Masaru Tanokura

List of Publications by Citations

Source: https://exaly.com/author-pdf/34596/masaru-tanokura-publications-by-citations.pdf

Version: 2024-04-10

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

13,756 496 102 54 h-index g-index citations papers 6.1 516 15,547 5.1 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
496	Mitochondrial DNA mutations, oxidative stress, and apoptosis in mammalian aging. <i>Science</i> , 2005 , 309, 481-4	33.3	1573
495	Sirt3 mediates reduction of oxidative damage and prevention of age-related hearing loss under caloric restriction. <i>Cell</i> , 2010 , 143, 802-12	56.2	860
494	Molecular basis of the core regulatory network in ABA responses: sensing, signaling and transport. <i>Plant and Cell Physiology</i> , 2010 , 51, 1821-39	4.9	612
493	Structural basis of abscisic acid signalling. <i>Nature</i> , 2009 , 462, 609-14	50.4	406
492	Arabidopsis DREB2A-interacting proteins function as RING E3 ligases and negatively regulate plant drought stress-responsive gene expression. <i>Plant Cell</i> , 2008 , 20, 1693-707	11.6	361
491	Synergistic activation of the Arabidopsis NADPH oxidase AtrbohD by Ca2+ and phosphorylation. Journal of Biological Chemistry, 2008 , 283, 8885-92	5.4	344
490	Molecular mechanism of strigolactone perception by DWARF14. <i>Nature Communications</i> , 2013 , 4, 2613	17.4	245
489	Age-related hearing loss in C57BL/6J mice is mediated by Bak-dependent mitochondrial apoptosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 19432-7	11.5	238
488	Mitochondrial DNA mutations induce mitochondrial dysfunction, apoptosis and sarcopenia in skeletal muscle of mitochondrial DNA mutator mice. <i>PLoS ONE</i> , 2010 , 5, e11468	3.7	196
487	A novel Call+-activated, thermostabilized polyesterase capable of hydrolyzing polyethylene terephthalate from Saccharomonospora viridis AHK190. <i>Applied Microbiology and Biotechnology</i> , 2014 , 98, 10053-64	5.7	155
486	Biochemical characterization of NfsA, the Escherichia coli major nitroreductase exhibiting a high amino acid sequence homology to Frp, a Vibrio harveyi flavin oxidoreductase. <i>Journal of Bacteriology</i> , 1996 , 178, 4508-14	3.5	154
485	Single-molecule paleoenzymology probes the chemistry of resurrected enzymes. <i>Nature Structural and Molecular Biology</i> , 2011 , 18, 592-6	17.6	149
484	Crystal structure of the terminal oxygenase component of biphenyl dioxygenase derived from Rhodococcus sp. strain RHA1. <i>Journal of Molecular Biology</i> , 2004 , 342, 1041-52	6.5	127
483	Structure and function of abscisic acid receptors. <i>Trends in Plant Science</i> , 2013 , 18, 259-66	13.1	126
482	Gene cloning, purification, and characterization of NfsB, a minor oxygen-insensitive nitroreductase from Escherichia coli, similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri. <i>Journal of Biochemistry</i> , 1996 , 120, 736-44	3.1	119
481	Experimental evidence for the thermophilicity of ancestral life. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 11067-72	11.5	112
480	Optimisation of hydrolysis conditions for the production of the angiotensin-I converting enzyme (ACE) inhibitory peptides from whey protein using response surface methodology. <i>Food Chemistry</i> , 2009 , 114, 328-333	8.5	111

479	Roasting process of coffee beans as studied by nuclear magnetic resonance: time course of changes in composition. <i>Journal of Agricultural and Food Chemistry</i> , 2012 , 60, 1005-12	5.7	103	
478	Caloric restriction suppresses apoptotic cell death in the mammalian cochlea and leads to prevention of presbycusis. <i>Neurobiology of Aging</i> , 2007 , 28, 1613-22	5.6	103	
477	1H-NMR study on the tautomerism of the imidazole ring of histidine residues. I. Microscopic pK values and molar ratios of tautomers in histidine-containing peptides. <i>BBA - Proteins and Proteomics</i> , 1983 , 742, 576-85		103	
476	Role of mitochondrial dysfunction and mitochondrial DNA mutations in age-related hearing loss. <i>Hearing Research</i> , 2007 , 226, 185-93	3.9	101	
475	Structural basis for Ca2+-regulated muscle relaxation at interaction sites of troponin with actin and tropomyosin. <i>Journal of Molecular Biology</i> , 2005 , 352, 178-201	6.5	98	
474	Three-dimensional solution structure of oryzacystatin-I, a cysteine proteinase inhibitor of the rice, Oryza sativa L. japonica. <i>Biochemistry</i> , 2000 , 39, 14753-60	3.2	97	
473	1H nuclear magnetic resonance studies of histidine-containing di- and tripeptides. Estimation of the effects of charged groups on the pKa value of the imidazole ring. <i>Biopolymers</i> , 1976 , 15, 393-401	2.2	96	
472	Antihypertensive peptides from skimmed milk hydrolysate digested by cell-free extract of Lactobacillus helveticus JCM1004. <i>Food Chemistry</i> , 2005 , 91, 123-129	8.5	93	
471	Structure of Alcaligenes faecalis nitrite reductase and a copper site mutant, M150E, that contains zinc. <i>Biochemistry</i> , 1995 , 34, 12107-17	3.2	92	
470	(13)C NMR-based metabolomics for the classification of green coffee beans according to variety and origin. <i>Journal of Agricultural and Food Chemistry</i> , 2012 , 60, 10118-25	5.7	89	
469	Calcium binding to calmodulin: effects of ionic strength, Mg2+, pH and temperature. <i>Journal of Biochemistry</i> , 1984 , 95, 19-28	3.1	86	
468	Nondestructive quantification of organic compounds in whole milk without pretreatment by two-dimensional NMR spectroscopy. <i>Journal of Agricultural and Food Chemistry</i> , 2007 , 55, 4307-11	5.7	84	
467	Chymotryptic subfragments of troponin T from rabbit skeletal muscle. Interaction with tropomyosin, troponin I and troponin C. <i>Journal of Biochemistry</i> , 1983 , 93, 331-7	3.1	83	
466	Infrared studies of interaction between metal ions and Ca(2+)-binding proteins. Marker bands for identifying the types of coordination of the side-chain COO- groups to metal ions in pike parvalbumin (pI = 4.10). <i>FEBS Letters</i> , 1994 , 349, 84-8	3.8	81	
465	Isolation of an antihypertensive peptide from alcalase digest of Spirulina platensis. <i>Journal of Agricultural and Food Chemistry</i> , 2010 , 58, 7166-71	5.7	80	
464	Coordination to divalent cations by calcium-binding proteins studied by FTIR spectroscopy. Biochimica Et Biophysica Acta - Biomembranes, 2013 , 1828, 2319-27	3.8	77	
463	Structure and site-directed mutagenesis of a flavoprotein from Escherichia coli that reduces nitrocompounds: alteration of pyridine nucleotide binding by a single amino acid substitution. <i>Journal of Biological Chemistry</i> , 2001 , 276, 2816-23	5.4	76	
462	1.8 A crystal structure of the major NAD(P)H:FMN oxidoreductase of a bioluminescent bacterium, Vibrio fischeri: overall structure, cofactor and substrate-analog binding, and comparison with related flavoproteins. <i>Journal of Molecular Biology</i> , 1998 , 280, 259-73	6.5	74	

461	Structural basis for the Ca(2+)-enhanced thermostability and activity of PET-degrading cutinase-like enzyme from Saccharomonospora viridis AHK190. <i>Applied Microbiology and Biotechnology</i> , 2015 , 99, 429	9 7 :⁄307	70
460	The role of mtDNA mutations in the pathogenesis of age-related hearing loss in mice carrying a mutator DNA polymerase gamma. <i>Neurobiology of Aging</i> , 2008 , 29, 1080-92	5.6	70
459	The solution structure of molt-inhibiting hormone from the Kuruma prawn Marsupenaeus japonicus. <i>Journal of Biological Chemistry</i> , 2003 , 278, 9620-3	5.4	70
458	Comparative study of catechin compositions in five Japanese persimmons (). <i>Food Chemistry</i> , 2005 , 93, 149-152	8.5	70
457	Discovery of a novel restriction endonuclease by genome comparison and application of a wheat-germ-based cell-free translation assay: PabI (5MGTA/C) from the hyperthermophilic archaeon Pyrococcus abyssi. <i>Nucleic Acids Research</i> , 2005 , 33, e112	20.1	67
456	Homodimeric structure and double-stranded RNA cleavage activity of the C-terminal RNase III domain of human dicer. <i>Journal of Molecular Biology</i> , 2007 , 374, 106-20	6.5	66
455	Complex mixture analysis of organic compounds in green coffee bean extract by two-dimensional NMR spectroscopy. <i>Magnetic Resonance in Chemistry</i> , 2010 , 48, 857-65	2.1	65
454	Drastic Ca2+ sensitization of myofilament associated with a small structural change in troponin I in inherited restrictive cardiomyopathy. <i>Biochemical and Biophysical Research Communications</i> , 2005 , 338, 1519-26	3.4	65
453	Comprehensive NMR analysis of compositional changes of black garlic during thermal processing. Journal of Agricultural and Food Chemistry, 2015 , 63, 683-91	5.7	64
452	Autism spectrum disorder is related to endoplasmic reticulum stress induced by mutations in the synaptic cell adhesion molecule, CADM1. <i>Cell Death and Disease</i> , 2010 , 1, e47	9.8	64
451	Siamycin attenuates fsr quorum sensing mediated by a gelatinase biosynthesis-activating pheromone in Enterococcus faecalis. <i>Journal of Bacteriology</i> , 2007 , 189, 1358-65	3.5	64
45°	Aspartate kinase-independent lysine synthesis in an extremely thermophilic bacterium, Thermus thermophilus: lysine is synthesized via alpha-aminoadipic acid not via diaminopimelic acid. <i>Journal of Bacteriology</i> , 1999 , 181, 1713-8	3.5	64
449	Three-dimensional structure of AzoR from Escherichia coli. An oxidereductase conserved in microorganisms. <i>Journal of Biological Chemistry</i> , 2006 , 281, 20567-76	5.4	63
448	Nondestructive observation of bovine milk by NMR spectroscopy: analysis of existing States of compounds and detection of new compounds. <i>Journal of Agricultural and Food Chemistry</i> , 2004 , 52, 496	9 ⁵ 7 4	61
447	Calaxin drives sperm chemotaxis by Call+-mediated direct modulation of a dynein motor. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20497-502	11.5	60
446	Isolation and tyrosinase inhibitory effects of polyphenols from the leaves of persimmon, Diospyros kaki. <i>Journal of Agricultural and Food Chemistry</i> , 2011 , 59, 6011-7	5.7	59
445	Infrared spectroscopic study of the metal-coordination structures of calcium-binding proteins. <i>Biochemical and Biophysical Research Communications</i> , 2008 , 369, 225-39	3.4	57
444	Three-dimensional solution structure of an archaeal FKBP with a dual function of peptidyl prolyl cis-trans isomerase and chaperone-like activities. <i>Journal of Molecular Biology</i> , 2003 , 328, 1149-60	6.5	56

(2013-2001)

443	The three-dimensional structure of septum site-determining protein MinD from Pyrococcus horikoshii OT3 in complex with Mg-ADP. <i>Structure</i> , 2001 , 9, 817-26	5.2	56
442	Purification, characterization, and molecular gene cloning of an antifungal protein from Ginkgo biloba seeds. <i>Biological Chemistry</i> , 2007 , 388, 273-80	4.5	53
441	Common mechanism of ligand recognition by group II/III WW domains: redefining their functional classification. <i>Journal of Biological Chemistry</i> , 2004 , 279, 31833-41	5.4	53
440	Expansion of substrate specificity and catalytic mechanism of azoreductase by X-ray crystallography and site-directed mutagenesis. <i>Journal of Biological Chemistry</i> , 2008 , 283, 13889-96	5.4	52
439	Biological actions of green tea catechins on cardiac troponin C. <i>British Journal of Pharmacology</i> , 2010 , 161, 1034-43	8.6	51
438	Metabolic discrimination of mango juice from various cultivars by band-selective NMR spectroscopy. <i>Journal of Agricultural and Food Chemistry</i> , 2012 , 60, 1158-66	5.7	50
437	Studies on protein-protein interaction between copper-containing nitrite reductase and pseudoazurin from Alcaligenes faecalis S-6. <i>Journal of Biological Chemistry</i> , 1996 , 271, 13680-3	5.4	49
436	Heat capacity and entropy changes of calmodulin induced by calcium binding. <i>Journal of Biochemistry</i> , 1984 , 95, 643-9	3.1	48
435	A pilot study of NMR-based sensory prediction of roasted coffee bean extracts. <i>Food Chemistry</i> , 2014 , 152, 363-9	8.5	47
	Novel protein fold discovered in the PabI family of restriction enzymes. <i>Nucleic Acids Research</i> ,		
434	2007 , 35, 1908-18	20.1	45
434		3.5	44
	2007, 35, 1908-18 Conversion of NfsB, a minor Escherichia coli nitroreductase, to a flavin reductase similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri, by a single amino		
433	2007, 35, 1908-18 Conversion of NfsB, a minor Escherichia coli nitroreductase, to a flavin reductase similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri, by a single amino acid substitution. <i>Journal of Bacteriology</i> , 1996, 178, 4731-3 1H-NMR study on the tautomerism of the imidazole ring of histidine residues. II. Microenvironments of histidine-12 and histidine-119 of bovine pancreatic ribonuclease A. <i>BBA</i> -		44
433	Conversion of NfsB, a minor Escherichia coli nitroreductase, to a flavin reductase similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri, by a single amino acid substitution. <i>Journal of Bacteriology</i> , 1996 , 178, 4731-3 1H-NMR study on the tautomerism of the imidazole ring of histidine residues. II. Microenvironments of histidine-12 and histidine-119 of bovine pancreatic ribonuclease A. <i>BBA</i> - <i>Proteins and Proteomics</i> , 1983 , 742, 586-96 A secreted protein with plant-specific cysteine-rich motif functions as a mannose-binding lectin	3.5	44
433 432 431	Conversion of NfsB, a minor Escherichia coli nitroreductase, to a flavin reductase similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri, by a single amino acid substitution. <i>Journal of Bacteriology</i> , 1996 , 178, 4731-3 1H-NMR study on the tautomerism of the imidazole ring of histidine residues. II. Microenvironments of histidine-12 and histidine-119 of bovine pancreatic ribonuclease A. <i>BBA - Proteins and Proteomics</i> , 1983 , 742, 586-96 A secreted protein with plant-specific cysteine-rich motif functions as a mannose-binding lectin that exhibits antifungal activity. <i>Plant Physiology</i> , 2014 , 166, 766-78 Structural analysis of HTL and D14 proteins reveals the basis for ligand selectivity in Striga. <i>Nature</i>	3.5	44 44 43
433 432 431 430	 2007, 35, 1908-18 Conversion of NfsB, a minor Escherichia coli nitroreductase, to a flavin reductase similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri, by a single amino acid substitution. <i>Journal of Bacteriology</i>, 1996, 178, 4731-3 1H-NMR study on the tautomerism of the imidazole ring of histidine residues. II. Microenvironments of histidine-12 and histidine-119 of bovine pancreatic ribonuclease A. <i>BBA-Proteins and Proteomics</i>, 1983, 742, 586-96 A secreted protein with plant-specific cysteine-rich motif functions as a mannose-binding lectin that exhibits antifungal activity. <i>Plant Physiology</i>, 2014, 166, 766-78 Structural analysis of HTL and D14 proteins reveals the basis for ligand selectivity in Striga. <i>Nature Communications</i>, 2018, 9, 3947 Two-dimensional 1H-13C nuclear magnetic resonance (NMR)-based comprehensive analysis of 	3.5 6.6	44 44 43 43
433 432 431 430 429	Conversion of NfsB, a minor Escherichia coli nitroreductase, to a flavin reductase similar in biochemical properties to FRase I, the major flavin reductase in Vibrio fischeri, by a single amino acid substitution. <i>Journal of Bacteriology</i> , 1996, 178, 4731-3 1H-NMR study on the tautomerism of the imidazole ring of histidine residues. II. Microenvironments of histidine-12 and histidine-119 of bovine pancreatic ribonuclease A. <i>BBA-Proteins and Proteomics</i> , 1983, 742, 586-96 A secreted protein with plant-specific cysteine-rich motif functions as a mannose-binding lectin that exhibits antifungal activity. <i>Plant Physiology</i> , 2014, 166, 766-78 Structural analysis of HTL and D14 proteins reveals the basis for ligand selectivity in Striga. <i>Nature Communications</i> , 2018, 9, 3947 Two-dimensional 1H-13C nuclear magnetic resonance (NMR)-based comprehensive analysis of roasted coffee bean extract. <i>Journal of Agricultural and Food Chemistry</i> , 2011, 59, 9065-73 NMR-based metabolomics for simultaneously evaluating multiple determinants of primary beef	3.5 6.6 17.4 5.7	4444434342

425	Multiple parallel-pathway folding of proline-free Staphylococcal nuclease. <i>Journal of Molecular Biology</i> , 2003 , 332, 1143-53	6.5	40
424	Addition of exogenous NAD+ prevents mefloquine-induced neuroaxonal and hair cell degeneration through reduction of caspase-3-mediated apoptosis in cochlear organotypic cultures. <i>PLoS ONE</i> , 2013 , 8, e79817	3.7	39
423	Structural characterization of the N-linked carbohydrate chains of the zona pellucida glycoproteins from bovine ovarian and fertilized eggs. <i>FEBS Journal</i> , 1996 , 240, 448-53		39
422	Structural basis of unique ligand specificity of KAI2-like protein from parasitic weed Striga hermonthica. <i>Scientific Reports</i> , 2016 , 6, 31386	4.9	38
421	Structural basis for controlling the enzymatic properties of polymannuronate preferred alginate lyase FlAlyA from the PL-7 family. <i>Chemical Communications</i> , 2018 , 54, 555-558	5.8	37
420	Purification and characterization of NfrA1, a Bacillus subtilis nitro/flavin reductase capable of interacting with the bacterial luciferase. <i>Bioscience, Biotechnology and Biochemistry</i> , 1998 , 62, 1978-87	2.1	37
419	A new target region for changing the substrate specificity of amine transaminases. <i>Scientific Reports</i> , 2015 , 5, 10753	4.9	36
418	Effects of caloric restriction on age-related hearing loss in rodents and rhesus monkeys. <i>Current Aging Science</i> , 2010 , 3, 20-5	2.2	36
417	Purification and gene cloning of Fundulus heteroclitus hatching enzyme. A hatching enzyme system composed of high choriolytic enzyme and low choriolytic enzyme is conserved between two different teleosts, Fundulus heteroclitus and medaka Oryzias latipes. <i>FEBS Journal</i> , 2005 , 272, 4315-26	5.7	36
416	Conversion of NfsA, the major Escherichia coli nitroreductase, to a flavin reductase with an activity similar to that of Frp, a flavin reductase in Vibrio harveyi, by a single amino acid substitution. <i>Journal of Bacteriology</i> , 1998 , 180, 422-5	3.5	36
415	Crystal structure and desulfurization mechanism of 2Mhydroxybiphenyl-2-sulfinic acid desulfinase. Journal of Biological Chemistry, 2006 , 281, 32534-9	5.4	35
414	Coordination structures of Ca2+ and Mg2+ in Akazara scallop troponin C in solution. FTIR spectroscopy of side-chain COO- groups. <i>FEBS Journal</i> , 2001 , 268, 6284-90		35
413	One-week antihypertensive effect of Ile-Gln-Pro in spontaneously hypertensive rats. <i>Journal of Agricultural and Food Chemistry</i> , 2011 , 59, 559-63	5.7	34
412	Proteinase inhibitor from ginkgo seeds is a member of the plant nonspecific lipid transfer protein gene family. <i>Plant Physiology</i> , 2008 , 146, 1909-19	6.6	34
411	Direct demonstration of the cross-bridge recovery stroke in muscle thick filaments in aqueous solution by using the hydration chamber. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 17396-401	11.5	34
410	pH-dependent unfolding of aspergillopepsin II studied by small-angle X-ray scattering. <i>Biochemistry</i> , 2000 , 39, 1364-72	3.2	34
409	Solution structure of bromelain inhibitor IV from pineapple stem: structural similarity with Bowman-Birk trypsin/chymotrypsin inhibitor from soybean. <i>Biochemistry</i> , 1996 , 35, 5379-84	3.2	34
408	Crystal structure of Ehexachlorocyclohexane Dehydrochlorinase LinA from Sphingobium japonicum UT26. <i>Journal of Molecular Biology</i> , 2010 , 403, 260-9	6.5	33

407	A 26K fragment of troponin T from rabbit skeletal muscle. <i>Journal of Biochemistry</i> , 1984 , 95, 1337-42	3.1	33
406	Crystal structure of zebrafish hatching enzyme 1 from the zebrafish Danio rerio. <i>Journal of Molecular Biology</i> , 2010 , 402, 865-78	6.5	32
405	A calorimetric study of Ca2+- and Mg2+-binding by calmodulin. <i>Journal of Biochemistry</i> , 1983 , 94, 607-9	3.1	32
404	Structure-activity relationship of gelatinase biosynthesis-activating pheromone of Enterococcus faecalis. <i>Journal of Bacteriology</i> , 2009 , 191, 641-50	3.5	31
403	Genes encoding mitochondrial respiratory chain components are profoundly down-regulated with aging in the cochlea of DBA/2J mice. <i>Brain Research</i> , 2007 , 1182, 26-33	3.7	31
402	Crystal structures of the short-chain flavin reductase HpaC from Sulfolobus tokodaii strain 7 in its three states: NAD(P)(+)(-)free, NAD(+)(-)bound, and NADP(+)(-)bound. <i>Biochemistry</i> , 2006 , 45, 5103-10	3.2	31
401	Effects of Long-Term Exercise on Age-Related Hearing Loss in Mice. <i>Journal of Neuroscience</i> , 2016 , 36, 11308-11319	6.6	30
400	Re-Evaluation of the PBAN Receptor Molecule: Characterization of PBANR Variants Expressed in the Pheromone Glands of Moths. <i>Frontiers in Endocrinology</i> , 2012 , 3, 6	5.7	30
399	A comparative study of the binding effects of Mg2+, Ca2+, Sr2+, and Cd2+ on calmodulin by fourier-transform infrared spectroscopy. <i>Biospectroscopy</i> , 1995 , 1, 47-54		30
398	Crystal structure of ginkbilobin-2 with homology to the extracellular domain of plant cysteine-rich receptor-like kinases. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009 , 77, 247-51	4.2	29
397	Improvement of 2Mhydroxybiphenyl-2-sulfinate desulfinase, an enzyme involved in the dibenzothiophene desulfurization pathway, from Rhodococcus erythropolis KA2-5-1 by site-directed mutagenesis. <i>Bioscience, Biotechnology and Biochemistry</i> , 2007 , 71, 2815-21	2.1	29
396	Recombinant porcine zona pellucida glycoproteins expressed in Sf9 cells bind to bovine sperm but not to porcine sperm. <i>Journal of Biological Chemistry</i> , 2005 , 280, 20189-96	5.4	29
395	1H nuclear magnetic resonance titration curves and microenvironments of aromatic residues in bovine pancreatic ribonuclease A. <i>Journal of Biochemistry</i> , 1983 , 94, 51-62	3.1	29
394	Optimization of the ultrafiltration-assisted extraction of Chinese yam polysaccharide using response surface methodology and its biological activity. <i>International Journal of Biological Macromolecules</i> , 2019 , 121, 1186-1193	7.9	29
393	Development of an Azoreductase-based Reporter System with Synthetic Fluorogenic Substrates. <i>ACS Chemical Biology</i> , 2017 , 12, 558-563	4.9	28
392	Redesign of a novel D-allulose 3-epimerase from Staphylococcus aureus for thermostability and efficient biocatalytic production of D-allulose. <i>Microbial Cell Factories</i> , 2019 , 18, 59	6.4	28
391	A panel set for epitope analysis of myeloperoxidase (MPO)-specific antineutrophil cytoplasmic antibody MPO-ANCA using recombinant hexamer histidine-tagged MPO deletion mutants. <i>Journal of Clinical Immunology</i> , 1998 , 18, 142-52	5.7	28
390	The flavoenzyme azobenzene reductase AzoR from Escherichia coli binds roseoflavin mononucleotide (RoFMN) with high affinity and is less active in its RoFMN form. <i>Biochemistry</i> , 2013 , 52, 4288-95	3.2	27

389	Functional heterologous expression and characterization of mannuronan C5-epimerase from the brown alga Saccharina japonica. <i>Algal Research</i> , 2016 , 16, 282-291	5	26
388	Phosphorylation of proteins by dry-heating in the presence of pyrophosphate and some characteristics of introduced phosphate groups. <i>Food Chemistry</i> , 2009 , 114, 1036-1041	8.5	26
387	Multiplication of a restriction-modification gene complex. <i>Molecular Microbiology</i> , 2003 , 48, 417-27	4.1	26
386	Enzymes useful for chiral compound synthesis: structural biology, directed evolution, and protein engineering for industrial use. <i>Applied Microbiology and Biotechnology</i> , 2016 , 100, 5747-57	5.7	26
385	Improvement in Quality of Protein Crystals Grown in a High Magnetic Field Gradient. <i>Crystal Growth and Design</i> , 2012 , 12, 1141-1150	3.5	25
384	Steady-state properties of calcium binding to parvalbumins from bullfrog skeletal muscle: effects of Mg2+, pH, ionic strength, and temperature. <i>Journal of Biochemistry</i> , 1986 , 99, 73-80	3.1	25
383	Chymotryptic subfragments of troponin T from rabbit skeletal muscle. I. Determination of the primary structure. <i>Journal of Biochemistry</i> , 1982 , 91, 1257-65	3.1	25
382	NMR STUDY ON THE PROTONATION OF IMIDAZOLE RING OFN-ACETYL-L-HISTIDINE METHYLAMIDE, A MODEL FOR HISTIDINE RESIDUES EXPOSED TO AQUEOUS SOLVENT. <i>Chemistry Letters</i> , 1978 , 7, 739-742	1.7	25
381	Methyl phenlactonoates are efficient strigolactone analogs with simple structure. <i>Journal of Experimental Botany</i> , 2018 , 69, 2319-2331	7	25
380	Biochemical characterization and biocatalytic application of a novel d-tagatose 3-epimerase from sp <i>RSC Advances</i> , 2019 , 9, 2919-2927	3.7	24
379	Comparative studies on the functional roles of N- and C-terminal regions of molluskan and vertebrate troponin-I. <i>FEBS Journal</i> , 2005 , 272, 4475-86	5.7	24
378	Kinetic studies of calcium binding to parvalbumins from bullfrog skeletal muscle. <i>Journal of Biochemistry</i> , 1986 , 99, 81-9	3.1	24
377	A sequence-specific DNA glycosylase mediates restriction-modification in Pyrococcus abyssi. <i>Nature Communications</i> , 2014 , 5, 3178	17.4	23
376	Cooperative DNA-binding and sequence-recognition mechanism of aristaless and clawless. <i>EMBO Journal</i> , 2010 , 29, 1613-23	13	23
375	Structure of the histone chaperone CIA/ASF1-double bromodomain complex linking histone modifications and site-specific histone eviction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 8153-8	11.5	23
374	Isolation and Caenorhabditis elegans lifespan assay of flavonoids from onion. <i>Journal of Agricultural and Food Chemistry</i> , 2011 , 59, 5927-34	5.7	23
373	Studies of the structure of multiferric ion-bound lactoferrin: A new antianemic edible material. <i>International Dairy Journal</i> , 2008 , 18, 1051-1056	3.5	23
372	Site-directed mutagenesis of azurin from Pseudomonas aeruginosa enhances the formation of an electron-transfer complex with a copper-containing nitrite reductase from Alcaligenes faecalis S-6.	3.8	23

(2006-2019)

371	Triazole Ureas Covalently Bind to Strigolactone Receptor and Antagonize Strigolactone Responses. <i>Molecular Plant</i> , 2019 , 12, 44-58	14.4	23
370	Crystal structures of apo-DszC and FMN-bound DszC from Rhodococcus erythropolis D-1. <i>FEBS Journal</i> , 2015 , 282, 3126-35	5.7	22
369	Structural basis for cyclization specificity of two Azotobacter type III polyketide synthases: a single amino acid substitution reverses their cyclization specificity. <i>Journal of Biological Chemistry</i> , 2013 , 288, 34146-34157	5.4	22
368	Separation and characterization of the colored material from sugarcane molasses. <i>Chemosphere</i> , 2008 , 71, 1730-7	8.4	22
367	Crystal structure of the PIN domain of human telomerase-associated protein EST1A. <i>Proteins:</i> Structure, Function and Bioinformatics, 2007 , 68, 980-9	4.2	22
366	A calorimetric study of Ca2+ binding by the parvalbumin of the toad (Bufo): distinguishable binding sites in the molecule. <i>FEBS Letters</i> , 1986 , 209, 77-82	3.8	22
365	The effect of acyclic retinoid on the metabolomic profiles of hepatocytes and hepatocellular carcinoma cells. <i>PLoS ONE</i> , 2013 , 8, e82860	3.7	22
364	Structure and Polymannuronate Specificity of a Eukaryotic Member of Polysaccharide Lyase Family 14. <i>Journal of Biological Chemistry</i> , 2017 , 292, 2182-2190	5.4	21
363	Loss of IDH2 Accelerates Age-related Hearing Loss in Male Mice. Scientific Reports, 2018, 8, 5039	4.9	21
362	NMR-based metabolic profiling of rice wines by F(2)-selective total correlation spectra. <i>Journal of Agricultural and Food Chemistry</i> , 2012 , 60, 4818-25	5.7	21
361	The structure of brazzein, a sweet-tasting protein from the wild African plant Pentadiplandra brazzeana. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013 , 69, 642-7		21
360	GSTA4 mediates reduction of cisplatin ototoxicity in female mice. <i>Nature Communications</i> , 2019 , 10, 415	5 0 7.4	20
359	Effects of different modification methods on the physicochemical and rheological properties of Chinese yam (Dioscorea opposita Thunb.) starch. <i>LWT - Food Science and Technology</i> , 2019 , 116, 108513	5.4	20
358	Control of the localization and function of a miRNA silencing component TNRC6A by Argonaute protein. <i>Nucleic Acids Research</i> , 2015 , 43, 9856-73	20.1	20
357	Structural and biochemical elucidation of mechanism for decarboxylative condensation of beta-keto acid by curcumin synthase. <i>Journal of Biological Chemistry</i> , 2011 , 286, 6659-68	5.4	20
356	Substrate recognition mechanism and substrate-dependent conformational changes of an ROK family glucokinase from Streptomyces griseus. <i>Journal of Bacteriology</i> , 2012 , 194, 607-16	3.5	20
355	Recombinant bovine zona pellucida glycoproteins ZP3 and ZP4 coexpressed in Sf9 cells form a sperm-binding active hetero-complex. <i>FEBS Journal</i> , 2007 , 274, 5390-405	5.7	20
354	Fourier transform infrared spectroscopic analysis of the intact zona pellucida of the mammalian egg: changes in the secondary structure of bovine zona pellucida proteins during fertilization. <i>Experimental Biology and Medicine</i> , 2006 , 231, 166-71	3.7	20

353	Phosphorus nuclear magnetic resonance studies on the effect of duration of contraction in bull-frog skeletal muscles. <i>Journal of Physiology</i> , 1988 , 407, 243-61	3.9	20
352	A calorimetric study of Ca2+ binding to two major isotypes of bullfrog parvalbumin. <i>FEBS Letters</i> , 1985 , 185, 165-9	3.8	20
351	An engineered old yellow enzyme that enables efficient synthesis of (4R,6R)-Actinol in a one-pot reduction system. <i>ChemBioChem</i> , 2015 , 16, 440-5	3.8	19
350	Crystal structure and site-directed mutagenesis analyses of haloalkane dehalogenase LinB from Sphingobium sp. strain MI1205. <i>Journal of Bacteriology</i> , 2013 , 195, 2642-51	3.5	19
349	Crystal structure of a novel N-substituted L-amino acid dioxygenase from Burkholderia ambifaria AMMD. <i>PLoS ONE</i> , 2013 , 8, e63996	3.7	19
348	siRNA-mediated knockdown of aryl hydrocarbon receptor nuclear translocator 2 affects hypoxia-inducible factor-1 regulatory signaling and metabolism in human breast cancer cells. <i>FEBS Letters</i> , 2011 , 585, 3310-5	3.8	19
347	Crystal structure of an archaeal homologue of multidrug resistance repressor protein, EmrR, from hyperthermophilic archaea Sulfolobus tokodaii strain 7. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007 , 67, 1138-46	4.2	19
346	Cell-free protein synthesis for structure determination by X-ray crystallography. <i>Methods in Molecular Biology</i> , 2010 , 607, 149-60	1.4	19
345	Chemical Changes in the Components of Coffee Beans during Roasting 2015 , 83-91		18
344	Hydrophobic patches on SMAD2 and SMAD3 determine selective binding to cofactors. <i>Science Signaling</i> , 2018 , 11,	8.8	18
343	Identification of functionally important residues of the silkmoth pheromone biosynthesis-activating neuropeptide receptor, an insect ortholog of the vertebrate neuromedin U receptor. <i>Journal of Biological Chemistry</i> , 2014 , 289, 19150-63	5.4	18
342	Cloning of genes and enzymatic characterizations of novel dioscorin isoforms from Dioscorea japonica. <i>Plant Science</i> , 2012 , 183, 14-9	5.3	18
341	Electron microscopic evidence for the myosin head lever arm mechanism in hydrated myosin filaments using the gas environmental chamber. <i>Biochemical and Biophysical Research Communications</i> , 2011 , 405, 651-6	3.4	18
340	On the tryptophan residue of smooth muscle myosin that responds to binding of nucleotide. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 11203-8	11.5	18
339	Heat capacity and entropy changes of the two major isotypes of bullfrog (Rana catesbeiana) parvalbumins induced by calcium binding. <i>Biochemistry</i> , 1987 , 26, 7668-74	3.2	18
338	Distinguishable in vitro binding mode of monomeric TRBP and dimeric PACT with siRNA. <i>PLoS ONE</i> , 2013 , 8, e63434	3.7	18
337	Different Call+-sensitivities between the EF-hands of T- and L-plastins. <i>Biochemical and Biophysical Research Communications</i> , 2012 , 429, 137-41	3.4	17
336	OTOTOXIC EFFECTS OF CARBOPLATIN IN ORGANOTYPIC CULTURES IN CHINCHILLAS AND RATS. Journal of Otology, 2012 , 7, 92-101	1.9	17

335	Interactions among chymotryptic troponin T subfragments, tropomyosin, troponin I and troponin C. <i>Journal of Biochemistry</i> , 1984 , 95, 1417-21	3.1	17
334	Overview of the mechanism of cytoskeletal motors based on structure. <i>Biophysical Reviews</i> , 2018 , 10, 571-581	3.7	17
333	Studies on the regulatory mechanism of isocitrate dehydrogenase 2 using acetylation mimics. <i>Scientific Reports</i> , 2017 , 7, 9785	4.9	16
332	Molecular mechanism of distinct salt-dependent enzyme activity of two halophilic nucleoside diphosphate kinases. <i>Biophysical Journal</i> , 2009 , 96, 4692-700	2.9	16
331	Structural basis for brassinosteroid response by BIL1/BZR1. <i>Nature Plants</i> , 2018 , 4, 771-776	11.5	16
330	Structure and function of a pepstatin-insensitive acid proteinase from Aspergillus niger var. Macrosporus. <i>Advances in Experimental Medicine and Biology</i> , 1991 , 306, 203-11	3.6	16
329	Laminarinase from Flavobacterium sp. reveals the structural basis of thermostability and substrate specificity. <i>Scientific Reports</i> , 2017 , 7, 11425	4.9	15
328	Structure of conjugated polyketone reductase from Candida parapsilosis IFO 0708 reveals conformational changes for substrate recognition upon NADPH binding. <i>Applied Microbiology and Biotechnology</i> , 2014 , 98, 243-9	5.7	15
327	Complex structure of the DNA-binding domain of AdpA, the global transcription factor in Streptomyces griseus, and a target duplex DNA reveals the structural basis of its tolerant DNA sequence specificity. <i>Journal of Biological Chemistry</i> , 2013 , 288, 31019-29	5.4	15
326	Crystal structure of the ATPPase subunit and its substrate-dependent association with the GATase subunit: a novel regulatory mechanism for a two-subunit-type GMP synthetase from Pyrococcus horikoshii OT3. <i>Journal of Molecular Biology</i> , 2010 , 395, 417-29	6.5	15
325	An archaeal Dim2-like protein, aDim2p, forms a ternary complex with a/eIF2 alpha and the 3Mend fragment of 16S rRNA. <i>Journal of Molecular Biology</i> , 2010 , 398, 774-85	6.5	15
324	Fourier transform infrared spectroscopic study on the binding of Mg2+ to a mutant Akazara scallop troponin C (E142Q). <i>Biopolymers</i> , 2004 , 74, 77-81	2.2	15
323	Crystal structure and structural stability of acylphosphatase from hyperthermophilic archaeon Pyrococcus horikoshii OT3. <i>Proteins: Structure, Function and Bioinformatics</i> , 2005 , 61, 196-205	4.2	15
322	Structural optimization of SadA, an Fe(II)- and Exetoglutarate-dependent dioxygenase targeting biocatalytic synthesis of N-succinyl-L-threo-3,4-dimethoxyphenylserine. <i>Biochemical and Biophysical Research Communications</i> , 2014 , 450, 1458-61	3.4	14
321	L-allo-threonine aldolase with an H128Y/S292R mutation from Aeromonas jandaei DK-39 reveals the structural basis of changes in substrate stereoselectivity. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2014 , 70, 1695-703		14
320	The crystal structure of l-sorbose reductase from Gluconobacter frateurii complexed with NADPH and l-sorbose. <i>Journal of Molecular Biology</i> , 2011 , 407, 543-55	6.5	14
319	Characterization of genomic sequence coding for bromelain inhibitors in pineapple and expression of its recombinant isoform. <i>Journal of Biological Chemistry</i> , 2002 , 277, 28222-7	5.4	14
318	Effect of the DNA topoisomerase II inhibitor VP-16 on illegitimate recombination in yeast chromosomes. <i>Gene</i> , 2002 , 291, 251-7	3.8	14

317	Primary Structure, Sequence-Specific 1H-NMR Assignments and Secondary Structure in Solution of Bromelain Inhibitor VI from Pineapple Stem. <i>FEBS Journal</i> , 1995 , 232, 335-343		14
316	Calorimetric studies on calcium and magnesium binding by troponin C from bullfrog skeletal muscle. <i>Journal of Biochemistry</i> , 1990 , 107, 127-32	3.1	14
315	Changes in intracellular pH and inorganic phosphate concentration during and after muscle contraction as studied by time-resolved 31P-NMR. Alkalinization by contraction. <i>FEBS Letters</i> , 1984 , 171, 165-8	3.8	14
314	Complex Mixture Analysis of Organic Compounds in Yogurt by NMR Spectroscopy. <i>Metabolites</i> , 2016 , 6,	5.6	14
313	Rationally Designed Strigolactone Analogs as Antagonists of the D14 Receptor. <i>Plant and Cell Physiology</i> , 2018 , 59, 1545-1554	4.9	13
312	Structure of the microtubule-binding domain of flagellar dynein. <i>Structure</i> , 2014 , 22, 1628-38	5.2	13
311	Crystal structure of TTHA1264, a putative M16-family zinc peptidase from Thermus thermophilus HB8 that is homologous to the beta subunit of mitochondrial processing peptidase. <i>Proteins:</i> Structure, Function and Bioinformatics, 2009 , 75, 774-80	4.2	13
310	Crystal structure of KaiC-like protein PH0186 from hyperthermophilic archaea Pyrococcus horikoshii OT3. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009 , 75, 1035-9	4.2	13
309	Hydrolysis of AMPPNP by the motor domain of ncd, a kinesin-related protein. <i>FEBS Letters</i> , 1997 , 409, 29-32	3.8	13
308	Fourier transform infrared spectroscopic study on the Ca2+ -bound coordination structures of synthetic peptide analogues of the calcium-binding site III of troponin C. <i>Biopolymers</i> , 2006 , 82, 339-43	2.2	13
307	Solution structure and binding specificity of FBP11/HYPA WW domain as Group-II/III. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006 , 63, 227-34	4.2	13
306	Expression, oxidative refolding, and characterization of six-histidine-tagged recombinant human LECT2, a 16-kDa chemotactic protein with three disulfide bonds. <i>Protein Expression and Purification</i> , 2003 , 27, 272-8	2	13
305	Crystal structure of the YjgF/YER057c/UK114 family protein from the hyperthermophilic archaeon Sulfolobus tokodaii strain 7. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006 , 62, 557-61	4.2	13
304	Complexes of myosin subfragment 1 with pyrophosphate and with adenosine diphosphate as studied by phosphorus-31 nuclear magnetic resonance. <i>Journal of Biochemistry</i> , 1993 , 113, 19-21	3.1	13
303	31P-nuclear magnetic resonance studies of intact plasmodia of Physarum polycephalum. <i>FEBS Letters</i> , 1984 , 176, 161-165	3.8	13
302	Structural bases of IMiD selectivity that emerges by 5-hydroxythalidomide. <i>Nature Communications</i> , 2020 , 11, 4578	17.4	13
301	Use of NMR-Based Metabolomics To Chemically Characterize the Roasting Process of Chicory Root. Journal of Agricultural and Food Chemistry, 2016 , 64, 6459-6465	5.7	13
300	Structural basis for receptor-regulated SMAD recognition by MAN1. <i>Nucleic Acids Research</i> , 2018 , 46, 12139-12153	20.1	13

29	99	Yam Tuber Storage Protein Reduces Plant Oxidants Using the Coupled Reactions as Carbonic Anhydrase and Dehydroascorbate Reductase. <i>Molecular Plant</i> , 2015 , 8, 1115-8	14.4	12	
29	98	Organic Compounds in Green Coffee Beans 2015 , 149-162		12	
29	97	Crystal structure of the novel haloalkane dehalogenase DatA from Agrobacterium tumefaciens C58 reveals a special halide-stabilizing pair and enantioselectivity mechanism. <i>Applied Microbiology and Biotechnology</i> , 2014 , 98, 8573-82	5.7	12	
29	96	Effect of low-dose thalidomide on dopaminergic neuronal differentiation of human neural progenitor cells: a combined study of metabolomics and morphological analysis. <i>NeuroToxicology</i> , 2012 , 33, 1375-80	4.4	12	
29	95	Structural units important for activity of a novel-type phosphoserine phosphatase from Hydrogenobacter thermophilus TK-6 revealed by crystal structure analysis. <i>Journal of Biological Chemistry</i> , 2013 , 288, 11448-58	5.4	12	
29	94	Solution structure of a short-chain insecticidal toxin LaIT1 from the venom of scorpion Liocheles australasiae. <i>Biochemical and Biophysical Research Communications</i> , 2011 , 411, 738-44	3.4	12	
29	93	Isolation, cDNA cloning, and structure-based functional characterization of oryctin, a hemolymph protein from the coconut rhinoceros beetle, Oryctes rhinoceros, as a novel serine protease inhibitor. <i>Journal of Biological Chemistry</i> , 2010 , 285, 30150-8	5.4	12	
29	92	Purification and characterization of CEP from Lactococcus lactis ssp. lactis. <i>Food Chemistry</i> , 2009 , 112, 533-538	8.5	12	
29	91	Modulation of actin filament sliding by mutations of the SH2 cysteine in Dictyostelium myosin II. <i>Biochemical and Biophysical Research Communications</i> , 1997 , 234, 701-6	3.4	12	
29	90	Specificity of the synergistic anion for iron binding by ferric binding protein from Neisseria gonorrhoeae. <i>Biochemistry</i> , 2004 , 43, 9195-203	3.2	12	
28	89	The effect of amphiphilic additives on the growth and morphology of Aspergillus niger acid proteinase A crystals. <i>Journal of Crystal Growth</i> , 2002 , 234, 247-254	1.6	12	
28	88	The three-dimensional structure of aspergilloglutamic peptidase from Aspergillus niger. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2004 , 80, 435-438	4	12	
28	87	Structure-function relationship of bromelain isoinhibitors from pineapple stem. <i>Biological Chemistry</i> , 2002 , 383, 1151-6	4.5	12	
28	86	Gene structure and amino acid sequence of Latimeria chalumnae (coelacanth) myelin DM20: phylogenetic relation of the fish. <i>Neurochemical Research</i> , 1999 , 24, 867-73	4.6	12	
28	85	Crystallization and preliminary X-ray investigation of proteinase A, a non-pepsin-type acid proteinase from Aspergillus niger var. macrosporus. <i>Journal of Molecular Biology</i> , 1992 , 223, 373-5	6.5	12	
2	84	Primary structure of chymotryptic subfragments from rabbit skeletal troponin T. <i>Journal of Biochemistry</i> , 1981 , 90, 263-5	3.1	12	
28	83	Post-contractile phosphocreatine splitting in muscle as revealed by time-resolved 31P nuclear magnetic resonance. <i>The Japanese Journal of Physiology</i> , 1983 , 33, 909-19		12	
28	82	Crystal structure of dibenzothiophene sulfone monooxygenase BdsA from Bacillus subtilis WU-S2B. <i>Proteins: Structure, Function and Bioinformatics</i> , 2017 , 85, 1171-1177	4.2	11	

281	Structural basis for high substrate-binding affinity and enantioselectivity of 3-quinuclidinone reductase AtQR. <i>Biochemical and Biophysical Research Communications</i> , 2014 , 446, 911-5	3.4	11
280	The Arginine Residue within the C-Terminal Active Core of Bombyx mori Pheromone Biosynthesis-Activating Neuropeptide is Essential for Receptor Binding and Activation. <i>Frontiers in Endocrinology</i> , 2012 , 3, 42	5.7	11
279	Structure of TTHA1623, a novel metallo-beta-lactamase superfamily protein from Thermus thermophilus HB8. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 455-9		11
278	Crystallization and preliminary X-ray analyses of desulfurization enzyme DszB and its C27S mutant complexed with biphenyl-2-sulfinic acid. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 1636-8		11
277	Purification and characterization of an aminopeptidase from Lactobacillus helveticus JCM 1004. <i>Food Chemistry</i> , 2004 , 88, 511-516	8.5	11
276	Purification and characterization of a novel prolyl endopeptidase from Aspergillus niger. Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 2005, 81, 447-453	4	11
275	The amino acid sequences of isoforms of the bromelain inhibitor from pineapple stem. <i>Journal of Biochemistry</i> , 1998 , 124, 457-61	3.1	11
274	Oligomerization mechanisms of an H-NS family protein, Pmr, encoded on the plasmid pCAR1 provide a molecular basis for functions of H-NS family members. <i>PLoS ONE</i> , 2014 , 9, e105656	3.7	11
273	Multivariate analyses of the physicochemical properties of turnip (Brassica rapa L.) chips dried using different methods. <i>Drying Technology</i> , 2020 , 38, 411-419	2.6	11
272	Deficiency Does Not Affect the Cytosolic Glutathione or Thioredoxin Antioxidant Defense in Mouse Cochlea. <i>Journal of Neuroscience</i> , 2017 , 37, 5770-5781	6.6	10
271	Crystal Structure of Human Leukocyte Cell-derived Chemotaxin 2 (LECT2) Reveals a Mechanistic Basis of Functional Evolution in a Mammalian Protein with an M23 Metalloendopeptidase Fold. Journal of Biological Chemistry, 2016 , 291, 17133-42	5.4	10
270	Splenic stromal cells from aged mice produce higher levels of IL-6 compared to young mice. <i>Mediators of Inflammation</i> , 2014 , 2014, 826987	4.3	10
269	Calcium-dependent structural changes in human reticulocalbin-1. <i>Journal of Biochemistry</i> , 2014 , 155, 281-93	3.1	10
268	Establishment of Sf9 Transformants Constitutively Expressing PBAN Receptor Variants: Application to Functional Evaluation. <i>Frontiers in Endocrinology</i> , 2012 , 3, 56	5.7	10
267	Stimulatory effects of arachidonic acid on myosin ATPase activity and contraction of smooth muscle via myosin motor domain. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2010 , 298, H505-14	5.2	10
266	The structure of mAG, a monomeric mutant of the green fluorescent protein Azami-Green, reveals the structural basis of its stable green emission. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010 , 66, 485-9		10
265	Structure of FBP11 WW1-PL ligand complex reveals the mechanism of proline-rich ligand recognition by group II/III WW domains. <i>Journal of Biological Chemistry</i> , 2006 , 281, 40321-9	5.4	10
264	Stalk region of kinesin-related protein Unc104 has moderate ability to form coiled-coil dimer. <i>Biochemical and Biophysical Research Communications</i> , 2005 , 337, 868-74	3.4	10

263	Molecular cloning of the Lon protease gene from Thermus thermophilus HB8 and characterization of its gene product. <i>FEBS Journal</i> , 1999 , 266, 811-9		10
262	Preparation and characterization of two major isotypes of parvalbumins from skeletal muscle of bullfrog (Rana catesbeiana). <i>Journal of Biochemistry</i> , 1986 , 99, 1211-8	3.1	10
261	Biochemical characterization and structural analysis of ulvan lyase from marine Alteromonas sp. reveals the basis for its salt tolerance. <i>International Journal of Biological Macromolecules</i> , 2020 , 147, 13	30 9 -93 [.]	17 ¹⁰
260	Real-Time Monitoring of Chemical Changes in Three Kinds of Fermented Milk Products during Fermentation Using Quantitative Difference Nuclear Magnetic Resonance Spectroscopy. <i>Journal of Agricultural and Food Chemistry</i> , 2018 , 66, 1479-1487	5.7	9
259	Leukocyte cell-derived chemotaxin 2 is a zinc-binding protein. FEBS Letters, 2013, 587, 404-9	3.8	9
258	High pressure refolding, purification, and crystallization of flavin reductase from Sulfolobus tokodaii strain 7. <i>Protein Expression and Purification</i> , 2012 , 84, 214-8	2	9
257	Expression, high-pressure refolding and purification of human leukocyte cell-derived chemotaxin 2 (LECT2). <i>Protein Expression and Purification</i> , 2013 , 88, 221-9	2	9
256	F2-selective two-dimensional NMR spectroscopy for the analysis of minor components in foods. <i>Magnetic Resonance in Chemistry</i> , 2011 , 49, 710-6	2.1	9
255	Reversible and irreversible coiled coils in the stalk domain of ncd motor protein. <i>Biochemistry</i> , 2007 , 46, 9523-32	3.2	9
254	Crystallization and preliminary X-ray analysis of ginkbilobin-2 from Ginkgo biloba seeds: a novel antifungal protein with homology to the extracellular domain of plant cysteine-rich receptor-like kinases. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007 , 63, 737-9		9
253	The dynein stalk head, the microtubule binding-domain of dynein: NMR assignment and ligand binding. <i>Journal of Biomolecular NMR</i> , 2008 , 41, 89-96	3	9
252	Structure of calcium-bound human S100A13 at pH 7.5 at 1.8 A resolution. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2008 , 64, 70-6		9
251	Atomically detailed description of the unfolding of alpha-lactalbumin by the combined use of experiments and simulations. <i>Journal of Molecular Biology</i> , 2005 , 354, 164-72	6.5	9
250	Functional importance of Ca2+-deficient N-terminal lobe of molluscan troponin C in troponin regulation. <i>Archives of Biochemistry and Biophysics</i> , 2005 , 436, 83-90	4.1	9
249	Crystallization and preliminary X-ray analysis of HCE-1, a hatching enzyme of medaka fish, Oryzias latipes. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 725-6		9
248	Crystal structure of monomeric sarcosine oxidase from Bacillus sp. NS-129 reveals multiple conformations at the active-site loop. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2005 , 81, 220-224	4	9
247	Gene organization for nitric oxide reduction in Alcaligenes faecalis S-6. <i>Bioscience, Biotechnology and Biochemistry</i> , 2000 , 64, 852-7	2.1	9
246	Heat capacity and entropy changes of troponin C from bullfrog skeletal muscle induced by calcium binding. <i>FEBS Journal</i> , 1990 , 192, 275-81		9

245	A novel mode of ferric ion coordination by the periplasmic ferric ion-binding subunit FbpA of an ABC-type iron transporter from Thermus thermophilus HB8. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2014 , 70, 196-202		9
244	Increased burden of mitochondrial DNA deletions and point mutations in early-onset age-related hearing loss in mitochondrial mutator mice. <i>Experimental Gerontology</i> , 2019 , 125, 110675	4.5	8
243	Structural analysis of enzymes used for bioindustry and bioremediation. <i>Bioscience, Biotechnology and Biochemistry</i> , 2015 , 79, 1391-401	2.1	8
242	NMR-based analysis of the chemical composition of Japanese persimmon aqueous extracts. <i>Magnetic Resonance in Chemistry</i> , 2016 , 54, 213-21	2.1	8
241	Application of High-Field Superconducting Magnet to Protein Crystallization. <i>Physics Procedia</i> , 2012 , 36, 953-957		8
240	Investigation of chemical structure of nonprotein proteinase inhibitors from dried figs. <i>Food Chemistry</i> , 2008 , 107, 305-311	8.5	8
239	Metal-free and Ca2+-bound structures of a multidomain EF-hand protein, CBP40, from the lower eukaryote Physarum polycephalum. <i>Structure</i> , 2003 , 11, 75-85	5.2	8
238	Crystallization and preliminary X-ray analysis of AzoR (azoreductase) from Escherichia coli. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2005 , 61, 399-402		8
237	Rapid determination of parvalbumin amino acid sequence from Rana catesbeiana (pl 4.78) by combination of ESI mass spectrometry, protein sequencing, and amino acid analysis. <i>Journal of Biochemistry</i> , 2000 , 127, 723-9	3.1	8
236	Differential scanning calorimetric studies of the thermal unfolding of acid proteinase A from Aspergillus niger at various pHs. <i>Thermochimica Acta</i> , 1995 , 267, 373-378	2.9	8
235	Enhancement of the turnover number of thermostable malate dehydrogenase by deleting hydrogen bonds around the catalytic site. <i>Biochemical and Biophysical Research Communications</i> , 1996 , 225, 844-8	3.4	8
234	A new crystal form of proteinase A, a non-pepsin-type acid proteinase from Aspergillus niger var. macrosporus. <i>Journal of Biochemistry</i> , 1993 , 114, 457-8	3.1	8
233	Protein sequence determination by ESI-MS and LSI-MS tandem mass spectrometry: parvalbumin primary structures from cat, gerbil and monkey skeletal muscle. <i>BBA - Proteins and Proteomics</i> , 1992 , 1160, 1-7		8
232	Fourier-transform infrared and circular dichroism studies on temperature dependence of the secondary structure of troponin C from bullfrog skeletal muscle. <i>Canadian Journal of Chemistry</i> , 1991 , 69, 1679-1684	0.9	8
231	Preparation and characterization of the major isotype of parvalbumin from skeletal muscle of the toad (Bufo bufo japonicus). <i>Journal of Biochemistry</i> , 1987 , 102, 1133-9	3.1	8
230	Structural basis for calcium-regulated relaxation of striated muscles at interaction sites of troponin with actin and tropomyosin. <i>Advances in Experimental Medicine and Biology</i> , 2007 , 592, 71-86	3.6	8
229	Different DNA-binding specificities of NLP and NIN transcription factors underlie nitrate-induced control of root nodulation. <i>Plant Cell</i> , 2021 , 33, 2340-2359	11.6	8
228	Functional properties of Chinese yam (Thunb. cv. Baiyu) soluble protein. <i>Journal of Food Science and Technology</i> , 2018 , 55, 381-388	3.3	8

(2008-2003)

227	NMR structural study of troponin C C-terminal domain complexed with troponin I fragment from Akazara scallop. <i>Advances in Experimental Medicine and Biology</i> , 2003 , 538, 195-200; discussion 201	3.6	8
226	Engineering a short-chain dehydrogenase/reductase for the stereoselective production of (2S,3R,4S)-4-hydroxyisoleucine with three asymmetric centers. <i>Scientific Reports</i> , 2017 , 7, 13703	4.9	7
225	Structure and reaction mechanism of a novel enone reductase. FEBS Journal, 2015, 282, 1526-37	5.7	7
224	Expression, purification, refolding, and enzymatic characterization of two secretory phospholipases Alfrom Neurospora crassa. <i>Protein Expression and Purification</i> , 2015 , 115, 69-75	2	7
223	Structural basis for substrate recognition and processive cleavage mechanisms of the trimeric exonuclease PhoExo I. <i>Nucleic Acids Research</i> , 2015 , 43, 7122-36	20.1	7
222	Functional production of human antibody by the filamentous fungus. <i>Fungal Biology and Biotechnology</i> , 2020 , 7, 7	7.5	7
221	Quantification of terpene trilactones in Ginkgo biloba with a H NMR method. <i>Journal of Natural Medicines</i> , 2018 , 72, 793-797	3.3	7
220	Crystal structure of a crustacean hyperglycemic hormone (CHH) precursor suggests structural variety in the C-terminal regions of CHH superfamily members. <i>FEBS Journal</i> , 2016 , 283, 4325-4339	5.7	7
219	In-situ and real-time growth observation of high-quality protein crystals under quasi-microgravity on earth. <i>Scientific Reports</i> , 2016 , 6, 22127	4.9	7
218	Structural basis of stereospecific reduction by quinuclidinone reductase. <i>AMB Express</i> , 2014 , 4, 6	4.1	7
217	Evaluation of nonionic adsorbent resins for removal of inhibitory compounds from corncob hydrolysate for ethanol fermentation. <i>Bioresource Technology</i> , 2013 , 149, 541-5	11	7
216	Electron microscopic recording of myosin head power stroke in hydrated myosin filaments. <i>Scientific Reports</i> , 2015 , 5, 15700	4.9	7
215	Attenuation of migration properties of CD4+ T cells from aged mice correlates with decrease in chemokine receptor expression, response to retinoic acid, and RALDH expression compared to young mice. <i>Bioscience, Biotechnology and Biochemistry</i> , 2014 , 78, 976-80	2.1	7
214	Ototoxic Model of Oxaliplatin and Protection from Nicotinamide Adenine Dinucleotide. <i>Journal of Otology</i> , 2013 , 8, 63-71	1.9	7
213	Structure of the Ca(2+)-saturated C-terminal domain of scallop troponin C in complex with a troponin I fragment. <i>Biological Chemistry</i> , 2013 , 394, 55-68	4.5	7
212	Antifungal and Lipid Transfer Proteins from Ginkgo (Ginkgo biloba) Seeds 2011 , 527-534		7
211	Regulatory mechanism of abscisic acid signaling. <i>Biophysics (Nagoya-shi, Japan)</i> , 2011 , 7, 123-128		7
210	Infrared spectroscopic study of the binding of divalent cations to Akazara scallop troponin C: the effect of the methylene side chain of glutamate residue. <i>Biopolymers</i> , 2008 , 89, 595-9	2.2	7

209	Nuclear magnetic resonance studies on the pKa values and interaction of ionizable groups in bromelain inhibitor VI from pineapple stem. <i>Biological Chemistry</i> , 2003 , 384, 93-104	4.5	7
208	A proposed catalytic mechanism of aspergilloglutamic peptidase from Aspergillus niger. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2005 , 81, 441-446	4	7
207	A phosphorus-31 nuclear magnetic resonance study on the complex of chicken gizzard myosin subfragment 1 with adenosine diphosphate 1999 , 190, 75-78		7
206	Crystal structure of ribonuclease T1 carboxymethylated at Glu58 in complex with 2MGMP. <i>Biochemistry</i> , 1996 , 35, 8329-34	3.2	7
205	Tertiary Structure Comparison of Bromelain Inhibitor VI from Pineapple Stem with Bovine Pancreatic Trypsin Inhibitor <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1996 , 72, 104-107	4	7
204	Preparation and characterization of troponin C from bullfrog skeletal muscle. <i>Journal of Biochemistry</i> , 1992 , 112, 800-3	3.1	7
203	Tetrameric structure of the restriction DNA glycosylase R.Pabl in complex with nonspecific double-stranded DNA. <i>Scientific Reports</i> , 2016 , 6, 35197	4.9	7
202	Comprehensive NMR analysis of two kinds of post-fermented tea and their anti-glycation activities in vitro. <i>Food Chemistry</i> , 2019 , 277, 735-743	8.5	7
201	Foliar application is an effective method for incorporating selenium into peanut leaf proteins with antioxidant activities. <i>Food Research International</i> , 2019 , 126, 108617	7	6
200	GSR is not essential for the maintenance of antioxidant defenses in mouse cochlea: Possible role of the thioredoxin system as a functional backup for GSR. <i>PLoS ONE</i> , 2017 , 12, e0180817	3.7	6
199	REFOLDdb: a new and sustainable gateway to experimental protocols for protein refolding. <i>BMC Structural Biology</i> , 2017 , 17, 4	2.7	6
198	Human CTP:phosphoethanolamine cytidylyltransferase: enzymatic properties and unequal catalytic roles of CTP-binding motifs in two cytidylyltransferase domains. <i>Biochemical and Biophysical Research Communications</i> , 2014 , 449, 26-31	3.4	6
197	Definite differences between in vitro actin-myosin sliding and muscle contraction as revealed using antibodies to myosin head. <i>PLoS ONE</i> , 2014 , 9, e93272	3.7	6
196	Crystallization and preliminary X-ray analysis of human leukocyte cell-derived chemotaxin 2 (LECT2). <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013 , 69, 316-9		6
195	Structure of flap endonuclease 1 from the hyperthermophilic archaeon Desulfurococcus amylolyticus. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2011 , 67, 209-13		6
194	Crystal structure of curcuminoid synthase CUS from Oryza sativa. <i>Proteins: Structure, Function and Bioinformatics</i> , 2011 , 79, 669-73	4.2	6
193	Reduction of MPO-ANCA epitopes in SCG/Kj mice by 15-deoxyspergualin treatment restricted by IgG2b associated with crescentic glomerulonephritis. <i>Rheumatology</i> , 2010 , 49, 1245-56	3.9	6
192	Crystallization and preliminary X-ray analysis of ZHE1, a hatching enzyme from the zebrafish Danio rerio. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 1018-20		6

191	Evidence for existence of multiple conformations of kinesin and ncd motor domains in solution revealed by 31P-NMR of the tightly bound ADP. <i>FEBS Journal</i> , 1998 , 257, 466-71		6
190	Crystal structure of thioredoxin domain of ST2123 from thermophilic archaea Sulfolobus tokodaii strain7. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007 , 69, 204-8	4.2	6
189	Characterization of the acidic and basic limbs of a bell-shaped pH profile in the inhibitory activity of bromelain inhibitor VI. <i>Biopolymers</i> , 2006 , 81, 309-19	2.2	6
188	Expression and purification of active WW domains of FBP11/HYPA and FBP28/CA150. <i>Protein and Peptide Letters</i> , 2006 , 13, 197-201	1.9	6
187	Coiled coil in the stalk region of ncd motor protein is nonlocally sustained. <i>Biochemistry</i> , 2006 , 45, 3315-	-3,42	6
186	Infrared spectroscopic study on Ca2+ binding to Akazara scallop troponin C in comparison with peptide analogues of its Ca2+-binding Site IV. <i>Vibrational Spectroscopy</i> , 2006 , 42, 188-191	2.1	6
185	Determination of the NMR structure of Gln25-ribonuclease T1. <i>Biological Chemistry</i> , 2003 , 384, 1173-83	4.5	6
184	Crystallization and preliminary crystallographic analysis of the major NAD(P)H: FMN oxidoreductase of Vibrio fischeri ATCC 7744. <i>Journal of Structural Biology</i> , 1996 , 117, 70-2	3.4	6
183	Crystallization and preliminary X-ray analysis of brazzein, a new sweet protein. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1996 , 52, 577-8		6
182	In vivo Antihypertensive Effect of Val-Glu-Pro in Spontaneously Hypertensive Rats*. <i>Progress in Biochemistry and Biophysics</i> , 2011 , 38, 353-360		6
181	Antioxidant properties and inhibition of angiotensin-converting enzyme by highly active peptides from wheat gluten. <i>Scientific Reports</i> , 2021 , 11, 5206	4.9	6
180	Insight into the transition between the open and closed conformations of Thermus thermophilus carboxypeptidase. <i>Biochemical and Biophysical Research Communications</i> , 2017 , 484, 787-793	3.4	5
179	Characterization of the Ca-coordination structures of L- and T-plastins in combination with their synthetic peptide analogs by FTIR spectroscopy. <i>Scientific Reports</i> , 2019 , 9, 4217	4.9	5
178	Structural Basis for Action of the External Chaperone for a Propeptide-deficient Serine Protease from Aeromonas sobria. <i>Journal of Biological Chemistry</i> , 2015 , 290, 11130-43	5.4	5
177	Crystal structure of the novel lesion-specific endonuclease PfuEndoQ from Pyrococcus furiosus. <i>Nucleic Acids Research</i> , 2018 , 46, 4807-4818	20.1	5
176	Crystal structure of a Ca-dependent regulator of flagellar motility reveals the open-closed structural transition. <i>Scientific Reports</i> , 2018 , 8, 2014	4.9	5
175	Broadband WET: a novel technique for quantitative characterization of minor components in foods. <i>Magnetic Resonance in Chemistry</i> , 2014 , 52, 333-8	2.1	5
174	Crystal structure of conjugated polyketone reductase (CPR-C1) from Candida parapsilosis IFO 0708 complexed with NADPH. <i>Proteins: Structure, Function and Bioinformatics</i> , 2013 , 81, 2059-63	4.2	5

173	Preparation of recombinant six-histidine-tagged human LECT2, a chemotactic protein to neutrophils, in Escherichia coli. <i>Cytotechnology</i> , 1997 , 25, 235-8	2.2	5
172	Cloning, expression, purification, crystallization and preliminary crystallographic analysis of selenomethionine-labelled KaiC-like protein PH0186 from Pyrococcus horikoshii OT3. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007 , 63, 327-9		5
171	Crystal structure of TTHA0303 (TT2238), a four-helix bundle protein with an exposed histidine triad from Thermus thermophilus HB8 at 2.0 A. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 70, 1103	47	5
170	A unique catalytic triad revealed by the crystal structure of APE0912, a short-chain dehydrogenase/reductase family protein from Aeropyrum pernix K1. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 70, 1640-5	4.2	5
169	Crystal structure of STS042, a stand-alone RAM module protein, from hyperthermophilic archaeon Sulfolobus tokodaii strain 7. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 71, 1557-62	4.2	5
168	Structural characterisation of PinA WW domain and a comparison with other group IV WW domains, Pin1 and Ess1. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2008 , 1784, 1208-14	4	5
167	Crystallization and preliminary X-ray analysis of the PIN domain of human EST1A. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006 , 62, 656-8		5
166	Susceptibility of the interchain peptide of a bromelain inhibitor precursor to the target proteases bromelain, chymotrypsin, and trypsin. <i>Biological Chemistry</i> , 2005 , 386, 491-8	4.5	5
165	Kinetic and mutation analyses of aspartate kinase from Thermus flavus. <i>Journal of Bioscience and Bioengineering</i> , 1999 , 87, 739-45	3.3	5
164	UV-laser-induced changes in the infrared spectrum of calmodulin in the presence of a photosensitive Ca2+ chelator. <i>Journal of Molecular Structure</i> , 1994 , 324, 107-112	3.4	5
163	Heat capacity and entropy changes of the major isotype of the toad (Bufo) parvalbumin induced by calcium binding. <i>FEBS Journal</i> , 1990 , 188, 23-8		5
162	The complete amino acid sequence of bullfrog (Rana catesbeiana) parvalbumin pl4.97. <i>FEBS Letters</i> , 1990 , 268, 249-51	3.8	5
161	New approach to generating insights for aging research based on literature mining and knowledge integration. <i>PLoS ONE</i> , 2017 , 12, e0183534	3.7	5
160	Purification and Characterization of an Angiotensin I-converting Enzyme Inhibitory Peptide Derived From Spirulina platensis*. <i>Progress in Biochemistry and Biophysics</i> , 2010 , 37, 568-574		5
159	Treatment with hydrogen peroxide improves the physicochemical properties of dietary fibres from Chinese yam peel. <i>International Journal of Food Science and Technology</i> , 2020 , 55, 1289-1297	3.8	5
158	Development of Engineered Ferredoxin Reductase Systems for the Efficient Hydroxylation of Steroidal Substrates. <i>ACS Sustainable Chemistry and Engineering</i> , 2020 , 8, 16720-16730	8.3	5
157	Effect of chemical and enzymatic modifications on the structural and physicochemical properties of dietary fiber from purple turnip (Brassica rapa L.). <i>LWT - Food Science and Technology</i> , 2021 , 145, 111313	5.4	5
156	Structural basis for the regulation of phytohormone receptors. <i>Bioscience, Biotechnology and Biochemistry</i> , 2017 , 81, 1261-1273	2.1	4

155	Quantitation of Minor Components in Mango Juice with Band-Selective Excitation NMR Spectroscopy. <i>Journal of Agricultural and Food Chemistry</i> , 2017 , 65, 9547-9552	5.7	4
154	BC-Box Motif-Mediated Neuronal Differentiation of Somatic Stem Cells. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	4
153	NMR-based metabolic profiling and comparison of Japanese persimmon cultivars. <i>Scientific Reports</i> , 2019 , 9, 15011	4.9	4
152	Purification, crystallization and preliminary X-ray analysis of the DNA-binding domain of AdpA, the central transcription factor in the A-factor regulatory cascade in the filamentous bacterium Streptomyces griseus, in complex with a duplex DNA. <i>Acta Crystallographica Section F: Structural</i>		4
151	Expression, purification, crystallization and X-ray analysis of 3-quinuclidinone reductase from Agrobacterium tumefaciens. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 1237-9		4
150	Coordination structures of Mg2+ and Ca2+ in three types of tobacco calmodulins in solution: Fourier-transform infrared spectroscopic studies of side-chain COO- groups. <i>Biopolymers</i> , 2013 , 99, 472-	- 83	4
149	Crystallization and preliminary X-ray analysis of crustacean hyperglycaemic hormone from the kuruma prawn Marsupenaeus japonicus in its weakly active precursor form. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2011 , 67, 1586-9		4
148	Crystallization and preliminary X-ray analysis of gamma-hexachlorocyclohexane dehydrochlorinase LinA from Sphingobium japonicum UT26. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 822-4		4
147	Crystallization and preliminary X-ray analysis of a monomeric mutant of Azami-Green (mAG), an Aequorea victoria green fluorescent protein-like green-emitting fluorescent protein from the stony coral Galaxea fascicularis. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 ,		4
146	65, 1292-5 Binding of an RNA pol II Ligand to the WW Domain of Pin1 Using Molecular Dynamics Docking Simulations. <i>Journal of Chemical Theory and Computation</i> , 2009 , 5, 2886-97	6.4	4
145	Detection of apoptosis by RT B CR array in mefloquineInduced cochlear damage. <i>Journal of Otology</i> , 2011 , 6, 1-9	1.9	4
144	The crystal structure of an intermediate dimer of aspergilloglutamic peptidase that mimics the enzyme-activation product complex produced upon autoproteolysis. <i>Journal of Biochemistry</i> , 2012 , 152, 45-52	3.1	4
143	Spectroscopic and ITC study of the conformational change upon Ca2+-binding in TnC C-lobe and TnI peptide complex from Akazara scallop striated muscle. <i>Biochemical and Biophysical Research Communications</i> , 2008 , 369, 109-14	3.4	4
142	Crystal structure of Dim2p: a preribosomal RNA processing factor, from Pyrococcus horikoshii OT3 at 2.30 A. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007 , 69, 428-32	4.2	4
141	Crystal structure of TTHA1429, a novel metallo-beta-lactamase superfamily protein from Thermus thermophilus HB8. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 73, 1053-7	4.2	4
140	Temperature-induced chirality reversal of induced circular dichroism of premicellar aggregates of acridine orange derivatives and dodecanoyl-l-threonine in aqueous solution. <i>Colloid and Polymer Science</i> , 2006 , 284, 1453-1458	2.4	4
139	Crystal structure of azoreductase AzoR from Escherichia coli. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2005 , 81, 225-228	4	4
138	Preliminary X-ray crystallographic analysis of thermolysin in the presence of 4 M NaCl. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2005 , 61, 710-2		4

137	Phase diagram of crystallization of Aspergillus niger acid proteinase A, a non-pepsin-type acid proteinase. <i>Journal of Crystal Growth</i> , 1996 , 168, 118-123	1.6	4
136	Thermal stabilization of ribonuclease T1 by carboxymethylation at Glu-58 as revealed by 1H nuclear magnetic resonance spectroscopy. <i>FEBS Letters</i> , 1994 , 351, 389-92	3.8	4
135	Effect of Dimethylsulfoxide on the Crystallization of Aspergillus niger Proteinase A <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1991 , 67, 209-212	4	4
134	1H-Nuclear magnetic resonance study on imidazole imidazole interaction in L-histidyl-L-histidine and D-histidyl-L-histidine: Analysis including microscopic dissociation series. <i>Biopolymers</i> , 1983 , 22, 2563	- 2 2 5 76	4
133	Location of troponin I-binding on troponin T sequence. FEBS Letters, 1982, 145, 147-9	3.8	4
132	NIGT1 family proteins exhibit dual mode DNA recognition to regulate nutrient response-associated genes in Arabidopsis. <i>PLoS Genetics</i> , 2020 , 16, e1009197	6	4
131	Crystal structure of the complex of the interaction domains of Escherichia coli DnaB helicase and DnaC helicase loader: structural basis implying a distortion-accumulation mechanism for the DnaB ring opening caused by DnaC binding. <i>Journal of Biochemistry</i> , 2020 , 167, 1-14	3.1	4
130	Structural basis for transcriptional coactivator recognition by SMAD2 in TGF-Isignaling. <i>Science Signaling</i> , 2020 , 13,	8.8	4
129	Metabolic profiling of natural and cultured Cordyceps by NMR spectroscopy. <i>Scientific Reports</i> , 2019 , 9, 7735	4.9	3
128	Expression, high-pressure refolding, purification, crystallization and preliminary X-ray analysis of a novel single-strand-specific 3M5Mexonuclease PhoExo I from Pyrococcus horikoshii OT3. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2014 , 70, 1076-9	1.1	3
127	Purification, crystallization and preliminary X-ray analysis of OsAREB8 from rice, a member of the AREB/ABF family of bZIP transcription factors, in complex with its cognate DNA. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 491-4		3
126	A thermoacidophile-specific protein family, DUF3211, functions as a fatty acid carrier with novel binding mode. <i>Journal of Bacteriology</i> , 2013 , 195, 4005-12	3.5	3
125	Infrared study of synthetic peptide analogues of the calcium-binding site III of troponin C: The role of helix F of an EF-hand motif. <i>Biopolymers</i> , 2013 , 99, 342-7	2.2	3
124	Purification, crystallization and preliminary X-ray analysis of L-sorbose reductase from Gluconobacter frateurii complexed with L-sorbose or NADPH. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 562-4		3
123	Crystallization and preliminary X-ray analysis of the NADPH-dependent 3-quinuclidinone reductase from Rhodotorula rubra. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 645-7		3
122	Absolute side-chain structure at position 13 is required for the inhibitory activity of bromein. Journal of Biological Chemistry, 2008 , 283, 36338-43	5.4	3
121	PinA from Aspergillus nidulans binds to pS/pT-P motifs using the same Loop I and XP groove as mammalian Pin1. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2007 , 1774, 1208-12	4	3
120	Crystal structure of the proline iminopeptidase-related protein TTHA1809 from Thermus thermophilus HB8. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 70, 1646-9	4.2	3

119	Crystallization and preliminary X-ray analysis of CTP:phosphoethanolamine cytidylyltransferase (ECT) from Saccharomyces cerevisiae. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006 , 62, 1003-5		3
118	Cloning, purification, crystallization and preliminary crystallographic analysis of acylphosphatase from Pyrococcus horikoshii OT3. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 1135-6		3
117	Type I restriction enzyme with RecA protein promotes illegitimate recombination. <i>Plasmid</i> , 2003 , 50, 202-12	3.3	3
116	Identification of arginine residues important for the activity of Escherichia coli signal peptidase I. <i>Biological Chemistry</i> , 2004 , 385, 381-8	4.5	3
115	Crystallization and preliminary X-ray analysis of the YjgF/YER057c/UK114-family protein ST0811 from Sulfolobus tokodaii strain 7. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2005 , 61, 828-30		3
114	Characterization of native and recombinant peptidyl prolyl cis-trans isomerases derived from Methanococcus thermolithotrophicus based on cDNA sequence. <i>Electrophoresis</i> , 2000 , 21, 1733-9	3.6	3
113	Crystal growth and solubility studies of Aspergillus niger acid proteinase A in the presence of both a salt and an organic solvent. <i>Journal of Crystal Growth</i> , 1999 , 196, 291-296	1.6	3
112	Crystallization and preliminary X-ray diffraction studies of a 40 kDa calcium binding protein specifically expressed in plasmodia of Physarum polycephalum. <i>Journal of Biochemistry</i> , 1999 , 126, 7-9	3.1	3
111	Identification of novel interacting regions involving calcineurin and nuclear factor of activated T cells. <i>FASEB Journal</i> , 2020 , 34, 3197-3208	0.9	3
110	Solubility and emulsifying activity of yam soluble protein. <i>Journal of Food Science and Technology</i> , 2020 , 57, 1619-1627	3.3	3
109	Gene co-expression network analysis identifies BEH3 as a stabilizer of secondary vascular development in Arabidopsis. <i>Plant Cell</i> , 2021 , 33, 2618-2636	11.6	3
108	Molecular basis of strigolactone perception in root-parasitic plants: aiming to control its germination with strigolactone agonists/antagonists. <i>Cellular and Molecular Life Sciences</i> , 2020 , 77, 110.	3 ¹ 71713	3 ³
107	Intestinal regulatory T cell induction by \Box -elemene alleviates the formation of fat tissue-related inflammation. <i>IScience</i> , 2021 , 24, 101883	6.1	3
106	Hydrogen peroxide modification affects the structure and physicochemical properties of dietary fibers from white turnip (Brassica Rapa L.). <i>Scientific Reports</i> , 2021 , 11, 1024	4.9	3
105	NMR spectra of PB2 627, the RNA-binding domain in influenza A virus RNA polymerase that contains the pathogenicity factor lysine 627, and improvement of the spectra by small osmolytes. <i>Biochemistry and Biophysics Reports</i> , 2017 , 12, 129-134	2.2	2
104	Recovery of Yam Soluble Protein from Yam Starch Processing Wastewater. <i>Scientific Reports</i> , 2020 , 10, 5384	4.9	2
103	Expression, purification, crystallization and preliminary X-ray analysis of carbonyl reductase S1 from Candida magnoliae. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 540-2		2
102	Crystallization and preliminary X-ray analysis of the haloalkane dehalogenase DatA from Agrobacterium tumefaciens C58. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 652-4		2

101	Expression, purification, crystallization and preliminary X-ray analysis of a novel N-substituted branched-chain L-amino-acid dioxygenase from Burkholderia ambifaria AMMD. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 1067-9		2
100	Expression, purification, crystallization and preliminary X-ray analysis of the KaiC-like protein PH0187 from the hyperthermophilic archaeon Pyrococcus horikoshii OT3. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2011 , 67, 144-6		2
99	Crystallization and preliminary X-ray analysis of flap endonuclease 1 (FEN1) from Desulfurococcus amylolyticus. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 923-5		2
98	Expression, purification, crystallization and preliminary X-ray analysis of conjugated polyketone reductase C2 (CPR-C2) from Candida parapsilosis IFO 0708. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009 , 65, 1145-8		2
97	Crystallization and preliminary crystallographic analysis of the sarcosine oxidase from Bacillus sp. NS-129. <i>Journal of Structural Biology</i> , 1997 , 120, 109-11	3.4	2
96	Crystallization and preliminary X-ray analysis of the Ca2+-bound C-terminal lobe of troponin C in complex with a troponin I-derived peptide fragment from Akazara scallop. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007 , 63, 535-7		2
95	Crystal structure of hypothetical protein PH0734.1 from hyperthermophilic archaea Pyrococcus horikoshii OT3. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 73, 1068-71	4.2	2
94	Crystallization and preliminary X-ray analysis of the C-terminal RNase III domain of human Dicer. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006 , 62, 402-4		2
93	Conformational phase diagram of poly(l-lysine) in sodium alkanesulfonate aqueous solutions at various temperatures. <i>Colloid and Polymer Science</i> , 2006 , 284, 725-732	2.4	2
92	Backbone (1)H, (13)C and (15)N resonance assignment of the N-terminal domain of human eRF1. Journal of Biomolecular NMR, 2004 , 30, 109-10	3	2
91	Crystallization and preliminary X-ray crystallographic analysis of peptide deformylase from Thermus thermophilus HB8. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 1299	-300	2
90	Crystallization and preliminary X-ray analysis of carboxypeptidase 1 from Thermus thermophilus. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 1445-6		2
89	Kinetic studies of unfolding process of aspergillopepsin II by pH-jump methods. <i>Biochemical and Biophysical Research Communications</i> , 2003 , 301, 745-50	3.4	2
88	Crystal structure of glutamine amidotransferase from Pyrococcus horikoshii OT3. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2005 , 81, 459-462	4	2
87	Substrate specificity of non-pepsin-type acid proteinase, Aspergillus niger proteinase A. <i>Advances in Experimental Medicine and Biology</i> , 1998 , 436, 345-8	3.6	2
86	Crystallization and preliminary crystallographic analysis of major nitroreductase from Escherichia coli. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1999 , 55, 1901-2		2
85	Effects of replacement of Lys25 with Gln on the conformation of ribonuclease T1: sequence-specific 1H NMR resonance assignments of Gln25 ribonuclease T1 by two-dimensional NMR spectroscopy. <i>Journal of Biochemistry</i> , 1995 , 118, 710-6	3.1	2
84	Crystal structure of (S)-3-O-geranylgeranylglyceryl phosphate synthase from Thermoplasma acidophilum in complex with the substrate sn-glycerol 1-phosphate. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2019 , 75, 470-479	1.1	2

83	Crystal structure and substrate recognition mechanism of the prolyl endoprotease PEP from Aspergillus niger <i>Biochemical and Biophysical Research Communications</i> , 2022 , 591, 76-81	3.4	2
82	Discovery and Characterization of a Thermostable NADH Oxidase from Pyrococcus horikoshii OT3. <i>Bulletin of the Korean Chemical Society</i> , 2009 , 30, 2984-2988	1.2	2
81	Structural and functional analysis of troponins from scallop striated and human cardiac muscles. <i>Advances in Experimental Medicine and Biology</i> , 2007 , 592, 163-73	3.6	2
80	Molecular Basis for Substrate Recognition and Catalysis by a Marine Bacterial Laminarinase. <i>Applied and Environmental Microbiology</i> , 2020 , 86,	4.8	2
79	Evaluation of spice and herb as phyto-derived selective modulators of human retinaldehyde dehydrogenases using a simple in vitro method. <i>Bioscience Reports</i> , 2021 , 41,	4.1	2
78	A class-A GPCR solubilized under high hydrostatic pressure retains its ligand binding ability. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2016 , 1858, 2145-2151	3.8	2
77	Structural comparisons of phosphoenolpyruvate carboxykinases reveal the evolutionary trajectories of these phosphodiester energy conversion enzymes. <i>Journal of Biological Chemistry</i> , 2019 , 294, 19269-19278	5.4	2
76	Distortion of double-stranded DNA structure by the binding of the restriction DNA glycosylase R.Pabl. <i>Nucleic Acids Research</i> , 2020 , 48, 5106-5118	20.1	2
75	NMR-Based Metabolomics of Foods379-387		2
74	X-ray crystallographic study of a non-pepsin-type acid proteinase, Aspergillus niger proteinase A. <i>Advances in Experimental Medicine and Biology</i> , 1995 , 362, 605-9	3.6	2
74		3.6 1.7	1
	Advances in Experimental Medicine and Biology, 1995, 362, 605-9 Analysis of Weak Affinity of D-Fructofuranosyl-(2<-j1)-2-acetamido-2-deoxy-D-glucopyranoside for Yeast D-Fructofuranosidase Using NMR Spectroscopy. Journal of Carbohydrate Chemistry, 2014,		
73	Advances in Experimental Medicine and Biology, 1995, 362, 605-9 Analysis of Weak Affinity of I-D-Fructofuranosyl-(2<-j1)-2-acetamido-2-deoxy-I-D-glucopyranoside for Yeast I-Fructofuranosidase Using NMR Spectroscopy. Journal of Carbohydrate Chemistry, 2014, 33, 498-505 Crystallization and preliminary X-ray diffraction analysis of a novel type of phosphoserine phosphatase from Hydrogenobacter thermophilus TK-6. Acta Crystallographica Section F: Structural		1
73 72	Advances in Experimental Medicine and Biology, 1995, 362, 605-9 Analysis of Weak Affinity of D-Fructofuranosyl-(2<-j1)-2-acetamido-2-deoxy-D-glucopyranoside for Yeast D-Fructofuranosidase Using NMR Spectroscopy. Journal of Carbohydrate Chemistry, 2014, 33, 498-505 Crystallization and preliminary X-ray diffraction analysis of a novel type of phosphoserine phosphatase from Hydrogenobacter thermophilus TK-6. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 911-3 Crystallization and preliminary X-ray crystallographic analysis of dioscorin from Dioscorea japonica.		1
73 72 71	Advances in Experimental Medicine and Biology, 1995, 362, 605-9 Analysis of Weak Affinity of ID-Fructofuranosyl-(2<-j1)-2-acetamido-2-deoxy-ID-glucopyranoside for Yeast II-Fructofuranosidase Using NMR Spectroscopy. Journal of Carbohydrate Chemistry, 2014, 33, 498-505 Crystallization and preliminary X-ray diffraction analysis of a novel type of phosphoserine phosphatase from Hydrogenobacter thermophilus TK-6. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 911-3 Crystallization and preliminary X-ray crystallographic analysis of dioscorin from Dioscorea japonica. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 193-5 Purification, crystallization and preliminary X-ray analysis of glucokinase from Streptomyces griseus in complex with glucose. Acta Crystallographica Section F: Structural Biology Communications, 2011,		1 1
73 72 71 70	Advances in Experimental Medicine and Biology, 1995, 362, 605-9 Analysis of Weak Affinity of I-D-Fructofuranosyl-(2<-j1)-2-acetamido-2-deoxy-ID-glucopyranoside for Yeast I-Fructofuranosidase Using NMR Spectroscopy. Journal of Carbohydrate Chemistry, 2014, 33, 498-505 Crystallization and preliminary X-ray diffraction analysis of a novel type of phosphoserine phosphatase from Hydrogenobacter thermophilus TK-6. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 911-3 Crystallization and preliminary X-ray crystallographic analysis of dioscorin from Dioscorea japonica. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 193-5 Purification, crystallization and preliminary X-ray analysis of glucokinase from Streptomyces griseus in complex with glucose. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 914-6 Crystal structure of the DUF54 family protein PH1010 from hyperthermophilic archaea Pyrococcus	1.7	1 1 1
73 72 71 70 69	Analysis of Weak Affinity of I-D-Fructofuranosyl-(2<-j1)-2-acetamido-2-deoxy-ID-glucopyranoside for Yeast II-Fructofuranosidase Using NMR Spectroscopy. <i>Journal of Carbohydrate Chemistry</i> , 2014 , 33, 498-505 Crystallization and preliminary X-ray diffraction analysis of a novel type of phosphoserine phosphatase from Hydrogenobacter thermophilus TK-6. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 911-3 Crystallization and preliminary X-ray crystallographic analysis of dioscorin from Dioscorea japonica. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 193-5 Purification, crystallization and preliminary X-ray analysis of glucokinase from Streptomyces griseus in complex with glucose. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2011 , 67, 914-6 Crystal structure of the DUF54 family protein PH1010 from hyperthermophilic archaea Pyrococcus horikoshii OT3. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009 , 74, 256-60 Crystallization and Preliminary Crystallographic Studies of Dimethyl Sulfoxide Reductase from Rhodobacter sphaeroides f. sp. denitrificans <i>Proceedings of the Japan Academy Series B: Physical</i>	4.2	1 1 1 1 1

65	1H, 13C, 15N resonance assignments of the cytokine LECT2. <i>Journal of Biomolecular NMR</i> , 2004 , 29, 543	- 4	1
64	Assignment of 1H, 13C and 15N resonances of FKBP. from Methanococcus thermolithotrophicus. <i>Journal of Biomolecular NMR</i> , 2000 , 17, 183-4	3	1
63	Crystallization and preliminary X-ray diffraction studies of a rice cysteine proteinase inhibitor, Oryzacystatin-I. <i>Journal of Biochemistry</i> , 1998 , 123, 568-70	3.1	1
62	Purification and characterization of the parvalbumin from monkey skeletal muscle. <i>Comparative Biochemistry and Physiology Part B: Comparative Biochemistry</i> , 1990 , 96, 665-9		1
61	Bromein, a Bromelain Inhibitor from Pineapple Stem: Structural and Functional Characteristics. <i>Protein and Peptide Letters</i> , 2018 , 25, 838-852	1.9	1
60	Recent Progress and Development of Crystal Structure Analysis of Enzymes and Other Proteins. Nihon Kessho Gakkaishi, 2010 , 52, 8-13	Ο	1
59	Age-Dependent Decrease in the Induction of Regulatory T Cells Is Associated With Decreased Expression of RALDH2 in Mesenteric Lymph Node Dendritic Cells. <i>Frontiers in Immunology</i> , 2020 , 11, 1555	8.4	1
58	☐ Elemene Suppresses Obesity-Induced Imbalance in the Microbiota-Gut-Brain Axis. <i>Biomedicines</i> , 2021 , 9,	4.8	1
57	Coordination to Divalent Cations by Calcium-Binding Proteins. <i>Methods in Molecular Biology</i> , 2019 , 1929, 127-134	1.4	1
56	Structural basis of different substrate preferences of two old yellow enzymes from yeasts in the asymmetric reduction of enone compounds. <i>Bioscience, Biotechnology and Biochemistry</i> , 2019 , 83, 456-4	62 ¹	1
55	Crystallization and melting properties studied by DSC and FTIR spectroscopy of goldenberry (Physalis peruviana) oil. <i>Food Chemistry</i> , 2022 , 366, 130645	8.5	1
54	Comparison of explosion puffing drying with other methods on the physicochemical properties and volatiles of yam (Dioscorea opposita thunb.) chips through multivariate analysis. <i>Drying Technology</i> ,1-16	2.6	1
53	Regulation of Th2 responses by Lactococcus lactis subsp. cremoris YRC3780 alleviates DNCB-induced atopic dermatitis in the mouse model. <i>Journal of Functional Foods</i> , 2022 , 94, 105091	5.1	1
52	Txn2 haplodeficiency does not affect cochlear antioxidant defenses or accelerate the progression of cochlear cell loss or hearing loss across the lifespan. <i>Experimental Gerontology</i> , 2020 , 141, 111078	4.5	O
51	Ginkgo biloba Seeds 2020 , 241-254		0
50	Expression, purification, crystallization and preliminary X-ray analysis of 4-hydroxy-3-methyl-2-keto-pentanoate aldolase (asHPAL) from Arthrobacter simplex strain AKU 626. Acta Crystallographica Section F: Structural Biology Communications, 2012 , 68, 958-61		O
49	Crystallization and preliminary X-ray analysis of a putative sensor histidine kinase domain: the C-terminal domain of HksP4 from Aquifex aeolicus VF5. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2011 , 67, 803-7		О
48	Crystallization and preliminary X-ray analysis of 5-keto-D-gluconate reductase from Gluconobacter suboxydans IFO12528 complexed with 5-keto-D-gluconate and NADPH. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010 , 66, 1680-2		O

(2010-1989)

47	Potentiometric studies on calcium and magnesium binding by bullfrog skeletal muscle troponin C. Effect of proton on binding of calcium and magnesium <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1989 , 65, 215-218	4	O
46	Induction of Oral Tolerance by Pepsin-Digested Gliadin Retaining T Cell Reactivity in a Mouse Model of Wheat Allergy. <i>International Archives of Allergy and Immunology</i> , 2020 , 181, 446-455	3.7	O
45	Isolation and characterization of oligopeptides with vascular disease suppression effects derived from wheat gluten. <i>Journal of Food Science and Technology</i> , 2021 , 58, 3504-3513	3.3	O
44	Highlighting the potential utility of MBP crystallization chaperone for Arabidopsis BIL1/BZR1 transcription factor-DNA complex. <i>Scientific Reports</i> , 2021 , 11, 3879	4.9	O
43	Identification of the Effects of Chondroitin Sulfate on Inhibiting CDKs in Colorectal Cancer Based on Bioinformatic Analysis and Experimental Validation. <i>Frontiers in Oncology</i> , 2021 , 11, 705939	5.3	0
42	Preparation of the Extracellular Domain of Recombinant Human Toll-like Receptor 6. <i>Protein Journal</i> , 2017 , 36, 28-35	3.9	
41	Application of Liquid Cell Microscopy to Study Function of Muscle Proteins371-390		
40	2P077 Different Ca2+-sensitivities between the EF-hands of T- and L-plastins(01D. Protein: Function,Poster,The 52nd Annual Meeting of the Biophysical Society of Japan(BSJ2014)). <i>Seibutsu Butsuri</i> , 2014 , 54, S207	Ο	
39	Development of Quasi-Microgravity Environment for Protein Crystallization. <i>Nihon Kessho Gakkaishi</i> , 2017 , 59, 182-187	Ο	
38	Purification, crystallization and preliminary X-ray analysis of SGR6054, a Streptomyces homologue of the mycobacterial integration host factor mIHF. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012 , 68, 1085-8		
37	1P024 Coordination to divalent cations by calcium-binding proteins and calcium-binding peptide analogues studied by FTIR spectroscopy(01B. Protein:Structure & Function,Poster). <i>Seibutsu Butsuri</i> , 2013 , 53, S109	0	
36	???????????ሚ??????????? ABA?????????????	О	
35	Ca2+??????on/off?????????????. Kagaku To Seibutsu, 2009 , 47, 838-844	O	
34	3P-028 X-ray structural analysis of N-terminal domain of KaiC (KaiCl) for understanding of restrained ATPase activity(Protein:Structure & Function,The 47th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2009 , 49, S155	Ο	
33	1P-146 Crystal Structure of Nucleotide-free Kinesin-1 Motor Domain Explains Coordinated Walking Mechanism(Molecular motor, The 47th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2009 , 49, S85	Ο	
32	1TP4-06 Crystal Structure of Nucleotide-free Kinesin-1 Motor Domain Explains Coordinated Walking Mechanism(The 47th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2009 , 49, S36	Ο	
31	1P012 Structural analysis for substrate recognition of carbonyl reductase S1(Protein:Structure,The 48th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2010 , 50, S21	О	
30	2P030 The crystal structure of L-sorbose reductase from Gluconobacter frateurii complexed with NADPH and L-sorbose(The 48th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2010 , 50, S87	О	

29	1P046 1YA1045 Structural basis of abscisic acid signaling(Protein:Structure & Function,Early Research in Biophysics Award Candidate Presentations,Early Research in Biophysics Award,The 48th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2010 , 50, S27	0
28	2P165 Nucleotide-dependent changes in the stability of the ncd coiled-coil stalk domain as monitored by FRET(Molecular motors,Oral Presentations). <i>Seibutsu Butsuri</i> , 2007 , 47, S154	O
27	S06A5 Crystal structure and dsRNA cleavage activity of the RNase III domain of Dicer(Structural Biology of ncRNA Processing and Function). <i>Seibutsu Butsuri</i> , 2007 , 47, S9	O
26	Crystallization and preliminary X-ray analysis of PH1566, a putative ribosomal RNA-processing factor from the hyperthermophilic archaeon Pyrococcus horikoshii OT3. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006 , 62, 47-8	
25	1H, 13C and 15N assignments of the tandem WW domains of human MAGI-1/BAP-1. <i>Journal of Biomolecular NMR</i> , 2004 , 29, 539-40	3
24	Crystal structure of acylphosphatase from hyperthermophilic archaeon Pyrococcus horikoshii OT3. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2004 , 80, 439-442	4
23	Crystal structure of the short-chain flavin reductase HpaC from Sulfolobus tokodaii strain 7. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2005 , 81, 229-232	4
22	Effects of ionization on chemical shifts of the surrounding protons of bromelain inhibitor VI. <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 2005 , 81, 454-458	4
21	JAACT Issue. Preface. <i>Cytotechnology</i> , 2001 , 35, 1	2.2
20	Conformation analysis of non-pepsin-type acid proteinase A from the fungus Aspergillus niger by NMR. <i>Advances in Experimental Medicine and Biology</i> , 1995 , 362, 611-5	3.6
19	Temperature Dependences Of The Infrared And Circular Dichroism Spectra Of Ribonuclease A And Troponin C 1989 , 1145, 269	
18	Potentiometric studies on calcium and magnesium binding by bullfrog skeletal muscle troponin C. Potentiometric titration curves <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1989 , 65, 211-214	4
17	Potentiometric titration curves Proceedings of the Japan Academy Series B: Physical and Biological	4
	Potentiometric titration curves <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1989 , 65, 211-214 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. An improved system for physiological measurements <i>Proceedings of the Japan Academy Series B:</i>	
17	Potentiometric titration curves <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1989 , 65, 211-214 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. An improved system for physiological measurements <i>Proceedings of the Japan Academy Series B: Physical and Biological Sciences</i> , 1988 , 64, 278-281 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. A post-contractile increase in NMR-visible phosphates <i>Proceedings of the Japan Academy Series B:</i>	4
17 16	Potentiometric titration curves Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1989, 65, 211-214 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. An improved system for physiological measurements Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1988, 64, 278-281 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. A post-contractile increase in NMR-visible phosphates Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1988, 64, 282-286 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. Mechanism of post-contractile increase in NMR-visible phosphates Proceedings of the Japan	4
17 16 15	Potentiometric titration curves Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1989, 65, 211-214 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. An improved system for physiological measurements Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1988, 64, 278-281 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. A post-contractile increase in NMR-visible phosphates Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1988, 64, 282-286 Time-resolved 31P-NMR studies on contraction and recovery of bullfrog sartorius muscles. Mechanism of post-contractile increase in NMR-visible phosphates Proceedings of the Japan Academy Series B: Physical and Biological Sciences, 1988, 64, 287-289 Crystallographer Challenge in the 21st Century. 3. Frontier of Protein Crystallography. Frontiers	4 4

LIST OF PUBLICATIONS

11	A phosphorus-31 nuclear magnetic resonance study on the complex of chicken gizzard myosin subfragment 1 with adenosine diphosphate 1999 , 75-78	
10	Chapter 1 Electron Microscopic Visualization and Recording of ATP-Induced Myosin Head Power Stroke Producing Muscle Contraction Using the Gas Environmental Chamber 2016 , 1-34	
9	Chapter 4 Limitations of in vitro Motility Assay Systems in Studying Molecular Mechanism of Muscle Contraction as Revealed by the Effect of Antibodies to Myosin Head 2016 , 117-142	
8	II-1. Structural biology of proteins from aquatic organisms. <i>Nippon Suisan Gakkaishi</i> , 2017 , 83, 819-819	0.2
7	1P190 1B1435 Structural basis for the coordinated processive movement of kinesin-1 studied by structural and single molecule analyses(Molecular motor,Oral Presentations,The 48th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2010 , 50, S52-S53	O
6	Effects of Nutraceutical Antioxidants on Age-Related Hearing Loss 2010 , 113-124	
5	Mechanistic Basis for Plant Responses to Drought Stress: Regulatory Mechanism of Abscisic Acid Signaling. <i>Nihon Kessho Gakkaishi</i> , 2011 , 53, 178-185	O
4	Recent Progress in Abscisic Acid Receptor Research. <i>Seibutsu Butsuri</i> , 2011 , 51, 026-027	O
3	Improved preparation of group-specific component (Gc) protein to derive macrophage activating factor. <i>Protein Expression and Purification</i> , 2020 , 175, 105714	2
2	Crystal structure and DNA cleavage mechanism of the restriction DNA glycosylase R.CcoLI from Campylobacter coli. <i>Scientific Reports</i> , 2021 , 11, 859	4.9
1	Structural Diversity of Transcription Factor Complexes in TGF-\(\Percapprox\) Signaling. <i>Nihon Kessho Gakkaishi</i> , 2022 , 64, 113-114	0