Masafumi Yohda

List of Publications by Citations

252 papers	5,523 citations	40 h-index	63 g-index
269 ext. papers	5,976 ext. citations	4.5 avg, IF	5.05 L-index

#	Paper	IF	Citations
252	Novel non-heme iron center of nitrile hydratase with a claw setting of oxygen atoms. *Nature Structural Biology*, **1998**, 5, 347-51		315
251	Heat-inactivated proteins are rescued by the DnaK.J-GrpE set and ClpB chaperones. *Proceedings of the National Academy of Sciences of the United States of America*, **1999**, 96, 7184-9	11.5	223
250	Chaperonin-mediated stabilization and ATP-triggered release of semiconductor nanoparticles. *Nature*, **2003**, 423, 628-32	50.4	211
249	Post-translational modification is essential for catalytic activity of nitrile hydratase. *Protein Science*, **2000**, 9, 1024-30	6.3	147
248	Sequence and over-expression of subunits of adenosine triphosphate synthase in thermophilic bacterium PS3. *Biochimica Et Biophysica Acta - Bioenergetics*, **1988**, 933, 141-55	4.6	136
247	Fe-type nitrile hydratase. *Journal of Inorganic Biochemistry*, **2001**, 83, 247-53	4.2	113
246	Activity Regulation of Photoreactive Nitrile Hydratase by Nitric Oxide. *Journal of the American Chemical Society*, **1997**, 119, 3785-3791	16.4	108
245	Cyclic RGD peptide-labeled upconversion nanophosphors for tumor cell-targeted imaging. *Biochemical and Biophysical Research Communications*, **2009**, 381, 54-8	3.4	94
244	Functional expression of nitrile hydratase in Escherichia coli: requirement of a nitrile hydratase activator and post-translational modification of a ligand cysteine. *Journal of Biochemistry*, **1999**, 125, 696-704	3.1	87
243	Crystal structures of the group II chaperonin from Thermococcus strain KS-1: steric hindrance by the substituted amino acid, and inter-subunit rearrangement between two crystal forms. *Journal of Molecular Biology*, **2004**, 335, 1265-78	6.5	80
242	Crystal Structures of the Lumazine Protein from Photobacterium kishitanii in Complexes with the Authentic Chromophore, 6,7-Dimethyl-8-(1?- d -Ribityl) Lumazine, and Its Analogues, Riboflavin and Flavin Mononucleotide, at High Resolution. *Journal of Bacteriology*, **2010**, 192, 1749-1749	3.5	78
241	Structure of the photoreactive iron center of the nitrile hydratase from Rhodococcus sp. N-771. Evidence of a novel post-translational modification in the cysteine ligand. *Journal of Biological Chemistry*, **1997**, 272, 29454-9	5.4	78
240	An enzyme controlled by light: the molecular mechanism of photoreactivity in nitrile hydratase. *Trends in Biotechnology*, **1999**, 17, 244-8	15.1	77
239	Structural and functional characterization of homo-oligomeric complexes of alpha and beta chaperonin subunits from the hyperthermophilic archaeum Thermococcus strain KS-1. *Journal of Molecular Biology*, **1997**, 273, 635-45	6.5	72
238	Effects of linear polyacrylamide concentrations and applied voltages on the separation of oligonucleotides and DNA sequencing fragments by capillary electrophoresis. *Analytical Chemistry*, **1994**, 66, 4243-52	7.8	70
237	Carbonyl sulfide hydrolase from Thiobacillus thioparus strain THI115 is one of the β-carbonic anhydrase family enzymes. *Journal of the American Chemical Society*, **2013**, 135, 3818-25	16.4	69
236	Tertiary and quaternary structures of photoreactive Fe-type nitrile hydratase from Rhodococcus sp. N-771: roles of hydration water molecules in stabilizing the structures and the structural origin of the substrate specificity of the enzyme. *Biochemistry*, **1999**, 38, 9887-98	3.2	67

235	Site-directed mutagenesis of stable adenosine triphosphate synthase. *Biochimica Et Biophysica Acta - Bioenergetics*, **1988**, 933, 156-64	4.6	67
234	Crystal structure of aspartate racemase from Pyrococcus horikoshii OT3 and its implications for molecular mechanism of PLP-independent racemization. *Journal of Molecular Biology*, **2002**, 319, 479-89	6.5	64
233	Structure of thiocyanate hydrolase: a new nitrile hydratase family protein with a novel five-coordinate cobalt(III) center. *Journal of Molecular Biology*, **2007**, 366, 1497-509	6.5	62
232	A novel factor required for the assembly of the DnaK and DnaJ chaperones of Thermus thermophilus. *Journal of Biological Chemistry*, **1996**, 271, 17343-8	5.4	58
231	High speed polymerase chain reaction in constant flow. *Bioscience, Biotechnology and Biochemistry*, **1994**, 58, 349-52	2.1	58
230	In vitro mutated beta subunits from the F1-ATPase of the thermophilic bacterium, PS3, containing glutamine in place of glutamic acid in positions 190 or 201 assembles with the alpha and gamma subunits to produce inactive complexes. *Biochemical and Biophysical Research Communications*, **1987**, 146, 705-10	3.4	58
229	Catalytic mechanism of nitrile hydratase proposed by time-resolved X-ray crystallography using a novel substrate, tert-butylisonitrile. *Journal of Biological Chemistry*, **2008**, 283, 36617-23	5.4	55
228	Packaging guest proteins into the encapsulin nanocompartment from Rhodococcus erythropolis N771. *Biotechnology and Bioengineering*, **2015**, 112, 13-20	4.9	52
227	Formation of highly toxic soluble amyloid beta oligomers by the molecular chaperone prefoldin. *FEBS Journal*, **2008**, 275, 5982-93	5.7	51
226	Kinetics and binding sites for interaction of the prefoldin with a group II chaperonin: contiguous non-native substrate and chaperonin binding sites in the archaeal prefoldin. *Journal of Biological Chemistry*, **2004**, 279, 31788-95	5.4	51
225	Pyrococcus prefoldin stabilizes protein-folding intermediates and transfers them to chaperonins for correct folding. *Biochemical and Biophysical Research Communications*, **2002**, 291, 769-74	3.4	50
224	Nonequivalence observed for the 16-meric structure of a small heat shock protein, SpHsp16.0, from Schizosaccharomyces pombe. *Structure*, **2013**, 21, 220-8	5.2	49
223	Structure and direct electrochemistry of cytochrome P450 from the thermoacidophilic crenarchaeon, Sulfolobus tokodaii strain 7. *Journal of Inorganic Biochemistry*, **2004**, 98, 1194-9	4.2	48
222	Development of a novel method for operating magnetic particles, Magtration Technology, and its use for automating nucleic acid purification. *Journal of Bioscience and Bioengineering*, **2001**, 91, 500-503	3.3	47
221	Structure and characterization of amidase from Rhodococcus sp. N-771: Insight into the molecular mechanism of substrate recognition. *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2010**, 1804, 184-92	4	46
220	Structure and molecular dynamics simulation of archaeal prefoldin: the molecular mechanism for binding and recognition of nonnative substrate proteins. *Journal of Molecular Biology*, **2008**, 376, 1130-41	6.5	46
219	Arginine 56 mutation in the beta subunit of nitrile hydratase: importance of hydrogen bonding to the non-heme iron center. *Journal of Inorganic Biochemistry*, **2000**, 80, 283-8	4.2	45
218	Molecular cloning and nucleotide sequencing of the aspartate racemase gene from lactic acid bacteria Streptococcus thermophilus. *Biochimica Et Biophysica Acta Gene Regulatory Mechanisms*, **1991**, 1089, 234-40		45

217	Facilitated release of substrate protein from prefoldin by chaperonin. *FEBS Letters*, **2005**, 579, 3718-24	3.8	43
216	Thiocyanate hydrolase is a cobalt-containing metalloenzyme with a cysteine-sulfinic acid ligand. *Journal of the American Chemical Society*, **2006**, 128, 728-9	16.4	43
215	Archaeal group II chaperonin mediates protein folding in the cis-cavity without a detachable GroES-like co-chaperonin. *Journal of Molecular Biology*, **2002**, 315, 73-85	6.5	43
214	Cobalt-substituted Fe-type nitrile hydratase of Rhodococcus sp. N-771. *FEBS Letters*, **2000**, 465, 173-7	3.8	42
213	ATP binding is critical for the conformational change from an open to closed state in archaeal group II chaperonin. *Journal of Biological Chemistry*, **2003**, 278, 44959-65	5.4	41
212	Localization of prefoldin interaction sites in the hyperthermophilic group II chaperonin and correlations between binding rate and protein transfer rate. *Journal of Molecular Biology*, **2006**, 364, 110-20	6.5	40
211	Natural chaperonin of the hyperthermophilic archaeum, Thermococcus strain KS-1: a hetero-oligomeric chaperonin with variable subunit composition. *Molecular Microbiology*, **2001**, 39, 1406-13	4.1	38
210	Role of the helical protrusion in the conformational change and molecular chaperone activity of the archaeal group II chaperonin. *Journal of Biological Chemistry*, **2004**, 279, 18834-9	5.4	37
209	Structural basis for catalytic activation of thiocyanate hydrolase involving metal-ligated cysteine modification. *Journal of the American Chemical Society*, **2009**, 131, 14838-43	16.4	35
208	Occurrence of free D-amino acids and aspartate racemases in hyperthermophilic archaea. *Journal of Bacteriology*, **1999**, 181, 6560-3	3.5	33
207	Vapor detection and discrimination with a panel of odorant receptors. *Nature Communications*, **2018**, 9, 4556	17.4	33
206	Distribution and purification of aspartate racemase in lactic acid bacteria. *BBA - Proteins and Proteomics*, **1991**, 1078, 377-82		32
205	Photoreactive nitrile hydratase: the photoreaction site is located on the alpha subunit. *Journal of Biochemistry*, **1996**, 119, 407-13	3.1	31
204	Single-site catalysis of F1-ATPase from thermophilic bacterium PS3 and its dominance in steady-state catalysis at low ATP concentration. *Journal of Biochemistry*, **1987**, 102, 875-83	3.1	31
203	Dimer structure and conformational variability in the N-terminal region of an archaeal small heat shock protein, StHsp14.0. *Journal of Structural Biology*, **2011**, 174, 92-9	3.4	30
202	Gene for aspartate racemase from the sulfur-dependent hyperthermophilic archaeum, Desulfurococcus strain SY. *Journal of Biological Chemistry*, **1996**, 271, 22017-21	5.4	30
201	Role of the IXI/V motif in oligomer assembly and function of StHsp14.0, a small heat shock protein from the acidothermophilic archaeon, Sulfolobus tokodaii strain 7. *Proteins: Structure, Function and Bioinformatics*, **2008**, 71, 771-82	4.2	29
200	Molecular characterization of the group II chaperonin from the hyperthermophilic archaeum Pyrococcus horikoshii OT3. *Extremophiles*, **2005**, 9, 127-34	3	29

199	Characterization of archaeal group II chaperonin-ADP-metal fluoride complexes: implications that group II chaperonins operate as a "two-stroke engine". *Journal of Biological Chemistry*, **2005**, 280, 40375-83	5.4	29
198	Occurrence of D-Amino Acids and a pyridoxal 5Pphosphate-dependent aspartate racemase in the acidothermophilic archaeon, Thermoplasma acidophilum. *Biochemical and Biophysical Research Communications*, **2001**, 281, 317-21	3.4	29
197	Glycine at the 65th position plays an essential role in ATP-dependent protein folding by Archael group II chaperonin. *Biochemical and Biophysical Research Communications*, **2001**, 289, 1118-24	3.4	29
196	Location of the non-heme iron center on the alpha subunit of photoreactive nitrile hydratase from Rhodococcus sp. N-771. *Biochemical and Biophysical Research Communications*, **1996**, 221, 146-50	3.4	29
195	Structural insight into gene duplication, gene fusion and domain swapping in the evolution of PLP-independent amino acid racemases. *FEBS Letters*, **2002**, 528, 114-8	3.8	28
194	The N-terminal replacement of an olfactory receptor for the development of a yeast-based biomimetic odor sensor. *Biotechnology and Bioengineering*, **2012**, 109, 205-12	4.9	27
193	FtsH recognizes proteins with unfolded structure and hydrolyzes the carboxyl side of hydrophobic residues. *Journal of Biochemistry*, **2000**, 127, 931-7	3.1	27
192	Structural studies on the oligomeric transition of a small heat shock protein, StHsp14.0. *Journal of Molecular Biology*, **2012**, 422, 100-8	6.5	26
191	Crystal structure of an extensively simplified variant of bovine pancreatic trypsin inhibitor in which over one-third of the residues are alanines. *Proceedings of the National Academy of Sciences of the United States of America*, **2008**, 105, 15334-9	11.5	26
190	Overexpression of prefoldin from the hyperthermophilic archaeum Pyrococcus horikoshii OT3 endowed Escherichia coli with organic solvent tolerance. *Applied Microbiology and Biotechnology*, **2008**, 79, 443-9	5.7	26
189	Cloning and functional characterization of Arabidopsis thaliana D-amino acid aminotransferase--D-aspartate behavior during germination. *FEBS Journal*, **2008**, 275, 1188-200	5.7	26
188	Mutational study on alphaGln90 of Fe-type nitrile hydratase from Rhodococcus sp. N771. *Bioscience, Biotechnology and Biochemistry*, **2006**, 70, 881-9	2.1	26
187	A novel chiral thiol reagent for automated precolumn derivatization and high-performance liquid chromatographic enantioseparation of amino acids and its application to the aspartate racemase assay. *Analytical Biochemistry*, **2003**, 315, 262-9	3.1	26
186	ATP dependent rotational motion of group II chaperonin observed by X-ray single molecule tracking. *PLoS ONE*, **2013**, 8, e64176	3.7	25
185	Expression and biochemical characterization of two small heat shock proteins from the thermoacidophilic crenarchaeon Sulfolobus tokodaii strain 7. *Protein Science*, **2004**, 13, 134-44	6.3	25
184	Kinetic and structural studies on roles of the serine ligand and a strictly conserved tyrosine residue in nitrile hydratase. *Journal of Biological Inorganic Chemistry*, **2010**, 15, 655-65	3.7	24
183	Biophysical characterization of highly active recombinant Gaussia luciferase expressed in Escherichia coli. *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2010**, 1804, 1902-7	4	23
182	Sequential action of ATP-dependent subunit conformational change and interaction between helical protrusions in the closure of the built-in lid of group II chaperonins. *Journal of Biological Chemistry*, **2008**, 283, 34773-84	5.4	23

181	Development of an integrated automation system with a magnetic bead-mediated nucleic acid purification device for genetic analysis and gene manipulation. *Biotechnology and Bioengineering*, **2004**, 86, 667-71	4.9	23
180	Two kinds of archaeal group II chaperonin subunits with different thermostability in Thermococcus strain KS-1. *Molecular Microbiology*, **2002**, 44, 761-9	4.1	23
179	Solubilization and folding of a fully active recombinant Gaussia luciferase with native disulfide bonds by using a SEP-Tag. *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2011**, 1814, 1775-8	4	22
178	Crystal structure of 1-deoxy-d-xylulose 5-phosphate reductoisomerase from the hyperthermophile Thermotoga maritima for insights into the coordination of conformational changes and an inhibitor binding. *Journal of Structural Biology*, **2010**, 170, 532-9	3.4	22
177	Functional characterization of recombinant prefoldin complexes from a hyperthermophilic archaeon, Thermococcus sp. strain KS-1. *Journal of Molecular Biology*, **2008**, 377, 972-83	6.5	22
176	Structural and molecular characterization of the prefoldin beta subunit from Thermococcus strain KS-1. *Journal of Molecular Biology*, **2008**, 383, 465-74	6.5	22
175	Molecular cloning, expression, and characterization of chaperonin-60 and chaperonin-10 from a thermophilic bacterium, Thermus thermophilus HB8. *Journal of Biochemistry*, **1995**, 118, 347-54	3.1	22
174	Improving the odorant sensitivity of olfactory receptor-expressing yeast with accessory proteins. *Analytical Biochemistry*, **2015**, 471, 1-8	3.1	21
173	Modification of the response of olfactory receptors to acetophenone by CYP1a2. *Scientific Reports*, **2017**, 7, 10167	4.9	21
172	K+ is an indispensable cofactor for GrpE stimulation of ATPase activity of DnaK x DnaJ complex from Thermus thermophilus. *FEBS Letters*, **1997**, 412, 633-6	3.8	21
171	Structure of aspartate racemase complexed with a dual substrate analogue, citric acid, and implications for the reaction mechanism. *Proteins: Structure, Function and Bioinformatics*, **2008**, 70, 1167-74	4.2	21
170	Development of a novel method for operating magnetic particles, Magtration Technology, and its use for automating nucleic acid purification. *Journal of Bioscience and Bioengineering*, **2001**, 91, 500-3	3.3	21
169	Crystal structures of the lumazine protein from Photobacterium kishitanii in complexes with the authentic chromophore, 6,7-dimethyl- 8-(1PD-ribityl) lumazine, and its analogues, riboflavin and flavin mononucleotide, at high resolution. *Journal of Bacteriology*, **2010**, 192, 127-33	3.5	20
168	Single nucleotide polymorphism genotyping of CYP2C19 using a new automated system. *Analytical Biochemistry*, **2007**, 370, 121-3	3.1	20
167	Characterization of a thermostable enzyme with phosphomannomutase/phosphoglucomutase activities from the hyperthermophilic archaeon Pyrococcus horikoshii OT3. *Journal of Biochemistry*, **2005**, 138, 159-66	3.1	20
166	The synthesis of enzyme-bound ATP by the F1-ATPase from the thermophilic bacterium PS3 in the presence of organic solvents. *Biochimica Et Biophysica Acta - Bioenergetics*, **1986**, 850, 429-435	4.6	20
165	Olfactory receptor accessory proteins play crucial roles in receptor function and gene choice. *ELife*, **2017**, 6,	8.9	20
164	Purification by dye-ligand chromatography and a crystallization study of the F1-ATPase and its major subunits, beta and alpha, from a thermophilic bacterium, PS3. *Journal of Biochemistry*, **1991**, 109, 466-71	3.1	19

163	High resolution crystal structure of dengue-3 envelope protein domain III suggests possible molecular mechanisms for serospecific antibody recognition. *Proteins: Structure, Function and Bioinformatics*, **2013**, 81, 1090-5	4.2	18
162	F0F1-ATPase genes from an archaebacterium, Methanosarcina barkeri. *Biochemical and Biophysical Research Communications*, **1997**, 241, 427-33	3.4	18
161	Modulation of redox potential and alteration in reactivity via the peroxide shunt pathway by mutation of cytochrome P450 around the proximal heme ligand. *Biochemistry*, **2008**, 47, 4834-42	3.2	18
160	Role of the N-terminal region of the crenarchaeal sHsp, StHsp14.0, in thermal-induced disassembly of the complex and molecular chaperone activity. *Biochemical and Biophysical Research Communications*, **2004**, 315, 113-8	3.4	18
159	Structural and functional characterization of homo-oligomeric complexes of □and □chaperonin subunits from the hyperthermophilic archaeum Thermococcus strain KS-1. *Journal of Molecular Biology*, **2000**, 299, 1399-1400	6.5	18
158	Thermodynamic and structural analysis of highly stabilized BPTIs by single and double mutations. *Proteins: Structure, Function and Bioinformatics*, **2009**, 77, 962-70	4.2	17
157	Contribution of the C-terminal region to the thermostability of the archaeal group II chaperonin from Thermococcus sp. strain KS-1. *Extremophiles*, **2006**, 10, 451-9	3	17
156	Properties and crystal structure of methylenetetrahydrofolate reductase from Thermus thermophilus HB8. *PLoS ONE*, **2011**, 6, e23716	3.7	17
155	Expression and characterization of the Plasmodium translocon of the exported proteins component EXP2. *Biochemical and Biophysical Research Communications*, **2017**, 482, 700-705	3.4	16
154	Three-dimensional structures of OSW-1 and its congener. *Organic Letters*, **2010**, 12, 5732-5	6.2	16
153	Thermodynamic characterization of the interaction between prefoldin and group II chaperonin. *Journal of Molecular Biology*, **2010**, 399, 628-36	6.5	16
152	Interaction of a small heat shock protein of the fission yeast, Schizosaccharomyces pombe, with a denatured protein at elevated temperature. *Journal of Biological Chemistry*, **2005**, 280, 32586-93	5.4	16
151	Recent developments in laboratory automation using magnetic particles for genome analysis. *Pharmacogenomics*, **2002**, 3, 697-708	2.6	16
150	Analysis and Control of Protein Crystallization Using Short Peptide Tags That Change Solubility without Affecting Structure, Thermal Stability, and Function. *Crystal Growth and Design*, **2015**, 15, 2703-2711	3.5	15
149	Prefoldin, a jellyfish-like molecular chaperone: functional cooperation with a group II chaperonin and beyond. *Biophysical Reviews*, **2018**, 10, 339-345	3.7	15
148	Time-Resolved Crystallography of the Reaction Intermediate of Nitrile Hydratase: Revealing a Role for the Cysteinesulfenic Acid Ligand as a Catalytic Nucleophile. *Angewandte Chemie - International Edition*, **2015**, 54, 10763-7	16.4	15
147	An improved bioluminescence-based signaling assay for odor sensing with a yeast expressing a chimeric olfactory receptor. *Biotechnology and Bioengineering*, **2012**, 109, 3143-51	4.9	15
146	Roles of conserved basic amino acid residues and activation mechanism of the hyperthermophilic aspartate racemase at high temperature. *Proteins: Structure, Function and Bioinformatics*, **2006**, 64, 502-12	4.2	15

145	Small heat shock protein of a hyperthermophilic archaeum, Thermococcus sp. strain KS-1, exists as a spherical 24 mer and its expression is highly induced under heat-stress conditions. *Journal of Bioscience and Bioengineering*, **2001**, 92, 161-166	3.3	15
144	Crystal structure of chaperonin-60 from Paracoccus denitrificans. *Journal of Molecular Biology*, **2001**, 312, 501-9	6.5	15
143	Successful PEGylation of hollow encapsulin nanoparticles from Rhodococcus erythropolis N771 without affecting their disassembly and reassembly properties. *Biomaterials Science*, **2017**, 5, 1082-1089	7.4	14
142	Computational prediction and experimental characterization of a "size switch type repacking" during the evolution of dengue envelope protein domain III (ED3). *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2014**, 1844, 585-92	4	14
141	Genome sequence determination and metagenomic characterization of a Dehalococcoides mixed culture grown on cis-1,2-dichloroethene. *Journal of Bioscience and Bioengineering*, **2015**, 120, 69-77	3.3	13
140	Detection and identification of Dehalococcoides species responsible for in situ dechlorination of trichloroethene to ethene enhanced by hydrogen-releasing compounds. *Biotechnology and Applied Biochemistry*, **2008**, 51, 1-7	2.8	13
139	Topological relation of chick thalamofugal visual projections with hyper pallium revealed by three color tracers. *Neuroscience Research*, **2005**, 52, 235-42	2.9	13
138	Functional expression of thiocyanate hydrolase is promoted by its activator protein, P15K. *FEBS Letters*, **2006**, 580, 4667-72	3.8	13
137	A novel method for direct electrochemistry of a thermoacidophilic cytochrome P450. *Electrochemistry Communications*, **2006**, 8, 1245-1249	5.1	13
136	Gene of heat shock protein of sulfur-dependent archaeal hyperthermophile Desulfurococcus. *Biochemical and Biophysical Research Communications*, **1995**, 214, 730-6	3.4	13
135	Use of Candida rugosa lipase as a biocatalyst for L-lactide ring-opening polymerization and polylactic acid production. *Biocatalysis and Agricultural Biotechnology*, **2018**, 16, 683-691	4.2	13
134	Structure and function of archaeal prefoldin, a co-chaperone of group II chaperonin. *Frontiers in Bioscience - Landmark*, **2010**, 15, 708-17	2.8	12
133	High-resolution separation of oligonucleotides and DNA sequencing reaction products by capillary electrophoresis with linear polyacrylamide and laser-induced fluorescence detection. *Journal of Separation Science*, **1994**, 6, 539-543		12
132	Anticancer saponin OSW-1 is a novel class of selective Golgi stress inducer. *Bioorganic and Medicinal Chemistry Letters*, **2019**, 29, 1732-1736	2.9	11
131	Catalytic Mechanism of Nitrile Hydratase Subsequent to Cyclic Intermediate Formation: A QM/MM Study. *Journal of Physical Chemistry B*, **2016**, 120, 3259-66	3.4	11
130	Identification of the rate-limiting step of the peroxygenase reactions catalyzed by the thermophilic cytochrome P450 from Sulfolobus tokodaii strain 7. *FEBS Journal*, **2014**, 281, 1409-16	5.7	11
129	Structure-based mutational study of an archaeal DNA ligase towards improvement of ligation activity. *ChemBioChem*, **2012**, 13, 2575-82	3.8	11
128	The electrochemical properties of thermophilic cytochrome P450 CYP119A2 at extremely high temperatures in poly(ethylene oxide). *Electrochemistry Communications*, **2007**, 9, 361-364	5.1	11

127	MagSNiPer: a new single nucleotide polymorphism typing method based on single base extension, magnetic separation, and chemiluminescence. *Analytical Biochemistry*, **2005**, 341, 77-82	3.1	11
126	Characterization of homo-oligomeric complexes of alpha and beta chaperonin subunits from the acidothermophilic archaeon, Sulfolobus sp. strain 7. *Biochemical and Biophysical Research Communications*, **1998**, 242, 640-7	3.4	11
125	Structural instability and divergence from conserved residues underlie intracellular retention of mammalian odorant receptors. *Proceedings of the National Academy of Sciences of the United States of America*, **2020**, 117, 2957-2967	11.5	10
124	Dissection of the ATP-dependent conformational change cycle of a group II chaperonin. *Journal of Molecular Biology*, **2014**, 426, 447-59	6.5	10
123	Crystal structures of halohydrin hydrogen-halide-lyases from Corynebacterium sp. N-1074. *Proteins: Structure, Function and Bioinformatics*, **2015**, 83, 2230-9	4.2	10
122	Purification and molecular cloning of the group II chaperonin from the acidothermophilic archaeon, Sulfolobus sp. strain 7. *Biochemical and Biophysical Research Communications*, **1997**, 236, 727-32	3.4	10
121	Solubilization of heme proteins in low polar solvents by chemical modification on a protein surface. *Polymers for Advanced Technologies*, **2008**, 19, 1430-1435	3.2	10
120	Inter-ring communication is dispensable in the reaction cycle of group II chaperonins. *Journal of Molecular Biology*, **2014**, 426, 2667-78	6.5	9
119	Biochemical characterization and cooperation with co-chaperones of heat shock protein 90 from Schizosaccharomyces pombe. *Journal of Bioscience and Bioengineering*, **2013**, 116, 444-8	3.3	9
118	Isolation and genomic characterization of a Dehalococcoides strain suggests genomic rearrangement during culture. *Scientific Reports*, **2017**, 7, 2230	4.9	9
117	Specificity of MicroRNA Detection on a Power-free Microfluidic Chip with Laminar Flow-assisted Dendritic Amplification. *Analytical Sciences*, **2017**, 33, 171-177	1.7	9
116	Characterization of a sHsp of Schizosaccharomyces pombe, SpHsp15.8, and the implication of its functional mechanism by comparison with another sHsp, SpHsp16.0. *Proteins: Structure, Function and Bioinformatics*, **2009**, 74, 6-17	4.2	9
115	The stabilizing residues and the functional domains in the hyperthermophilic V-ATPase of Desulfurococcus. *Biochemical and Biophysical Research Communications*, **1997**, 234, 341-5	3.4	9
114	Novel catalytic activity of nitrile hydratase from Rhodococcus sp. N771. *Journal of Bioscience and Bioengineering*, **2008**, 106, 174-9	3.3	9
113	Contribution of the C-Terminal Region of a Group II Chaperonin to its Interaction with Prefoldin and Substrate Transfer. *Journal of Molecular Biology*, **2016**, 428, 2405-2417	6.5	9
112	Size-selective recognition of gold nanoparticles by a molecular chaperone. *Chemical Physics Letters*, **2010**, 501, 108-112	2.5	8
111	Rapid construction of a transcription map for a cosmid contig of Arabidopsis thaliana genome using a novel cDNA selection method. *Plant Journal*, **1997**, 12, 481-7	6.9	8
110	Protein refolding system using holo-chaperonin from the thermophilic bacterium Thermus thermophilus. *Journal of Bioscience and Bioengineering*, **1998**, 85, 564-570		8

109	Dynamics of group II chaperonin and prefoldin probed by 13C NMR spectroscopy. *Proteins: Structure, Function and Bioinformatics*, **2008**, 70, 1257-63	4.2	8
108	Development of an automation system for single nucleotide polymorphisms genotyping using bio-strand, a new three-dimensional microarray. *Journal of Bioscience and Bioengineering*, **2005**, 99, 120-4	3.3	8
107	Stability of Thermophilic Cytochrome P450 Modified with Poly(ethylene oxide) in Ionic Liquid. *Chemistry Letters*, **2006**, 35, 798-799	1.7	8
106	Inversion of the anatomical lateralization of chick thalamofugal visual pathway by light experience. *Neuroscience Letters*, **2002**, 318, 113-6	3.3	8
105	Single site catalysis of the F1-ATPase from Saccharomyces cerevisiae and the effect of inorganic phosphate on it. *Journal of Biochemistry*, **1987**, 102, 273-9	3.1	8
104	Asymmetry in the function and dynamics of the cytosolic group II chaperonin CCT/TRiC. *PLoS ONE*, **2017**, 12, e0176054	3.7	8
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99	Goniothalamin enhances the ATPase activity of the molecular chaperone Hsp90 but inhibits its chaperone activity. *Journal of Biochemistry*, **2015**, 157, 161-8	3.1	7
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1 Protein-Folding Systems209-223