Katsuya Gomi

List of Publications by Year
in descending order

127	8,667	41344	45317
papers	citations	49 h-index	90 g-index

132	132	132	5724
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Regulation of gliotoxin biosynthesis and protection in Aspergillus species. PLoS Genetics, 2022, 18, e1009965.	3.5	16
2	Visualization of polypeptides including fragmented Î±-amylase in rice koji grains using mass spectrometry imaging. Journal of Bioscience and Bioengineering, 2022, 134, 34-40.	2.2	1
3	Metaproteomics reveals protein composition of multiple saccharifying enzymes in nongxiangxing daqu and jiangxiangxing daqu under different thermophilic temperatures. International Journal of Food Science and Technology, 2022, 57, 5102-5113.	2.7	12
4	Visualization of dipeptidyl peptidase B enzymatic reaction in rice koji using mass spectrometry imaging. Journal of Bioscience and Bioengineering, 2022, 134, 133-137.	2.2	3
5	Analysis of fermentation control factors on volatile compounds of primary microorganisms in Jiangâ€flavor Daqu. Journal of Food Biochemistry, 2022, 46, .	2.9	12
6	Enzymatic degradation of xyloglucans by Aspergillus species: a comparative view of this genus. Applied Microbiology and Biotechnology, 2021, 105, 2701-2711.	3.6	5
7	Induction and Repression of Hydrolase Genes in Aspergillus oryzae. Frontiers in Microbiology, 2021, 12, 677603.	3.5	21
8	Crucial role of the intracellular Î±-glucosidase MalT in the activation of the transcription factor AmyR essential for amylolytic gene expression in Aspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2021, 85, 2076-2083.	1.3	6
9	Expression profiles of amylolytic genes in AmyR and CreA transcription factor deletion mutants of the black koji mold Aspergillus luchuensis. Journal of Bioscience and Bioengineering, 2021, 132, 321-326.	2.2	2
10	Identification and distinct regulation of three di/tripeptide transporters in Aspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2021, 85, 452-463.	1.3	8
11	Mapping haze-komi on rice koji grains using Î²-glucuronidase expressing Aspergillus oryzae and mass spectrometry imaging. Journal of Bioscience and Bioengineering, 2020, 129, 296-301.	2.2	13
12	Efficient production of recombinant tannase in Aspergillus oryzae using an improved glucoamylase gene promoter. Journal of Bioscience and Bioengineering, 2020, 129, 150-154.	2.2	9
13	Alternative transcription start sites of the enolase-encoding gene enoA are stringently used in glycolytic/gluconeogenic conditions in Aspergillus oryzae. Current Genetics, 2020, 66, 729-747.	1.7	7
14	Regulatory mechanisms for amylolytic gene expression in the koji moldAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2019, 83, 1385-1401.	1.3	43
15	AtrR Is an Essential Determinant of Azole Resistance in Aspergillus fumigatus. MBio, 2019, 10, .	4.1	59
16	Chaperone complex formation of the transcription factor MalR involved in maltose utilization and amylolytic enzyme production in Aspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2018, 82, 827-835.	1.3	3
17	Heterologous Biosynthesis of Fungal Indole Sesquiterpene Sespendole. ChemBioChem, 2018, 19, 1492-1497.	2.6	18
18	Increased production of biomass-degrading enzymes by double deletion of creA and creB genes involved in carbon catabolite repression in Aspergillus oryzae. Journal of Bioscience and Bioengineering, 2018, 125, 141-147.	2.2	25

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19	Subcellular localization of aphidicolin biosynthetic enzymes heterologously expressed in Aspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2018, 82, 139-147.	1.3	5
20	Total Biosynthesis of Brassicicenes: Identification of a Key Enzyme for Skeletal Diversification. Organic Letters, 2018, 20, 6178-6182.	4.6	21
21	The C-terminal region of the yeast monocarboxylate transporter Jen1 acts as a glucose signalâ€“responding degron recognized by the Î±-arrestin Rod1. Journal of Biological Chemistry, 2018, 293, 10926-10936.	3.4	24
22	Heterologous Production of a Novel Cyclic Peptide Compound, KK-1, in Aspergillus oryzae. Frontiers in Microbiology, 2018, 9, 690.	3.5	16
23	The PDR-type ABC transporters AtrA and AtrG are involved in azole drug resistance in Aspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2018, 82, 1840-1848.	1.3	10
24	Nuclear exportâ€dependent degradation of the carbon catabolite repressor CreA is regulated by a region located near the Câ€terminus in Aspergillus oryzae. Molecular Microbiology, 2018, 110, 176-190.	2.5	18
25	Cellular responses to the expression of unstable secretory proteins in the filamentous fungus Aspergillus oryzae. Applied Microbiology and Biotechnology, 2017, 101, 2437-2446.	3.6	13
26	Quantitative evaluation of haze formation of koji and progression of internal haze by drying of koji during koji making. Journal of Bioscience and Bioengineering, 2017, 124, 62-70.	2.2	9
27	Improved Î±-Amylase Production by Dephosphorylation Mutation of CreD, an Arrestin-Like Protein Required for Glucose-Induced Endocytosis of Maltose Permease and Carbon Catabolite Derepression in Aspergillusoryzae. Applied and Environmental Microbiology, 2017, 83, .	3.1	14
28	Cell wall Î±-1,3-glucan prevents Î±-amylase adsorption onto fungal cell in submerged culture of Aspergillus oryzae. Journal of Bioscience and Bioengineering, 2017, 124, 47-53.	2.2	30
29	Self-excising Cre/mutant lox marker recycling system for multiple gene integrations and consecutive gene deletions in Aspergillus oryzae. Journal of Bioscience and Bioengineering, 2017, 123, 403-411.	2.2	49
30	Biosynthetic Machinery of Diterpene Pleuromutilin Isolated from Basidiomycete Fungi. ChemBioChem, 2017, 18, 2317-2322.	2.6	35
31	Characterization of Cell Wall Î±-1,3-Glucanâ€“Deficient Mutants in Aspergillus oryzae Isolated by a Screening Method Based on Their Sensitivities to Congo Red or Lysing Enzymes. Journal of Applied Glycoscience (1999), 2017, 64, 65-73.	0.7	3
32	A Novel Zn2-Cys6 Transcription Factor AtrR Plays a Key Role in an Azole Resistance Mechanism of Aspergillus fumigatus by Co-regulating cyp51A and cdr1B Expressions. PLoS Pathogens, 2017, 13, e1006096.	4.7	104
33	Genome mining approach for harnessing the cryptic gene cluster in Alternaria solani: production of PKSâ€“NRPS hybrid metabolite, didymellamide B. Tetrahedron Letters, 2016, 57, 2793-2796.	1.4	18
34	Multiple Oxidative Modifications in the Ophiobolin Biosynthesis: P450 Oxidations Found in Genome Mining. Organic Letters, 2016, 18, 1980-1983.	4.6	36
35	Genome sequence ofAspergillus luchuensisNBRC 4314. DNA Research, 2016, 23, 507-515.	3.4	48
36	Biosynthesis of Shearinine: Diversification of a Tandem Prenyl Moiety of Fungal Indole Diterpenes. Organic Letters, 2016, 18, 5026-5029.	4.6	39

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37	Increased enzyme production under liquid culture conditions in the industrial fungus Aspergillus oryzae by disruption of the genes encoding cell wall Î±-1,3-glucan synthase. Bioscience, Biotechnology and Biochemistry, 2016, 80, 1853-1863.	1.3	42
38	Unveiling the Biosynthetic Pathway of the Ribosomally Synthesized and Postâ€translationally Modified Peptide Ustiloxin B in Filamentous Fungi. Angewandte Chemie, 2016, 128, 8204-8207.	2.0	7
39	Unveiling the Biosynthetic Pathway of the Ribosomally Synthesized and Postâ€translationally Modified Peptide Ustiloxin B in Filamentous Fungi. Angewandte Chemie - International Edition, 2016, 55, 8072-8075.	13.8	76
40	Reconstitution of biosynthetic machinery of fungal polyketides: unexpected oxidations of biosynthetic intermediates by expression host. Bioscience, Biotechnology and Biochemistry, 2016, 80, 426-431.	1.3	23
41	Substantial decrease in cell wall Î±-1,3-glucan caused by disruption of the kexB gene encoding a subtilisin-like processing protease in Aspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2016, 80, 1781-1791.	1.3	10
42	Signaling pathways for stress responses and adaptation in Aspergillus species: stress biology in the post-genomic era. Bioscience, Biotechnology and Biochemistry, 2016, 80, 1667-1680.	1.3	65
43	The C2H2-type transcription factor, FlbC, is involved in the transcriptional regulation of Aspergillus oryzae glucoamylase and protease genes specifically expressed in solid-state culture. Applied Microbiology and Biotechnology, 2016, 100, 5859-5868.	3.6	23
44	Purification and enzymatic characterization of a novel Î²-1,6-glucosidase from Aspergillus oryzae. Journal of Bioscience and Bioengineering, 2016, 121, 259-264.	2.2	28
45	Purification and enzymatic characterization of secretory glycoside hydrolase family 3 (GH3) aryl Î²-glucosidases screened from Aspergillus oryzae genome. Journal of Bioscience and Bioengineering, 2015, 120, 614-623.	2.2	25
46	Unfolded protein response is required for Aspergillus oryzae growth under conditions inducing secretory hydrolytic enzyme production. Fungal Genetics and Biology, 2015, 85, 1-6.	2.1	21
47	Endocytosis of a maltose permease is induced when amylolytic enzyme production is repressed in Aspergillus oryzae. Fungal Genetics and Biology, 2015, 82, 136-144.	2.1	21
48	Use of a biosynthetic intermediate to explore the chemical diversity of pseudo-natural fungal polyketides. Nature Chemistry, 2015, 7, 737-743.	13.6	74
49	Reconstitution of Biosynthetic Machinery for the Synthesis of the Highly Elaborated Indole Diterpene Penitrem. Angewandte Chemie - International Edition, 2015, 54, 5748-5752.	13.8	101
50	Genome Mining for Sesterterpenes Using Bifunctional Terpene Synthases Reveals a Unified Intermediate of Di/Sesterterpenes. Journal of the American Chemical Society, 2015, 137, 11846-11853.	13.7	141
51	Biosynthetic Study on Antihypercholesterolemic Agent Phomoidride: General Biogenesis of Fungal Dimeric Anhydrides. Organic Letters, 2015, 17, 5658-5661.	4.6	34
52	Distinct mechanism of activation of two transcription factors, AmyR and MalR, involved in amylolytic enzyme production in Aspergillus oryzae. Applied Microbiology and Biotechnology, 2015, 99, 1805-1815.	3.6	38
53	Change in enzyme production by gradually drying culture substrate during solid-state fermentation. Journal of Bioscience and Bioengineering, 2015, 119, 674-677.	2.2	4
54	Heterologous expression of highly reducing polyketide synthase involved in betaenone biosynthesis. Chemical Communications, 2015, 51, 1878-1881.	4.1	67

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55	Response and Adaptation to Cell Wall Stress and Osmotic Stress in Aspergillus Species. , 2015, , 199-218.		2
56	Effects of codon optimization on the mRNA levels of heterologous genes in filamentous fungi. Applied Microbiology and Biotechnology, 2014, 98, 3859-3867.	3.6	26
57	Improved Î±-amylase production by Aspergillus oryzae after a double deletion of genes involved in carbon catabolite repression. Applied Microbiology and Biotechnology, 2014, 98, 335-343.	3.6	55
58	Rapid Reconstitution of Biosynthetic Machinery for Fungal Metabolites in Aspergillus oryzae: Total Biosynthesis of Aflatrem. ChemBioChem, 2014, 15, 2076-2080.	2.6	76
59	Strategies for Increasing the Production Level of Heterologous Proteins in Aspergillus oryzae. , 2014, , 149-164.		4
60	Identification of Ophiobolin F Synthase by a Genome Mining Approach: A Sesterterpene Synthase from Aspergillus clavatus. Organic Letters, 2013, 15, 594-597.	4.6	160
61	Rapid enzyme production and mycelial growth in solid-state fermentation using the non-airflow box. Journal of Bioscience and Bioengineering, 2013, 116, 585-590.	2.2	12
62	Reconstitution of Biosynthetic Machinery for Indole-Diterpene Paxilline in Aspergillus oryzae. Journal of the American Chemical Society, 2013, 135, 1260-1263.	13.7	170
63	Structurally Diverse Chaetophenol Productions Induced by Chemically Mediated Epigenetic Manipulation of Fungal Gene Expression. Organic Letters, 2013, 15, 3346-3349.	4.6	55
64	ASPERGILLUS LUCHUENSIS, AN INDUSTRIALLY IMPORTANT BLACK ASPERGILLUS IN EAST ASIA. PLoS ONE, 2013, 8, e63769.	2.5	167
65	Modified Cre-loxPRecombination in Aspergillus oryzae by Direct Introduction of Cre Recombinase for Marker Gene Rescue. Applied and Environmental Microbiology, 2012, 78, 4126-4133.	3.1	66
66	Transcripts of a heterologous gene encoding mite allergen Der f 7 are stabilized by codon optimization in Aspergillus oryzae. Applied Microbiology and Biotechnology, 2012, 96, 1275-1282.	3.6	24
67	Functional analysis of FarA transcription factor in the regulation of the genes encoding lipolytic enzymes and hydrophobic surface binding protein for the degradation of biodegradable plastics in Aspergillus oryzae. Journal of Bioscience and Bioengineering, 2012, 113, 549-555.	2.2	23
68	Uniform culture in solid-state fermentation with fungi and its efficient enzyme production. Journal of Bioscience and Bioengineering, 2011, 111, 300-305.	2.2	21
69	Construction of a thiamine pyrophosphate high-producing strain of Aspergillus oryzae by overexpression of three genes involved in thiamine biosynthesis. Journal of Bioscience and Bioengineering, 2011, 111, 388-390.	2.2	11
70	Identification of potential cell wall component that allows Taka-amylase A adsorption in submerged cultures of Aspergillus oryzae. Applied Microbiology and Biotechnology, 2011, 92, 961-969.	3.6	27
71	Total Biosynthesis of Diterpene Aphidicolin, a Specific Inhibitor of DNA Polymerase Î±: Heterologous Expression of Four Biosynthetic Genes inAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2011, 75, 1813-1817.	1.3	79
72	In silico Analysis of 3'-End-Processing Signals in Aspergillus oryzae Using Expressed Sequence Tags and Genomic Sequencing Data. DNA Research, 2011, 18, 189-200.	3.4	13

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73	Characterization and expression analysis of a maltose-utilizing (MAL) cluster in Aspergillus oryzae. Fungal Genetics and Biology, 2010, 47, 1-9.	2.1	40
74	Alternative Processing of Proproteins in AspergillikexBGene Disruptants under Hyperosmotic Conditions. Bioscience, Biotechnology and Biochemistry, 2009, 73, 40-46.	1.3	7
75	Crawler, a novel Tc1/mariner-type transposable element in Aspergillus oryzae transposes under stress conditions. Fungal Genetics and Biology, 2009, 46, 441-449.	2.1	31
76	Aspergillus oryzae atfA controls conidial germination and stress tolerance. Fungal Genetics and Biology, 2009, 46, 887-897.	2.1	65
77	ç¨®ã„ã„ã•–ã‹ã‚‰å°ƒãŒã‚ã‚‹ç³¸çŠ¶èŒå†ç”Ÿå¤®ã‚¬ã‚¤ãƒ‰ã‚¬ã‚¤è€æ€§æ©Ÿæ§‹. Kagaku To Seibutsu, 2009, 47, 684-689.	0.0	0
78	A defect of LigD (human Lig4 homolog) for nonhomologous end joining significantly improves efficiency of gene-targeting in Aspergillus oryzae. Fungal Genetics and Biology, 2008, 45, 878-889.	2.1	132
79	Aspergillus oryzae atfB encodes a transcription factor required for stress tolerance in conidia. Fungal Genetics and Biology, 2008, 45, 922-932.	2.1	62
80	Genomics of Aspergillus oryzae: Learning from the History of Koji Mold and Exploration of Its Future. DNA Research, 2008, 15, 173-183.	3.4	328
81	Codon Optimization Increases Steady-State mRNA Levels in Aspergillus oryzae Heterologous Gene Expression. Applied and Environmental Microbiology, 2008, 74, 6538-6546.	3.1	61
82	Genomics ofAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2007, 71, 646-670.	1.3	163
83	Analysis of Expressed Sequence Tags from the Fungus Aspergillus oryzae Cultured Under Different Conditions. DNA Research, 2007, 14, 47-57.	3.4	73
84	Novel Hydrophobic Surface Binding Protein, HsbA, Produced by Aspergillus oryzae. Applied and Environmental Microbiology, 2006, 72, 2407-2413.	3.1	82
85	The fungal hydrophobin RolA recruits polyesterase and laterally moves on hydrophobic surfaces. Molecular Microbiology, 2005, 57, 1780-1796.	2.5	71
86	Genome sequencing and analysis of Aspergillus oryzae. Nature, 2005, 438, 1157-1161.	27.8	1,128
87	Genomic sequence of the pathogenic and allergenic filamentous fungus Aspergillus fumigatus. Nature, 2005, 438, 1151-1156.	27.8	1,272
88	Purification and characterization of a biodegradable plastic-degrading enzyme from Aspergillus oryzae. Applied Microbiology and Biotechnology, 2005, 67, 778-788.	3.6	195
89	Genomics of Economically Significant Aspergillus and Fusarium Species. Applied Mycology and Biotechnology, 2004, 4, 249-283.	0.3	19
90	Transcriptional analysis of genes for energy catabolism and hydrolytic enzymes in the filamentous fungus Aspergillus oryzae using cDNA microarrays and expressed sequence tags. Applied Microbiology and Biotechnology, 2004, 65, 74-83.	3.6	84

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91	Thiamine-regulated gene expression of Aspergillus oryzae thiA requires splicing of the intron containing a riboswitch-like domain in the 5â€²-UTR. FEBS Letters, 2003, 555, 516-520.	2.8	195
92	Cloning, nucleotide sequencing, and expression of the .BETA.-galactosidase-encoding gene (lacA) from Aspergillus oryzae.. Journal of General and Applied Microbiology, 2002, 48, 135-142.	0.7	18
93	chsZ , a gene for a novel class of chitin synthase from Aspergillus oryzae. Current Genetics, 2002, 41, 261-267.	1.7	54
94	Subtractive cloning of cDNA from Aspergillus oryzae differentially regulated between solid-state culture and liquid (submerged) culture. Current Genetics, 2002, 41, 275-281.	1.7	58
95	Deletion analysis of the enolase gene (enoA) promoter from the filamentous fungus Aspergillus oryzae. Current Genetics, 2001, 40, 260-267.	1.7	33
96	Characterization of the amyR gene encoding a transcriptional activator for the amylase genes in Aspergillus nidulans. Current Genetics, 2001, 39, 10-15.	1.7	81
97	Rapid detection of homologously integrated DNA fragments and accurate quantitation of their copy number in transgenic Aspergillus oryzae by PCR. Journal of Bioscience and Bioengineering, 2000, 90, 577-579.	2.2	3
98	Molecular Cloning and Characterization of a Transcriptional Activator Gene,amyR, Involved in the Amylolytic Gene Expression inAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 2000, 64, 816-827.	1.3	114
99	Cloning and functional analysis of the Aspergillus oryzae conidiation regulator gene brlA by its disruption and misscheduled expression. Journal of Bioscience and Bioengineering, 1999, 87, 424-429.	2.2	60
100	Insertion Analysis of Putative Functional Elements in the Promoter Region of theAspergillus oryzaeTaka-amylase A Gene (amyB) Using a HeterologousAspergillus nidulans amdS-lacZFusion Gene System. Bioscience, Biotechnology and Biochemistry, 1999, 63, 180-183.	1.3	28
101	Molecular and enzymic properties of recombinant 1,2-Î±-mannosidase from Aspergillus saitoi overexpressed in Aspergillus oryzae cells. Biochemical Journal, 1999, 339, 589-597.	3.7	37
102	Molecular and enzymic properties of recombinant 1,2-Î±-mannosidase from Aspergillus saitoi overexpressed in Aspergillus oryzae cells. Biochemical Journal, 1999, 339, 589.	3.7	27
103	Transformation System forAspergillus oryzaewith Double Auxotrophic Mutations,niaDandsC. Bioscience, Biotechnology and Biochemistry, 1997, 61, 1367-1369.	1.3	130
104	Deletion analysis of promoter elements of the Aspergillus oryzae agdA gene encoding Î± -glucosidase. Current Genetics, 1996, 30, 432-438.	1.7	87
105	Cloning and Nucleotide Sequence of the Ribonuclease T₁Gene (rntA) fromAspergillus oryzaeand Its Expression inSaccharomyces cerevisiaeandAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 1995, 59, 1869-1874.	1.3	92
106	Molecular Cloning and Heterologous Expression of the Gene Encoding Dihydrogeodin Oxidase, a Multicopper Blue Enzyme from Aspergillus terreus. Journal of Biological Chemistry, 1995, 270, 21495-21502.	3.4	66
107	Nucleotide Sequence and Expression ofÎ±-Glucosidase-encoding Gene (agdA) fromAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 1995, 59, 1516-1521.	1.3	53
108	A Novel Culture Method for High Level Production of Heterologous Protein inSaccharomyces cerevisiae. Bioscience, Biotechnology and Biochemistry, 1994, 58, 1292-1296.	1.3	8

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109	Electrophoretic Karyotype and Gene Assignment to Chromosomes ofAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 1994, 58, 1467-1470.	1.3	24
110	High Level Secretion of Calf Chymosin Using a Glucoamylase-prochymosin Fusion Gene inAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 1994, 58, 895-899.	1.3	99
111	Secretion of calf chymosin from the filamentous fungus Aspergillus oryzae. Applied Microbiology and Biotechnology, 1993, 40, 327-32.	3.6	45
112	Cloning and Nucleotide Sequence of the Acid Protease-encoding Gene (pepA) fromAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 1993, 57, 1095-1100.	1.3	60
113	Deletion Analysis of the Taka-amylase A Gene Promoter Using a Homologous Transformation System inAspergillus oryzae. Bioscience, Biotechnology and Biochemistry, 1992, 56, 1849-1853.	1.3	57
114	Functional elements of the promoter region of the Aspergillus oryzae glaA gene encoding glucoamylase. Current Genetics, 1992, 22, 85-91.	1.7	81
115	High level expression of the synthetic human lysozyme gene in Aspergillus oryzae. Applied Microbiology and Biotechnology, 1992, 38, 109-14.	3.6	73
116	Transformation of the industrial strain of Aspergillus oryzae with the homologous amdS gene as a dominant selectable marker. Journal of Bioscience and Bioengineering, 1992, 74, 389-391.	0.9	33
117	3 Genetic Transfer Applied to Traditional Sake Brewing. Biotechnology and Genetic Engineering Reviews, 1991, 9, 89-125.	6.2	2
118	Identification of the Promoter Region of the Taka-amylase A Gene Required for Starch Induction. Agricultural and Biological Chemistry, 1991, 55, 1939-1941.	0.3	15
119	Construction of a fusion gene comprising the Taka-amylase A promoter and the Escherichia coli Î²-glucuronidase gene and analysis of its expression in Aspergillus oryzae. Molecular Genetics and Genomics, 1991, 229, 301-306.	2.4	89
120	The Glucoamylase cDNA fromAspergillus oryzae: Its Cloning, Nucleotide Sequence, and Expression inSaccharomyces cerevisiae. Agricultural and Biological Chemistry, 1991, 55, 941-949.	0.3	24
121	Identification of the Promoter Region of the Taka-amylase A Gene Required for Starch Induction.. Agricultural and Biological Chemistry, 1991, 55, 1939-1941.	0.3	19
122	Cloning and nucleotide sequence of the genomic Taka-amylase A gene of Aspergillus oryzae.. Agricultural and Biological Chemistry, 1989, 53, 593-599.	0.3	81
123	Transformation ofAspergillus oryzaethrough Plasmid-mediated Complementation of the Methionine-auxotrophic Mutation. Agricultural and Biological Chemistry, 1987, 51, 323-328.	0.3	2
124	Integrative transformation of Aspergillus oryzae with a plasmid containing the Aspergillus nidulans argB gene.. Agricultural and Biological Chemistry, 1987, 51, 2549-2555.	0.3	239
125	Integrative Transformation ofAspergillus oryzaewith a Plasmid Containing theAspergillus nidulans argBGene. Agricultural and Biological Chemistry, 1987, 51, 2549-2555.	0.3	118
126	Transformation of Aspergillus oryzae through plasmid-mediated complementation of the methionine-auxotrophic mutation.. Agricultural and Biological Chemistry, 1987, 51, 323-328.	0.3	63

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127	Studies on application of fungal cell wall lytic enzyme produced by Oerskovia sp. CK. (Part II) Estimation of mycelial weight in rice-koji with use of fungal cell wall lytic enzyme.. Journal of the Society of Brewing Japan, 1987, 82, 130-133.	0.0	17