Cephalosporin Biosynthesis in <i>Acremonium chrysogenum</i> C10 but Not for Methionine Induction of the Cephalosporin Genes. <i>Journal of Bacteriology</i> , 2001, 183, 1765-1772.	1.0	38
87	Cloning and characterization of the gene <i>cah B</i> encoding a cephalosporin C acetylhydrolase from <i>Acremonium chrysogenum</i> . <i>Applied Microbiology and Biotechnology</i> , 2001, 57, 350-356.	1.7	16
88	Intrachromosomal recombination after targeted monocopy integration in <i>Penicillium chrysogenum</i> : stabilization of the direct repeats to prevent loss of the inserted gene. <i>Current Genetics</i> , 2001, 39, 231-236.	0.8	2
89	Characterization of the reverse transsulfuration gene <i>mecB</i> of <i>Acremonium chrysogenum</i> , which encodes a functional cystathionine- $\beta$ -lyase. <i>Molecular Genetics and Genomics</i> , 2001, 264, 746-754.	1.0	10
90	The specific transport system for lysine is fully inhibited by ammonium in <i>Penicillium chrysogenum</i> : an ammonium-insensitive system allows uptake in carbon-starved cells. <i>Antonie Van Leeuwenhoek</i> , 2000, 77, 91-100.	0.7	6

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91	Overexpression of the <i>lys1</i> gene in <i>Penicillium chrysogenum</i> : homocitrate synthase levels, $\hat{\pm}$ -aminoadipic acid pool and penicillin production. <i>Applied Microbiology and Biotechnology</i> , 2000, 54, 69-77.	1.7	18
92	Overexpression and lack of degradation of thaumatin in an aspergillopepsin A-defective mutant of <i>Aspergillus awamori</i> containing an insertion in the <i>pepA</i> gene. <i>Applied Microbiology and Biotechnology</i> , 2000, 54, 772-777.	1.7	33
93	A Novel Heptameric Sequence (TTAGTAA) Is the Binding Site for a Protein Required for High Level Expression of <i>pcbAB</i> , the First Gene of the Penicillin Biosynthesis in <i>Penicillium chrysogenum</i> . <i>Journal of Biological Chemistry</i> , 2000, 275, 2423-2430.	1.6	42
94	Transcription of the <i>pcbAB</i> , <i>pcbC</i> and <i>penDE</i> genes of <i>Penicillium chrysogenum</i> AS-P-78 is repressed by glucose and the repression is not reversed by alkaline pHs. <i>Microbiology (United Kingdom)</i> , 1999, 145, 317-324.	0.7	41
95	Penicillin and cephalosporin biosynthesis: mechanism of carbon catabolite regulation of penicillin production. <i>Antonie Van Leeuwenhoek</i> , 1999, 75, 21-31.	0.7	63
96	Gene organization and plasticity of the beta-lactam genes in different filamentous fungi. <i>Antonie Van Leeuwenhoek</i> , 1999, 75, 81-94.	0.7	61
97	Intrachromosomal recombination between direct repeats in <i>Penicillium chrysogenum</i> : gene conversion and deletion events. <i>Molecular Genetics and Genomics</i> , 1999, 261, 994-1000.	2.4	33
98	Characterization and lysine control of expression of the <i>lys1</i> gene of <i>Penicillium chrysogenum</i> encoding homocitrate synthase. <i>Gene</i> , 1999, 226, 51-59.	1.0	23
99	Molecular characterization of the <i>Acremonium chrysogenum</i> <i>cefG</i> gene product: the native deacetylcephalosporin C acetyltransferase is not processed into subunits. <i>Biochemical Journal</i> , 1999, 337, 379-385.	1.7	26
100	Molecular characterization of the <i>Acremonium chrysogenum</i> <i>cefG</i> gene product: the native deacetylcephalosporin C acetyltransferase is not processed into subunits. <i>Biochemical Journal</i> , 1999, 337, 379.	1.7	10
101	Thaumatin Production in <i>Aspergillus awamori</i> by Use of Expression Cassettes with Strong Fungal Promoters and High Gene Dosage. <i>Applied and Environmental Microbiology</i> , 1999, 65, 1168-1174.	1.4	53
102	Characterization of the <i>lys2</i> gene of <i>Penicillium chrysogenum</i> encoding $\hat{\pm}$ -aminoadipic acid reductase. <i>Molecular Genetics and Genomics</i> , 1998, 259, 549-556.	2.4	38
103	Characterization and nitrogen-source regulation at the transcriptional level of the <i>gdh A</i> gene of <i>Aspergillus awamori</i> encoding an NADP-dependent glutamate dehydrogenase. <i>Current Genetics</i> , 1998, 34, 50-59.	0.8	37
104	Characterization of the <i>bip</i> gene of <i>Aspergillus awamori</i> encoding a protein with an HDEL retention signal homologous to the mammalian BiP involved in polypeptide secretion. <i>Current Genetics</i> , 1997, 32, 139-146.	0.8	19
105	Expression of the <i>cefG</i> gene is limiting for cephalosporin biosynthesis in <i>Acremonium chrysogenum</i> . <i>Applied Microbiology and Biotechnology</i> , 1997, 48, 606-614.	1.7	74
106	Autonomously replicating plasmids carrying the <i>AMA1</i> region in <i>Penicillium chrysogenum</i> . <i>Current Genetics</i> , 1996, 29, 482-489.	0.8	65
107	Mutants blocked in penicillin biosynthesis show a deletion of the entire penicillin gene cluster at a specific site within a conserved hexanucleotide sequence. <i>Applied Microbiology and Biotechnology</i> , 1996, 44, 597-604.	1.7	90
108	Molecular genetics as a tool to remove bottlenecks in the biosynthesis of $\beta$ -lactam antibiotics. <i>World Journal of Microbiology and Biotechnology</i> , 1996, 12, 517-523.	1.7	2



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109	Mutants blocked in penicillin biosynthesis show a deletion of the entire penicillin gene cluster at a specific site within a conserved hexanucleotide sequence. <i>Applied Microbiology and Biotechnology</i> , 1996, 44, 597-604.	1.7	8
110	Autonomously replicating plasmids carrying the AMA1 region in <i>Penicillium chrysogenum</i> . <i>Current Genetics</i> , 1996, 29, 482-489.	0.8	4
111	The penicillin gene cluster is amplified in tandem repeats linked by conserved hexanucleotide sequences. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995, 92, 6200-6204.	3.3	174
112	Genes for $\beta$ -lactam antibiotic biosynthesis. <i>Antonie Van Leeuwenhoek</i> , 1995, 67, 181-200.	0.7	37
113	Three genes <i>hrdB</i> , <i>hrdD</i> and <i>hrdT</i> of <i>Streptomyces griseus</i> IMRU 3570, encoding sigma factor-like proteins, are differentially expressed under specific nutritional conditions. <i>Gene</i> , 1995, 153, 41-48.	1.0	17
114	Expression of genes and processing of enzymes for the biosynthesis of penicillins and cephalosporins. <i>Antonie Van Leeuwenhoek</i> , 1994, 65, 227-243.	0.7	19
115	Molecular characterization of three loss-of-function mutations in the isopenicillin N-acyltransferase gene ( <i>penDE</i> ) of <i>Penicillium chrysogenum</i> . <i>Journal of Bacteriology</i> , 1994, 176, 4941-4948.	1.0	28
116	Exogenous methionine increases levels of mRNAs transcribed from <i>pcbAB</i> , <i>pcbC</i> , and <i>cefEF</i> genes, encoding enzymes of the cephalosporin biosynthetic pathway, in <i>Acremonium chrysogenum</i> . <i>Journal of Bacteriology</i> , 1994, 176, 985-991.	1.0	53
117	Resolution of four large chromosomes in penicillin-producing filamentous fungi: the penicillin gene cluster is located on chromosome II (9.6 Mb) in <i>Penicillium notatum</i> and chromosome 1 (10.4 Mb) in <i>Penicillium chrysogenum</i> . <i>Molecular Genetics and Genomics</i> , 1993, 241-241, 573-578.	2.4	80
118	The isopenicillin-N acyltransferase of <i>Penicillium chrysogenum</i> has isopenicillin-N amidohydrolase, 6-aminopenicillanic acid acyltransferase and penicillin amidase activities, all of which are encoded by the single <i>penDE</i> gene. <i>FEBS Journal</i> , 1993, 215, 323-332.	0.2	80
119	The <i>cefG</i> gene of <i>Cephalosporium acremonium</i> is linked to the <i>cefEF</i> gene and encodes a deacetylcephalosporin C acetyltransferase closely related to homoserine O-acetyltransferase. <i>Journal of Bacteriology</i> , 1992, 174, 3056-3064.	1.0	92
120	Resolution of chromosomes III and VI of <i>Aspergillus nidulans</i> by pulsed-field gel electrophoresis shows that the penicillin biosynthetic pathway genes <i>pcbAB</i> , <i>pcbC</i> , and <i>penDE</i> are clustered on chromosome VI (3.0 megabases). <i>Journal of Bacteriology</i> , 1992, 174, 7063-7067.	1.0	47
121	Expression of the <i>penDE</i> gene of <i>Penicillium chrysogenum</i> encoding isopenicillin N acyltransferase in <i>Cephalosporium acremonium</i> : production of benzylpenicillin by the transformants. <i>Molecular Genetics and Genomics</i> , 1991, 225, 56-64.	2.4	63
122	Cloning, characterization of the acyl-CoA : 6-amino penicillanic acid acyltransferase gene of <i>Aspergillus nidulans</i> and linkage to the isopenicillin N synthase gene. <i>Molecular Genetics and Genomics</i> , 1990, 221, 322-330.	2.4	62
123	USO DE NUTRAGREEN® COMO TRANSPORTADOR COLOIDAL PARA REDUCIR EL USO DE FERTILIZANTES Y PESTICIDAS EN PERAL. , 0, , 220-227.		0
124	<em>Lippia</em> spp. Essential Oil as a Control Agent against <em>Acanthoscelides obtectus</em>, an Insect Pest in <em>Phaseolus vulgaris</em> Beans. , 0, , .		0