

Merja Penttil

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220
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221
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ext. citations

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L-index

#	Paper	IF	Citations
220	A versatile transformation system for the cellulolytic filamentous fungus <i>Trichoderma reesei</i> . <i>Gene</i> , 1987 , 61, 155-64	3.8	509
219	A consensus yeast metabolic network reconstruction obtained from a community approach to systems biology. <i>Nature Biotechnology</i> , 2008 , 26, 1155-60	44.5	471
218	Hydrophobins: the protein-amphiphiles of filamentous fungi. <i>FEMS Microbiology Reviews</i> , 2005 , 29, 877-96	9.1	453
217	Traffic jams reduce hydrolytic efficiency of cellulase on cellulose surface. <i>Science</i> , 2011 , 333, 1279-82	33.3	439
216	Transcriptional regulation of plant cell wall degradation by filamentous fungi. <i>FEMS Microbiology Reviews</i> , 2005 , 29, 719-39	15.1	374
215	Metabolic engineering applications to renewable resource utilization. <i>Current Opinion in Biotechnology</i> , 2000 , 11, 187-98	11.4	350
214	Swollenin, a <i>Trichoderma reesei</i> protein with sequence similarity to the plant expansins, exhibits disruption activity on cellulosic materials. <i>FEBS Journal</i> , 2002 , 269, 4202-11		320
213	Homology between cellulase genes of <i>Trichoderma reesei</i> : complete nucleotide sequence of the endoglucanase I gene. <i>Gene</i> , 1986 , 45, 253-63	3.8	267
212	<i>Saccharomyces cerevisiae</i> engineered to produce D-xylonate. <i>Applied Microbiology and Biotechnology</i> , 2010 , 88, 751-60	5.7	252
211	Xylose chemostat isolates of <i>Saccharomyces cerevisiae</i> show altered metabolite and enzyme levels compared with xylose, glucose, and ethanol metabolism of the original strain. <i>Applied Microbiology and Biotechnology</i> , 2005 , 67, 827-37	5.7	250
210	Bioconversion of d-xylose to d-xylonate with <i>Kluyveromyces lactis</i> . <i>Metabolic Engineering</i> , 2011 , 13, 383-91	9.7	248
209	ACEII, a novel transcriptional activator involved in regulation of cellulase and xylanase genes of <i>Trichoderma reesei</i> . <i>Journal of Biological Chemistry</i> , 2001 , 276, 24309-14	5.4	243
208	Identification in the mould <i>Hypocrea jecorina</i> of a gene encoding an NADP(+): d-xylose dehydrogenase. <i>FEMS Microbiology Letters</i> , 2007 , 277, 249-53	2.9	230
207	High speed atomic force microscopy visualizes processive movement of <i>Trichoderma reesei</i> cellobiohydrolase I on crystalline cellulose. <i>Journal of Biological Chemistry</i> , 2009 , 284, 36186-36190	5.4	214
206	ACEI of <i>Trichoderma reesei</i> is a repressor of cellulase and xylanase expression. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 56-65	4.8	208
205	Atomic resolution structure of the HFBII hydrophobin, a self-assembling amphiphile. <i>Journal of Biological Chemistry</i> , 2004 , 279, 534-9	5.4	191
204	Conversion of xylose to ethanol by recombinant <i>Saccharomyces cerevisiae</i> : importance of xylulokinase (XKS1) and oxygen availability. <i>Metabolic Engineering</i> , 2001 , 3, 236-49	9.7	191

203	Engineering redox cofactor regeneration for improved pentose fermentation in <i>Saccharomyces cerevisiae</i> . <i>Applied and Environmental Microbiology</i> , 2003 , 69, 5892-7	4.8	167
202	Efficient secretion of two fungal cellobiohydrolases by <i>Saccharomyces cerevisiae</i> . <i>Gene</i> , 1988 , 63, 103-123,8		161
201	Xylitol production by recombinant <i>Saccharomyces cerevisiae</i> . <i>Bio/technology</i> , 1991 , 9, 1090-5		159
200	Screening of candidate regulators for cellulase and hemicellulase production in <i>Trichoderma reesei</i> and identification of a factor essential for cellulase production. <i>Biotechnology for Biofuels</i> , 2014 , 7, 14	7.8	158
199	Isolation of the <i>ace1</i> gene encoding a Cys(2)-His(2) transcription factor involved in regulation of activity of the cellulase promoter <i>cbh1</i> of <i>Trichoderma reesei</i> . <i>Journal of Biological Chemistry</i> , 2000 , 275, 5817-25	5.4	152
198	Genetic modification of carbon catabolite repression in <i>Trichoderma reesei</i> for improved protein production. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 4853-60	4.8	145
197	Xylose transport studies with xylose-utilizing <i>Saccharomyces cerevisiae</i> strains expressing heterologous and homologous permeases. <i>Applied Microbiology and Biotechnology</i> , 2007 , 74, 1041-52	5.7	141
196	Effects of inactivation and constitutive expression of the unfolded- protein response pathway on protein production in the yeast <i>Saccharomyces cerevisiae</i> . <i>Applied and Environmental Microbiology</i> , 2003 , 69, 2065-72	4.8	141
195	cDNA cloning of a <i>Trichoderma reesei</i> cellulase and demonstration of endoglucanase activity by expression in yeast. <i>FEBS Journal</i> , 1997 , 249, 584-91		138
194	Molecular cloning and analysis of the yeast flocculation gene <i>FLO1</i> . <i>Yeast</i> , 1994 , 10, 211-25	3.4	137
193	Expression patterns of ten hemicellulase genes of the filamentous fungus <i>Trichoderma reesei</i> on various carbon sources. <i>Journal of Biotechnology</i> , 1997 , 57, 167-179	3.7	135
192	Expression in <i>Trichoderma reesei</i> and characterisation of a thermostable family 3 beta-glucosidase from the moderately thermophilic fungus <i>Talaromyces emersonii</i> . <i>Protein Expression and Purification</i> , 2004 , 38, 248-57	2	134
191	Re-annotation of the CAZy genes of <i>Trichoderma reesei</i> and transcription in the presence of lignocellulosic substrates. <i>Microbial Cell Factories</i> , 2012 , 11, 134	6.4	124
190	Enzymatic properties and intracellular localization of the novel <i>Trichoderma reesei</i> beta-glucosidase BGLII (<i>cel1A</i>). <i>Applied and Environmental Microbiology</i> , 2002 , 68, 4546-53	4.8	122
189	The hydrophobins HFBI and HFBII from <i>Trichoderma reesei</i> showing efficient interactions with nonionic surfactants in aqueous two-phase systems. <i>Biomacromolecules</i> , 2001 , 2, 511-7	6.9	122
188	A novel, small endoglucanase gene, <i>egl5</i> , from <i>Trichoderma reesei</i> isolated by expression in yeast. <i>Molecular Microbiology</i> , 1994 , 13, 219-28	4.1	122
187	Interaction and comparison of a class I hydrophobin from <i>Schizophyllum commune</i> and class II hydrophobins from <i>Trichoderma reesei</i> . <i>Biomacromolecules</i> , 2006 , 7, 1295-301	6.9	121
186	High level secretion of cellobiohydrolases by <i>Saccharomyces cerevisiae</i> . <i>Biotechnology for Biofuels</i> , 2011 , 4, 30	7.8	119

185	Transformation system for <i>Hypocrea jecorina</i> (<i>Trichoderma reesei</i>) that favors homologous integration and employs reusable bidirectionally selectable markers. <i>Applied and Environmental Microbiology</i> , 2011 , 77, 114-21	4.8	119
184	Laccase from the white-rot fungus <i>Trametes versicolor</i> : cDNA cloning of <i>lcc1</i> and expression in <i>Pichia pastoris</i> . <i>Current Genetics</i> , 1997 , 32, 425-30	2.9	119
183	Yeast oligo-mediated genome engineering (YOGE). <i>ACS Synthetic Biology</i> , 2013 , 2, 741-9	5.7	116
182	Transcriptional regulation of <i>xyn1</i> , encoding xylanase I, in <i>Hypocrea jecorina</i> . <i>Eukaryotic Cell</i> , 2006 , 5, 447-56		116
181	The effects of drugs inhibiting protein secretion in the filamentous fungus <i>Trichoderma reesei</i> . Evidence for down-regulation of genes that encode secreted proteins in the stressed cells. <i>Journal of Biological Chemistry</i> , 2003 , 278, 45011-20	5.4	113
180	Activation mechanisms of the HAC1-mediated unfolded protein response in filamentous fungi. <i>Molecular Microbiology</i> , 2003 , 47, 1149-61	4.1	112
179	Efficient production of L-lactic acid from xylose by <i>Pichia stipitis</i> . <i>Applied and Environmental Microbiology</i> , 2007 , 73, 117-23	4.8	109
178	Structural hierarchy in molecular films of two class II hydrophobins. <i>Biochemistry</i> , 2003 , 42, 5253-8	3.2	109
177	Homologous expression and characterization of Cel61A (EG IV) of <i>Trichoderma reesei</i> . <i>FEBS Journal</i> , 2001 , 268, 6498-507		102
176	Efficient purification of recombinant proteins using hydrophobins as tags in surfactant-based two-phase systems. <i>Biochemistry</i> , 2004 , 43, 11873-82	3.2	101
175	Recent advances in the malting and brewing industry ¹ Based on a lecture held at the symposium ¹ Biotechnology in advanced food and feed processing [†] at the 8th European Congress on Biotechnology (ECB8) in Budapest, Hungary, August 1997.1. <i>Journal of Biotechnology</i> , 1998 , 65, 85-98	3.7	100
174	Surface adhesion of fusion proteins containing the hydrophobins HFBI and HFBII from <i>Trichoderma reesei</i> . <i>Protein Science</i> , 2002 , 11, 2257-66	6.3	99
173	Lipid production in batch and fed-batch cultures of <i>Rhodospiridium toruloides</i> from 5 and 6 carbon carbohydrates. <i>BMC Biotechnology</i> , 2012 , 12, 26	3.5	96
172	Monitoring of transcriptional regulation in <i>Pichia pastoris</i> under protein production conditions. <i>BMC Genomics</i> , 2007 , 8, 179	4.5	96
171	Improvement of foreign-protein production in <i>Aspergillus niger</i> var. <i>awamori</i> by constitutive induction of the unfolded-protein response. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 6979-86	4.8	96
170	Endogenous xylose pathway in <i>Saccharomyces cerevisiae</i> . <i>Applied and Environmental Microbiology</i> , 2004 , 70, 3681-6	4.8	93
169	Oxygen dependence of metabolic fluxes and energy generation of <i>Saccharomyces cerevisiae</i> CEN.PK113-1A. <i>BMC Systems Biology</i> , 2008 , 2, 60	3.5	91
168	Metabolic flux analysis of xylose metabolism in recombinant <i>Saccharomyces cerevisiae</i> using continuous culture. <i>Metabolic Engineering</i> , 2003 , 5, 16-31	9.7	90

167	Enzyme production by recombinant <i>Trichoderma reesei</i> strains. <i>Journal of Biotechnology</i> , 1991 , 17, 35-49.	3.7	90
166	Production of ethanol from L-arabinose by <i>Saccharomyces cerevisiae</i> containing a fungal L-arabinose pathway. <i>FEMS Yeast Research</i> , 2003 , 3, 185-9	3.1	89
165	Glycolic acid production in the engineered yeasts <i>Saccharomyces cerevisiae</i> and <i>Kluyveromyces lactis</i> . <i>Microbial Cell Factories</i> , 2013 , 12, 82	6.4	84
164	Swollenin aids in the amorphogenesis step during the enzymatic hydrolysis of pretreated biomass. <i>Bioresource Technology</i> , 2013 , 142, 498-503	11	83
163	The effect of specific growth rate on protein synthesis and secretion in the filamentous fungus <i>Trichoderma reesei</i> . <i>Microbiology (United Kingdom)</i> , 2005 , 151, 135-143	2.9	83
162	The three-dimensional structure of a <i>Trichoderma reesei</i> beta-mannanase from glycoside hydrolase family 5. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2000 , 56, 3-13		80
161	Genetic and biochemical characterization of the <i>Trichoderma reesei</i> hydrophobin HFBI. <i>FEBS Journal</i> , 1996 , 235, 248-55		80
160	Integrated multilaboratory systems biology reveals differences in protein metabolism between two reference yeast strains. <i>Nature Communications</i> , 2010 , 1, 145	17.4	78
159	The role of xylulokinase in <i>Saccharomyces cerevisiae</i> xylulose catabolism. <i>FEMS Microbiology Letters</i> , 2000 , 190, 39-43	2.9	77
158	Cloning and expression of a fungal L-arabinitol 4-dehydrogenase gene. <i>Journal of Biological Chemistry</i> , 2001 , 276, 40631-7	5.4	76
157	The missing link in the fungal L-arabinose catabolic pathway, identification of the L-xylulose reductase gene. <i>Biochemistry</i> , 2002 , 41, 6432-7	3.2	76
156	Acetyl xylan esterase from <i>Trichoderma reesei</i> contains an active-site serine residue and a cellulose-binding domain. <i>FEBS Journal</i> , 1996 , 237, 553-60		76
155	Common features and interesting differences in transcriptional responses to secretion stress in the fungi <i>Trichoderma reesei</i> and <i>Saccharomyces cerevisiae</i> . <i>BMC Genomics</i> , 2006 , 7, 32	4.5	75
154	Genetic engineering of <i>Trichoderma</i> to produce strains with novel cellulase profiles. <i>Enzyme and Microbial Technology</i> , 1991 , 13, 227-33	3.8	75
153	Differential expression of the vegetative and spore-bound hydrophobins of <i>Trichoderma reesei</i> -cloning and characterization of the hfb2 gene. <i>FEBS Journal</i> , 1997 , 248, 415-23		74
152	Regulation of xylose metabolism in recombinant <i>Saccharomyces cerevisiae</i> . <i>Microbial Cell Factories</i> , 2008 , 7, 18	6.4	74
151	Evidence that the gene YLR070c of <i>Saccharomyces cerevisiae</i> encodes a xylitol dehydrogenase. <i>FEBS Letters</i> , 1999 , 457, 135-8	3.8	74
150	Challenges in enzymatic hydrolysis and fermentation of pretreated <i>Arundo donax</i> revealed by a comparison between SHF and SSF. <i>Process Biochemistry</i> , 2012 , 47, 1452-1459	4.8	73

149	Production of recombinant proteins in the filamentous fungus <i>Trichoderma reesei</i> . <i>Current Opinion in Biotechnology</i> , 1995 , 6, 534-7	11.4	73
148	Three alpha-galactosidase genes of <i>Trichoderma reesei</i> cloned by expression in yeast. <i>FEBS Journal</i> , 1996 , 240, 104-11		72
147	Microbial D-xylonate production. <i>Applied Microbiology and Biotechnology</i> , 2012 , 96, 1-8	5.7	71
146	Efficient production of antibody fragments by the filamentous fungus <i>Trichoderma reesei</i> . <i>Nature Biotechnology</i> , 1993 , 11, 591-5	44.5	70
145	Large-scale separation and production of engineered proteins, designed for facilitated recovery in detergent-based aqueous two-phase extraction systems. <i>Process Biochemistry</i> , 2004 , 39, 889-896	4.8	69
144	Array comparative genomic hybridization analysis of <i>Trichoderma reesei</i> strains with enhanced cellulase production properties. <i>BMC Genomics</i> , 2010 , 11, 441	4.5	68
143	The <i>Trichoderma reesei</i> hydrophobin genes <i>hfb1</i> and <i>hfb2</i> have diverse functions in fungal development. <i>FEMS Microbiology Letters</i> , 2005 , 253, 281-8	2.9	67
142	Metabolic engineering of <i>Saccharomyces cerevisiae</i> for bioconversion of D-xylose to D-xylonate. <i>Metabolic Engineering</i> , 2012 , 14, 427-36	9.7	65
141	Endoplasmic reticulum stress leads to the selective transcriptional downregulation of the glucoamylase gene in <i>Aspergillus niger</i> . <i>Molecular Microbiology</i> , 2004 , 53, 1731-42	4.1	65
140	Swollenin from <i>Trichoderma reesei</i> exhibits hydrolytic activity against cellulosic substrates with features of both endoglucanases and cellobiohydrolases. <i>Bioresource Technology</i> , 2015 , 181, 105-13	11	60
139	Comparative genome-scale reconstruction of gapless metabolic networks for present and ancestral species. <i>PLoS Computational Biology</i> , 2014 , 10, e1003465	5	59
138	Central carbon metabolism of <i>Saccharomyces cerevisiae</i> in anaerobic, oxygen-limited and fully aerobic steady-state conditions and following a shift to anaerobic conditions. <i>FEMS Yeast Research</i> , 2008 , 8, 140-54	3.1	59
137	Xylanase XYN IV from <i>Trichoderma reesei</i> showing exo- and endo-xylanase activity. <i>FEBS Journal</i> , 2013 , 280, 285-301	5.7	58
136	Metabolic engineering of fungal strains for conversion of D-galacturonate to meso-galactarate. <i>Applied and Environmental Microbiology</i> , 2010 , 76, 169-75	4.8	58
135	Low pH D-xylonate production with <i>Pichia kudriavzevii</i> . <i>Bioresource Technology</i> , 2013 , 133, 555-62	11	57
134	Cloning and characterization of the glucosidase II alpha subunit gene of <i>Trichoderma reesei</i> : a frameshift mutation results in the aberrant glycosylation profile of the hypercellulolytic strain Rut-C30. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 2910-24	4.8	57
133	Role of Ace2 (Activator of Cellulases 2) within the <i>xyn2</i> transcriptosome of <i>Hypocrea jecorina</i> . <i>Fungal Genetics and Biology</i> , 2008 , 45, 436-45	3.9	56
132	A novel NADH-linked l-xylulose reductase in the l-arabinose catabolic pathway of yeast. <i>Journal of Biological Chemistry</i> , 2004 , 279, 14746-51	5.4	55

131	Correlation of gene expression and protein production rate - a system wide study. <i>BMC Genomics</i> , 2011 , 12, 616	4.5	54
130	Pilot scale production of a heterologous <i>Trichoderma reesei</i> cellulase by <i>Saccharomyces cerevisiae</i> . <i>Journal of Biotechnology</i> , 1990 , 13, 267-78	3.7	54
129	Production and applications of carbohydrate-derived sugar acids as generic biobased chemicals. <i>Critical Reviews in Biotechnology</i> , 2016 , 36, 904-16	9.4	53
128	Spatially segregated SNARE protein interactions in living fungal cells. <i>Journal of Biological Chemistry</i> , 2007 , 282, 22775-85	5.4	53
127	Role of the bga1-encoded extracellular {beta}-galactosidase of <i>Hypocrea jecorina</i> in cellulase induction by lactose. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 851-7	4.8	52
126	Molecular cloning and enzymatic characterization of a <i>Trichoderma reesei</i> 1,2-alpha-D-mannosidase. <i>Journal of Biotechnology</i> , 2000 , 77, 255-63	3.7	52
125	Identification of the first fungal NADP-GAPDH from <i>Kluyveromyces lactis</i> . <i>Biochemistry</i> , 2002 , 41, 13833-32	3.2	51
124	Proteome analysis of recombinant xylose-fermenting <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 2003 , 20, 295-314	3.4	46
123	Isolation of <i>Trichoderma reesei</i> genes highly expressed on glucose-containing media: characterization of the tef1 gene encoding translation elongation factor 1 alpha. <i>Gene</i> , 1993 , 136, 313-8	3.8	46
122	Direct identification of hydrophobins and their processing in <i>Trichoderma</i> using intact-cell MALDI-TOF MS. <i>FEBS Journal</i> , 2007 , 274, 841-52	5.7	45
121	Enabling Low Cost Biopharmaceuticals: A Systematic Approach to Delete Proteases from a Well-Known Protein Production Host <i>Trichoderma reesei</i> . <i>PLoS ONE</i> , 2015 , 10, e0134723	3.7	45
120	Production of ethylene glycol or glycolic acid from D-xylose in <i>Saccharomyces cerevisiae</i> . <i>Applied Microbiology and Biotechnology</i> , 2017 , 101, 8151-8163	5.7	43
119	Low oxygen levels as a trigger for enhancement of respiratory metabolism in <i>Saccharomyces cerevisiae</i> . <i>BMC Genomics</i> , 2009 , 10, 461	4.5	43
118	Engineering filamentous fungi for conversion of D-galacturonic acid to L-galactonic acid. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 8676-83	4.8	42
117	Transcription analysis of recombinant <i>saccharomyces cerevisiae</i> reveals novel responses to xylose. <i>Applied Biochemistry and Biotechnology</i> , 2006 , 128, 237-61	3.2	42
116	The missing link in the fungal D-galacturonate pathway: identification of the L-threo-3-deoxy-hexulosonate aldolase. <i>Journal of Biological Chemistry</i> , 2007 , 282, 26195-201	5.4	41
115	A novel two-step extraction method with detergent/polymer systems for primary recovery of the fusion protein endoglucanase I-hydrophobin I. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2002 , 1569, 139-50	4	41
114	A universal gene expression system for fungi. <i>Nucleic Acids Research</i> , 2018 , 46, e111	20.1	41

113	Identification in the mold <i>Hypocrea jecorina</i> of the first fungal D-galacturonic acid reductase. <i>Biochemistry</i> , 2005 , 44, 11234-40	3.2	40
112	Identification in <i>Agrobacterium tumefaciens</i> of the D-galacturonic acid dehydrogenase gene. <i>Applied Microbiology and Biotechnology</i> , 2010 , 86, 901-9	5.7	39
111	Use of matrix-assisted laser desorption/ionization time-of-flight mass mapping and nanospray liquid chromatography/electrospray ionization tandem mass spectrometry sequence tag analysis for high sensitivity identification of yeast proteins separated by two-dimensional gel electrophoresis. <i>Rapid Communications in Mass Spectrometry</i> , 2001 , 15, 1685-92	2.2	39
110	Overexpression of PAD1 and FDC1 results in significant cinnamic acid decarboxylase activity in <i>Saccharomyces cerevisiae</i> . <i>AMB Express</i> , 2015 , 5, 12	4.1	38
109	Comparison of protein coding gene contents of the fungal phyla Pezizomycotina and Saccharomycotina. <i>BMC Genomics</i> , 2007 , 8, 325	4.5	38
108	Dual relationships of xylitol and alcohol dehydrogenases in families of two protein types. <i>FEBS Letters</i> , 1993 , 324, 9-14	3.8	38
107	Influence of growth temperature on the production of antibody Fab fragments in different microbes: a host comparative analysis. <i>Biotechnology Progress</i> , 2011 , 27, 38-46	2.8	37
106	L-galactonate dehydratase is part of the fungal path for D-galacturonic acid catabolism. <i>Molecular Microbiology</i> , 2006 , 61, 1060-8	4.1	37
105	Production of L-lactic acid by the yeast <i>Candida sonorensis</i> expressing heterologous bacterial and fungal lactate dehydrogenases. <i>Microbial Cell Factories</i> , 2013 , 12, 53	6.4	36
104	Identification of an L-arabinose reductase gene in <i>Aspergillus niger</i> and its role in L-arabinose catabolism. <i>Journal of Biological Chemistry</i> , 2010 , 285, 23622-8	5.4	36
103	Rapid and multiplexed transcript analysis of microbial cultures using capillary electrophoresis-detectable oligonucleotide probe pools. <i>Journal of Microbiological Methods</i> , 2006 , 65, 404-16	2.8	35
102	Transcriptional monitoring of steady state and effects of anaerobic phases in chemostat cultures of the filamentous fungus <i>Trichoderma reesei</i> . <i>BMC Genomics</i> , 2006 , 7, 247	4.5	34
101	The use of carbohydrate binding modules (CBMs) to monitor changes in fragmentation and cellulose fiber surface morphology during cellulase- and Swollenin-induced deconstruction of lignocellulosic substrates. <i>Journal of Biological Chemistry</i> , 2015 , 290, 2938-45	5.4	33
100	Metabolic engineering of the fungal D-galacturonate pathway for L-ascorbic acid production. <i>Microbial Cell Factories</i> , 2015 , 14, 2	6.4	33
99	Enzymes for the NADPH-dependent reduction of dihydroxyacetone and D-glyceraldehyde and L-glyceraldehyde in the mould <i>Hypocrea jecorina</i> . <i>FEBS Journal</i> , 2006 , 273, 4229-35	5.7	33
98	In vivo synthesis of complex N-glycans by expression of human N-acetylglucosaminyltransferase I in the filamentous fungus <i>Trichoderma reesei</i> . <i>FEBS Letters</i> , 1999 , 452, 365-70	3.8	33
97	Protein production and induction of the unfolded protein response in <i>Trichoderma reesei</i> strain Rut-C30 and its transformant expressing endoglucanase I with a hydrophobic tag. <i>Biotechnology and Bioengineering</i> , 2005 , 89, 335-44	4.9	32
96	Monitoring the kinetics of glycoprotein synthesis and secretion in the filamentous fungus <i>Trichoderma reesei</i> : cellobiohydrolase I (CBHI) as a model protein. <i>Microbiology (United Kingdom)</i> , 2000 , 146 (Pt 1), 223-232	2.9	32

95	Expression of <i>Trichoderma reesei</i> cellulases CBHI and EGI in <i>Ashbya gossypii</i> . <i>Applied Microbiology and Biotechnology</i> , 2010 , 87, 1437-46	5.7	31
94	Capillary electrophoresis for the monitoring of carboxylic acid production by <i>Gluconobacter oxydans</i> . <i>Journal of Chromatography A</i> , 2010 , 1217, 1537-42	4.5	31
93	Phase transitions as intermediate steps in the formation of molecularly engineered protein fibers. <i>Communications Biology</i> , 2018 , 1, 86	6.7	31
92	Engineering chimeric thermostable GH7 cellobiohydrolases in <i>Saccharomyces cerevisiae</i> . <i>Applied Microbiology and Biotechnology</i> , 2014 , 98, 2991-3001	5.7	30
91	The diverse role of Pdr12 in resistance to weak organic acids. <i>Yeast</i> , 2014 , 31, 219-32	3.4	30
90	Identification and quantitation of phosphorus metabolites in yeast neutral pH extracts by nuclear magnetic resonance spectroscopy. <i>Analytical Biochemistry</i> , 1999 , 272, 71-9	3.1	30
89	Metabolic engineering of <i>Saccharomyces cerevisiae</i> for conversion of D-glucose to xylitol and other five-carbon sugars and sugar alcohols. <i>Applied and Environmental Microbiology</i> , 2007 , 73, 5471-6	4.8	29
88	Expression of a fungal hydrophobin in the <i>Saccharomyces cerevisiae</i> cell wall: effect on cell surface properties and immobilization. <i>Applied and Environmental Microbiology</i> , 2002 , 68, 3385-91	4.8	29
87	Noninvasive high-throughput single-cell analysis of the intracellular pH of <i>Saccharomyces cerevisiae</i> by ratiometric flow cytometry. <i>Applied and Environmental Microbiology</i> , 2013 , 79, 7179-87	4.8	28
86	Expression of <i>Vitreoscilla</i> hemoglobin improves the metabolism of xylose in recombinant yeast <i>Saccharomyces cerevisiae</i> under low oxygen conditions. <i>Enzyme and Microbial Technology</i> , 2006 , 39, 6-14 ^{3.8}	3.8	28
85	Characterization of secretory genes <i>ypt1/yptA</i> and <i>nsf1/nsfA</i> from two filamentous fungi: induction of secretory pathway genes of <i>Trichoderma reesei</i> under secretion stress conditions. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 459-67	4.8	28
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