List of Publications by Year in descending order

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	57758	85541
7,134	44	71
citations	h-index	g-index
214	214	9233
docs citations	times ranked	citing authors
	7,134 citations 214 docs citations	7,13444citationsh-index214214docs citationstimes ranked

#	Article	IF	CITATIONS
1	Reduction of Recombinant Adeno-Associated Virus Vector Adsorption on Solid Surfaces by Polyionic Hydrophilic Complex Coating. Journal of Pharmaceutical Sciences, 2022, 111, 663-671.	3.3	6
2	Pro108Ser mutation of SARS-CoV-2 3CLpro reduces the enzyme activity and ameliorates the clinical severity of COVID-19. Scientific Reports, 2022, 12, 1299.	3.3	15
3	Selective targeting of multiple myeloma cells with a monoclonal antibody recognizing the ubiquitous protein CD98 heavy chain. Science Translational Medicine, 2022, 14, eaax7706.	12.4	10
4	The Fab portion of immunoglobulin G has sites in the CL domain that interact with Fc gamma receptor IIIa. MAbs, 2022, 14, 2038531.	5.2	7
5	Critical analysis of techniques and materials used in devices, syringes, and needles used for intravitreal injections. Progress in Retinal and Eye Research, 2021, 80, 100862.	15.5	51
6	Development of syringes and vials for delivery of biologics: current challenges and innovative solutions. Expert Opinion on Drug Delivery, 2021, 18, 459-470.	5.0	19
7	Glycyrrhizin Derivatives Suppress Cancer Chemoresistance by Inhibiting Progesterone Receptor Membrane Component 1. Cancers, 2021, 13, 3265.	3.7	16
8	Plate Reader-Based Analytical Method for the Size Distribution of Submicron-Sized Protein Aggregates Using Three-Dimensional Homodyne Light Detection. Journal of Pharmaceutical Sciences, 2021, 110, 3803-3810.	3.3	4
9	Mechanism of dimerization and structural features of human LI-cadherin. Journal of Biological Chemistry, 2021, 297, 101054.	3.4	4
10	Relation of Colloidal and Conformational Stabilities to Aggregate Formation in a Monoclonal Antibody. Journal of Pharmaceutical Sciences, 2020, 109, 308-315.	3.3	17
11	Assessment of the Injection Performance of a Tapered Needle for Use in Prefilled Biopharmaceutical Products. Journal of Pharmaceutical Sciences, 2020, 109, 515-523.	3.3	22
12	Automatic Identification of the Stress Sources of Protein Aggregates Using Flow Imaging Microscopy Images. Journal of Pharmaceutical Sciences, 2020, 109, 614-623.	3.3	36
13	Intrastrand backbone-nucleobase interactions stabilize unwound right-handed helical structures of heteroduplexes of L-aTNA/RNA and SNA/RNA. Communications Chemistry, 2020, 3, .	4.5	9
14	Current status and issues of protein solution biophysics—Session 1SDP. Biophysical Reviews, 2020, 12, 263-264.	3.2	1
15	Dataset of microbial community structure in alcohol sprayed banana associated with ripening process. Data in Brief, 2020, 29, 105216.	1.0	0
16	Supramolecular tholos-like architecture constituted by archaeal proteins without functional annotation. Scientific Reports, 2020, 10, 1540.	3.3	8
17	Recent Achievements and Current Interests in Research on the Characterization and Quality Control of Biopharmaceuticals in Japan. Journal of Pharmaceutical Sciences, 2020, 109, 1652-1661.	3.3	3
18	Effect of UVC Irradiation on the Oxidation of Histidine in Monoclonal Antibodies. Scientific Reports, 2020, 10, 6333.	3.3	20

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19	Allosteric regulation accompanied by oligomeric state changes of Trypanosoma brucei GMP reductase through cystathionine-β-synthase domain. Nature Communications, 2020, 11, 1837.	12.8	10
20	Incorporation of Pseudoâ€complementary Bases 2,6â€Diaminopurine and 2â€Thiouracil into Serinol Nucleic Acid (SNA) to Promote SNA/RNA Hybridization. Chemistry - an Asian Journal, 2020, 15, 1266-1271.	3.3	10
21	The Fab portion of immunoglobulin G contributes to its binding to FcÎ ³ receptor III. Scientific Reports, 2019, 9, 11957.	3.3	35
22	Structural characterization of HypX responsible for CO biosynthesis in the maturation of NiFe-hydrogenase. Communications Biology, 2019, 2, 385.	4.4	13
23	A head-to-toe dimerization has physiological relevance for ligand-induced inactivation of protein tyrosine receptor type Z. Journal of Biological Chemistry, 2019, 294, 14953-14965.	3.4	12
24	Interlaboratory comparison about feasibility of insoluble particulate matter test for injections with reduced test volume in light obscuration method. Biologicals, 2019, 57, 46-49.	1.4	8
25	Mutational and Combinatorial Control of Self-Assembling and Disassembling of Human Proteasome α Subunits. International Journal of Molecular Sciences, 2019, 20, 2308.	4.1	6
26	lonic liquids and protein folding—old tricks for new solvents. Biophysical Reviews, 2019, 11, 209-225.	3.2	19
27	SDS-induced oligomerization of Lys49-phospholipase A2 from snake venom. Scientific Reports, 2019, 9, 2330.	3.3	15
28	An Assessment of the Ability of Submicron- and Micron-Size Silicone Oil Droplets in Dropped Prefillable Syringes to Invoke Early- and Late-Stage Immune Responses. Journal of Pharmaceutical Sciences, 2019, 108, 2278-2287.	3.3	47
29	Identification of IgG1 Aggregation Initiation Region by Hydrogen Deuterium Mass Spectrometry. Journal of Pharmaceutical Sciences, 2019, 108, 2323-2333.	3.3	14
30	Temperature-controlled repeatable scrambling and induced-sorting of building blocks between cubic assemblies. Nature Communications, 2019, 10, 1440.	12.8	11
31	Crystal structure of the dog allergen Can f 6 and structure-based implications of its cross-reactivity with the cat allergen Fel d 4. Scientific Reports, 2019, 9, 1503.	3.3	15
32	Cooperative Binding of KaiB to the KaiC Hexamer Ensures Accurate Circadian Clock Oscillation in Cyanobacteria. International Journal of Molecular Sciences, 2019, 20, 4550.	4.1	18
33	Polarizability and isotope effects on dispersion interactions in water. Communications Chemistry, 2019, 2, .	4.5	4
34	Collaborative Study for Analysis of Subvisible Particles Using Flow Imaging and Light Obscuration: Experiences in Japanese Biopharmaceutical Consortium. Journal of Pharmaceutical Sciences, 2019, 108, 832-841.	3.3	40
35	Quantitative Laser Diffraction for Quantification of Protein Aggregates: Comparison With Resonant Mass Measurement, Nanoparticle Tracking Analysis, Flow Imaging, and Light Obscuration. Journal of Pharmaceutical Sciences, 2019, 108, 755-762.	3.3	25
36	ATP hydrolysis by KaiC promotes its KaiA binding in the cyanobacterial circadian clock system. Life Science Alliance, 2019, 2, e201900368.	2.8	14

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37	Inhalation effect of massoilactone from massoia essential oil on lipid profile, liver tissues, and body weight of Sprague Dawley rat. Journal of Applied Pharmaceutical Science, 2019, 9, 111-116.	1.0	1
38	Sweeping of Adsorbed Therapeutic Protein on Prefillable Syringes Promotes Micron Aggregate Generation. Journal of Pharmaceutical Sciences, 2018, 107, 1521-1529.	3.3	30
39	Sedimentation velocity analytical ultracentrifugation for characterization of therapeutic antibodies. Biophysical Reviews, 2018, 10, 259-269.	3.2	26
40	Insight into adaptive remodeling of the rotor ring complex of the bacterial flagellar motor. Biochemical and Biophysical Research Communications, 2018, 496, 12-17.	2.1	17
41	Hyperthermostable cube-shaped assembly in water. Communications Chemistry, 2018, 1, .	4.5	22
42	RecA requires two molecules of Mg2+ ions for its optimal strand exchange activity in vitro. Nucleic Acids Research, 2018, 46, 2548-2559.	14.5	12
43	Physicochemical improvement of rabbit derived single-domain antibodies by substitutions with amino acids conserved in camelid antibodies. Journal of Bioscience and Bioengineering, 2018, 125, 654-661.	2.2	1
44	Native mass spectrometry for understanding dynamic protein complex. Biochimica Et Biophysica Acta - General Subjects, 2018, 1862, 275-286.	2.4	20
45	Assembly of protein complexes restricts diffusion of Wnt3a proteins. Communications Biology, 2018, 1, 165.	4.4	23
46	Structural Dynamics of the PET-Degrading Cutinase-like Enzyme from <i>Saccharomonospora viridis</i> AHK190 in Substrate-Bound States Elucidates the Ca ²⁺ -Driven Catalytic Cycle. Biochemistry, 2018, 57, 5289-5300.	2.5	59
47	Dynamic structural states of ClpB involved in its disaggregation function. Nature Communications, 2018, 9, 2147.	12.8	55
48	Bifacial Nucleobases for Hexaplex Formation in Aqueous Solution. Journal of the American Chemical Society, 2018, 140, 8456-8462.	13.7	21
49	Recent Developments of Cyclo Olefin Polymers. Kobunshi Ronbunshu, 2018, 75, 477-485.	0.2	2
50	Analytical ultracentrifugation with fluorescence detection system reveals differences in complex formation between recombinant human TNF and different biological TNF antagonists in various environments. MAbs, 2017, 9, 664-679.	5.2	27
51	Friability Testing as a New Stress-Stability Assay for Biopharmaceuticals. Journal of Pharmaceutical Sciences, 2017, 106, 2966-2978.	3.3	27
52	A Comprehensive Study of the Interaction between Peptidoglycan Fragments and the Extracellular Domain of <i>Mycobacterium tuberculosis</i> Ser/Thr Kinase PknB. ChemBioChem, 2017, 18, 2094-2098.	2.6	12
53	Efficient generation of single domain antibodies with high affinities and enhanced thermal stabilities. Scientific Reports, 2017, 7, 5794.	3.3	8
54	Calcium depletion destabilises kinetochore fibres by the removal of CENP-F from the kinetochore. Scientific Reports, 2017, 7, 7335.	3.3	5

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55	Recent Topics of Research in the Characterization and Quality Control of Biopharmaceuticals in Japan. Journal of Pharmaceutical Sciences, 2017, 106, 3431-3437.	3.3	7
56	Synergistic Effect of Cavitation and Agitation on Protein Aggregation. Journal of Pharmaceutical Sciences, 2017, 106, 521-529.	3.3	59
57	NMR Detection of Semi-Specific Antibody Interactions in Serum Environments. Molecules, 2017, 22, 1619.	3.8	13
58	Interdependency and phosphorylation of KIF4 and condensin I are essential for organization of chromosome scaffold. PLoS ONE, 2017, 12, e0183298.	2.5	11
59	Calcium ions function as a booster of chromosome condensation. Scientific Reports, 2016, 6, 38281.	3.3	39
60	Structural and binding properties of laminarin revealed by analytical ultracentrifugation and calorimetric analyses. Carbohydrate Research, 2016, 431, 33-38.	2.3	11
61	Suppression of Methionine Oxidation of a Pharmaceutical Antibody Stored in a Polymer-Based Syringe. Journal of Pharmaceutical Sciences, 2016, 105, 623-629.	3.3	19
62	Structural Analysis Reveals that Toll-like Receptor 7 Is a Dual Receptor for Guanosine and Single-Stranded RNA. Immunity, 2016, 45, 737-748.	14.3	321
63	Interaction mode between catalytic and regulatory subunits in glucosidase II involved in ER glycoprotein quality control. Protein Science, 2016, 25, 2095-2101.	7.6	16
64	Mass Spectrometric Characterization of HIV-1 Reverse Transcriptase Interactions with Non-nucleoside Reverse Transcriptase Inhibitors. Biological and Pharmaceutical Bulletin, 2016, 39, 450-454.	1.4	5
65	Taste substance binding elicits conformational change of taste receptor T1r heterodimer extracellular domains. Scientific Reports, 2016, 6, 25745.	3.3	36
66	New insight into the dynamical system of αB-crystallin oligomers. Scientific Reports, 2016, 6, 29208.	3.3	32
67	Structural characterization of the circadian clock protein complex composed of KaiB and KaiC by inverse contrast-matching small-angle neutron scattering. Scientific Reports, 2016, 6, 35567.	3.3	24
68	Haem-dependent dimerization of PGRMC1/Sigma-2 receptor facilitates cancer proliferation and chemoresistance. Nature Communications, 2016, 7, 11030.	12.8	153
69	Small-molecule inhibition of PTPRZ reduces tumor growth in a rat model of glioblastoma. Scientific Reports, 2016, 6, 20473.	3.3	47
70	Mass spectrometric analysis of protein–ligand interactions. Biophysics and Physicobiology, 2016, 13, 87-95.	1.0	33
71	Novel helical assembly in arginine methyltransferase 8. Journal of Molecular Biology, 2016, 428, 1197-1208.	4.2	19
72	Structure of IZUMO1–JUNO reveals sperm–oocyte recognition during mammalian fertilization. Nature, 2016, 534, 566-569.	27.8	118

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73	Two Decades of Publishing Excellence in Pharmaceutical Biotechnology. Journal of Pharmaceutical Sciences, 2015, 104, 290-300.	3.3	15
74	Disassembly of the self-assembled, double-ring structure of proteasome α7 homo-tetradecamer by α6. Scientific Reports, 2015, 5, 18167.	3.3	23
75	Condensin in Chromatid Cohesion and Segregation. Cytogenetic and Genome Research, 2015, 147, 212-216.	1.1	6
76	Chromosome Scaffold is a Double-Stranded Assembly of Scaffold Proteins. Scientific Reports, 2015, 5, 11916.	3.3	37
77	Quantitative Laser Diffraction Method for the Assessment of Protein Subvisible Particles. Journal of Pharmaceutical Sciences, 2015, 104, 618-626.	3.3	23
78	Target Antigen Density Governs the Efficacy of Anti–CD20-CD28-CD3 ζ Chimeric Antigen Receptor–Modified Effector CD8+ T Cells. Journal of Immunology, 2015, 194, 911-920.	0.8	228
79	Structural basis of CpG and inhibitory DNA recognition by Toll-like receptor 9. Nature, 2015, 520, 702-705.	27.8	290
80	Structural basis for PPARÎ ³ transactivation by endocrine-disrupting organotin compounds. Scientific Reports, 2015, 5, 8520.	3.3	56
81	Crystal structure of extracellular domain of human lectinâ€like transcript 1 (LLT1), the ligand for natural killer receptorâ€₽1A. European Journal of Immunology, 2015, 45, 1605-1613.	2.9	16
82	Chemical mechanism of petal color development of Nemophila menziesii by a metalloanthocyanin, nemophilin. Tetrahedron, 2015, 71, 9123-9130.	1.9	15
83	Structural Basis for Dimer Formation of Human Condensin Structural Maintenance of Chromosome Proteins and Its Implications for Single-stranded DNA Recognition. Journal of Biological Chemistry, 2015, 290, 29461-29477.	3.4	18
84	Pepsin immobilization on an aldehyde-modified polymethacrylate monolith and its application for protein analysis. Journal of Bioscience and Bioengineering, 2015, 119, 505-510.	2.2	17
85	Effects of Syringe Material and Silicone Oil Lubrication on the Stability of Pharmaceutical Proteins. Journal of Pharmaceutical Sciences, 2015, 104, 527-535.	3.3	125
86	A Multilaboratory Comparison of Calibration Accuracy and the Performance of External References in Analytical Ultracentrifugation. PLoS ONE, 2015, 10, e0126420.	2.5	71
87	pH-Dependent Assembly and Segregation of the Coiled-Coil Segments of Yeast Putative Cargo Receptors Emp46p and Emp47p. PLoS ONE, 2015, 10, e0140287.	2.5	7
88	Detection of Histidine Oxidation in a Monoclonal Immunoglobulin Gamma (IgG) 1 Antibody. Analytical Chemistry, 2014, 86, 7536-7543.	6.5	41
89	Liquid formulation for antibody drugs. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 2041-2052.	2.3	81
90	The Effect of Magnesium Ions on Chromosome Structure as Observed by Helium Ion Microscopy. Microscopy and Microanalysis, 2014, 20, 184-188.	0.4	10

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91	Chromosome Interior Observation by Focused Ion Beam/Scanning Electron Microscopy (FIB/SEM) Using Ionic Liquid Technique. Microscopy and Microanalysis, 2014, 20, 1340-1347.	0.4	21
92	SuperNova, a monomeric photosensitizing fluorescent protein for chromophore-assisted light inactivation. Scientific Reports, 2013, 3, 2629.	3.3	132
93	Aggregation analysis of pharmaceutical human immunoglobulin preparations using size-exclusion chromatography and analytical ultracentrifugation sedimentation velocity. Journal of Bioscience and Bioengineering, 2013, 115, 104-110.	2.2	29
94	Effects of Ionic Strength and Sugars on the Aggregation Propensity of Monoclonal Antibodies: Influence of Colloidal and Conformational Stabilities. Pharmaceutical Research, 2013, 30, 1263-1280.	3.5	88
95	Characterization of the novel <i>Trypanosoma brucei</i> inosine 5′-monophosphate dehydrogenase. Parasitology, 2013, 140, 735-745.	1.5	11
96	Visualization of Oil Body Distribution inJatropha curcasL. by Four-Wave Mixing Microscopy. Japanese Journal of Applied Physics, 2013, 52, 062403.	1.5	0
97	ASURA (PHB2) Interacts with Scc1 through Chromatin. Cytogenetic and Genome Research, 2013, 139, 225-233.	1.1	2
98	Thermodynamic assessment of domainâ€domain interactions and in vitro activities of mesophilic and thermophilic ribosome recycling factors. Biopolymers, 2013, 100, 366-379.	2.4	3
99	An Archaeal Homolog of Proteasome Assembly Factor Functions as a Proteasome Activator. PLoS ONE, 2013, 8, e60294.	2.5	19
100	Archaeal ribosomal stalk protein interacts with translation factors in a nucleotide-independent manner via its conserved C terminus. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 3748-3753.	7.1	48
101	Creation of a Binuclear Purple Copper Site within a <i>de Novo</i> Coiled-Coil Protein. Biochemistry, 2012, 51, 7901-7907.	2.5	32
102	A nonâ€canonical UBA–UBL interaction forms the linearâ€ubiquitinâ€chain assembly complex. EMBO Reports, 2012, 13, 462-468.	4.5	52
103	RBMX: A Regulator for Maintenance and Centromeric Protection of Sister Chromatid Cohesion. Cell Reports, 2012, 1, 299-308.	6.4	75
104	Effects of rotational speed on the hydrodynamic properties of pharmaceutical antibodies measured by analytical ultracentrifugation sedimentation velocity. European Journal of Pharmaceutical Sciences, 2012, 47, 367-374.	4.0	6
105	Protein encapsulation within synthetic molecular hosts. Nature Communications, 2012, 3, 1093.	12.8	208
106	Domain 5 of high molecular weight kininogen inhibits collagen-mediated cancer cell adhesion and invasion in association with α-actinin-4. Biochemical and Biophysical Research Communications, 2012, 427, 497-502.	2.1	6
107	Chromosome observation by scanning electron microscopy using ionic liquid. Microscopy Research and Technique, 2012, 75, 1113-1118.	2.2	41
108	The triple helical structure and stability of collagen model peptide with 4(<i>s</i>)â€hydroxyprolylâ€proâ€gly units. Biopolymers, 2012, 98, 111-121.	2.4	18

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109	Polymorphism of Collagen Triple Helix Revealed by ¹⁹ F NMR of Model Peptide [Pro-4(<i>R</i>)-Hydroxyprolyl-Gly] ₃ -[Pro-4(<i>R</i>)-Fluoroprolyl-Gly]-[Pro-4(<i>R</i>)-Hydroxypro Journal of Physical Chemistry B, 2012, 116, 6908-6915.	olyl £Ga y]≺sı	ub <i>9</i> 3
110	Drug delivery system for poorly water-soluble compounds using lipocalin-type prostaglandin D synthase. Journal of Controlled Release, 2012, 159, 143-150.	9.9	23
111	Behavior of Monoclonal Antibodies: Relation Between the Second Virial Coefficient (B 2) at Low Concentrations and Aggregation Propensity and Viscosity at High Concentrations. Pharmaceutical Research, 2012, 29, 397-410.	3.5	131
112	Specific Racemization of Heavy-Chain Cysteine-220 in the Hinge Region of Immunoglobulin Gamma 1 as a Possible Cause of Degradation during Storage. Analytical Chemistry, 2011, 83, 3857-3864.	6.5	32
113	Assembly states of the nucleosome assembly protein 1 (NAP-1) revealed by sedimentation velocity and non-denaturing MS. Biochemical Journal, 2011, 436, 101-112.	3.7	21
114	ASURA (PHB2) Is Required for Kinetochore Assembly and Subsequent Chromosome Congression. Acta Histochemica Et Cytochemica, 2011, 44, 247-258.	1.6	7
115	Fc domain mediated self-association of an IgG1 monoclonal antibody under a low ionic strength condition. Journal of Bioscience and Bioengineering, 2011, 112, 326-332.	2.2	46
116	The nuclear scaffold protein SAF-A is required for kinetochore–microtubule attachment and contributes to the targeting of Aurora-A to mitotic spindles. Journal of Cell Science, 2011, 124, 394-404.	2.0	26
117	Thermal effects of added propanol on the helix–coil transition of (Pro-Pro-Gly)10 in D2O solution: An NMR study. Chemical Physics Letters, 2010, 491, 208-213.	2.6	1
118	Phase Separation of an IgG1 Antibody Solution under a Low Ionic Strength Condition. Pharmaceutical Research, 2010, 27, 1348-1360.	3.5	115
119	Characterization of HIV-1 resistance to a fusion inhibitor, N36, derived from the gp41 amino-terminal heptad repeat. Antiviral Research, 2010, 87, 179-186.	4.1	17
120	Cloning, expression, crystallization and preliminary X-ray crystallographic analysis of a human condensin SMC2 hinge domain with short coiled coils. Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 1067-1070.	0.7	2
121	Development of novel humanized anti D20 antibodies based on affinity constant and epitope. Cancer Science, 2010, 101, 201-209.	3.9	39
122	Structural basis for semaphorin signalling through the plexin receptor. Nature, 2010, 467, 1123-1127.	27.8	144
123	Structural basis for the cooperative interplay between the two causative gene products of combined factor V and factor VIII deficiency. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 4034-4039.	7.1	46
124	The Middle Region of an HP1-binding Protein, HP1-BP74, Associates with Linker DNA at the Entry/Exit Site of Nucleosomal DNA. Journal of Biological Chemistry, 2010, 285, 6498-6507.	3.4	21
125	Comparative Analysis of Highly HomologousShewanellaCytochromesc5for Stability and Function. Bioscience, Biotechnology and Biochemistry, 2010, 74, 1079-1083.	1.3	12
126	Assembly Modulation by Adjusting Countercharges of Heterobimetallic Supramolecular Polymers Composed of Tris(spiroborate) Twin Bowls. Journal of the American Chemical Society, 2010, 132, 15556-15558.	13.7	27

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127	Creation of a Type 1 Blue Copper Site within a de Novo Coiled-Coil Protein Scaffold. Journal of the American Chemical Society, 2010, 132, 18191-18198.	13.7	33
128	A nucleolar protein RRS1 contributes to chromosome congression. FEBS Letters, 2009, 583, 1951-1956.	2.8	35
129	The effect of the side chain length of Asp and Glu on coordination structure of Cu ²⁺ in a <i>de novo</i> designed protein. Biopolymers, 2009, 91, 907-916.	2.4	14
130	Stability enhancement of cytochrome c through heme deprotonation and mutations. Biophysical Chemistry, 2009, 139, 37-41.	2.8	1
131	Bioactive beads-mediated transformation of rice with large DNA fragments containing Aegilops tauschii genes. Plant Cell Reports, 2009, 28, 759-768.	5.6	26
132	Hyperstability and crystal structure of cytochrome <i>c</i> ₅₅₅ from hyperthermophilic <i>Aquifex aeolicus</i> . Acta Crystallographica Section D: Biological Crystallography, 2009, 65, 804-813.	2.5	22
133	Effects of antibody affinity and antigen valence on molecular forms of immune complexes. Molecular Immunology, 2009, 47, 357-364.	2.2	34
134	H2A.Z and H3.3 Histone Variants Affect Nucleosome Structure: Biochemical and Biophysical Studies. Biochemistry, 2009, 48, 10852-10857.	2.5	87
135	2DE for Proteome Analysis of Human Metaphase Chromosomes. Methods in Molecular Biology, 2009, 519, 259-271.	0.9	1
136	Thermal unfolding mechanism of lipocalinâ€ŧype prostaglandin D synthase. FEBS Journal, 2008, 275, 233-241.	4.7	9
137	Nucleophosmin is required for chromosome congression, proper mitotic spindle formation, and kinetochoreâ€microtubule attachment in HeLa cells. FEBS Letters, 2008, 582, 3839-3844.	2.8	46
138	The Arabidopsis SDG4 contributes to the regulation of pollen tube growth by methylation of histone H3 lysines 4 and 36 in mature pollen. Developmental Biology, 2008, 315, 355-368.	2.0	109
139	Stabilization Mechanism of Cytochromec552from a Moderately Thermophilic Bacterium,Hydrogenophilus thermoluteolus. Bioscience, Biotechnology and Biochemistry, 2008, 72, 2103-2109.	1.3	7
140	Live Cell Imaging Reveals Plant Aurora Kinase Has Dual Roles During Mitosis. Plant and Cell Physiology, 2008, 49, 1256-1261.	3.1	31
141	Depletion of nucleophosmin leads to distortion of nucleolar and nuclear structures in HeLa cells. Biochemical Journal, 2008, 415, 345-351.	3.7	88
142	Anti-Peptide Antibodies for Examining the Conformation, Molecular Assembly and Localization of an Intracellular Protein, Ribosomal Protein S6, In vivo. Journal of Biochemistry, 2007, 143, 325-332.	1.7	2
143	Histone H2A mobility is regulated by its tails and acetylation of core histone tails. Biochemical and Biophysical Research Communications, 2007, 357, 627-632.	2.1	30
144	Fibrillarin, a nucleolar protein, is required for normal nuclear morphology and cellular growth in HeLa cells. Biochemical and Biophysical Research Communications, 2007, 360, 320-326.	2.1	55

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145	H1.X with different properties from other linker histones is required for mitotic progression. FEBS Letters, 2007, 581, 3783-3788.	2.8	36
146	Nucleolin functions in nucleolus formation and chromosome congression. Journal of Cell Science, 2007, 120, 2091-2105.	2.0	112
147	Development of a multistage classifier for a monitoring system of cell activity based on imaging of chromosomal dynamics. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2007, 71A, 286-296.	1.5	18
148	Chromosome protein framework from proteome analysis of isolated human metaphase chromosomes. Chemical Record, 2007, 7, 230-237.	5.8	34
149	Crystal structure of Pyrococcus horikoshii PPC protein at 1.60 Ã resolution. Proteins: Structure, Function and Bioinformatics, 2007, 67, 505-507.	2.6	11
150	A comparative proteome analysis of human metaphase chromosomes isolated from two different cell lines reveals a set of conserved chromosomeâ€associated proteins. Genes To Cells, 2007, 12, 269-284.	1.2	52
151	Calciteâ€specific coupling protein in barnacle underwater cement. FEBS Journal, 2007, 274, 6436-6446.	4.7	71
152	PHB2 Protects Sister-Chromatid Cohesion in Mitosis. Current Biology, 2007, 17, 1356-1361.	3.9	44
153	Roles of a short connecting disulfide bond in the stability and function of psychrophilic Shewanella violacea cytochrome c 5*. Extremophiles, 2007, 11, 797-807.	2.3	25
154	Structure of Cytochromec552 from a Moderate Thermophilic Bacterium,Hydrogenophilus thermoluteolus: Comparative Study on the Thermostability of Cytochromec‡. Biochemistry, 2006, 45, 6115-6123.	2.5	22
155	Crystallization and preliminary X-ray analysis of the complex of NADH and 3α-hydroxysteroid dehydrogenase fromPseudomonassp. B-0831. Acta Crystallographica Section F: Structural Biology Communications, 2006, 62, 569-571.	0.7	7
156	Regional and segmental flexibility of antibodies in interaction with antigens of different size. FEBS Journal, 2006, 273, 1476-1487.	4.7	42
157	Aurora kinase is required for chromosome segregation in tobacco BY-2 cells. Plant Journal, 2006, 48, 572-580.	5.7	72
158	Characterization of a Splicing Variant of Plant Aurora Kinase. Plant and Cell Physiology, 2006, 48, 369-374.	3.1	11
159	Apo- and Holo-structures of 3α-Hydroxysteroid Dehydrogenase fromPseudomonassp. B-0831. Journal of Biological Chemistry, 2006, 281, 31876-31884.	3.4	16
160	Calreticulin as a new histone binding protein in mitotic chromosomes. Cytogenetic and Genome Research, 2006, 115, 10-15.	1.1	14
161	Fluorescent labeling of plant chromosomes in suspension by FISH. Genes and Genetic Systems, 2005, 80, 35-39.	0.7	16
162	Characterization and dynamic analysis of Arabidopsis condensin subunits, AtCAP-H and AtCAP-H2. Planta, 2005, 222, 293-300.	3.2	24

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163	Characterization of plant Aurora kinases during mitosis. Plant Molecular Biology, 2005, 58, 1-13.	3.9	100
164	Cloning, expression, crystallization and preliminary X-ray characterization of cytochromec552from a moderate thermophilic bacterium,Hydrogenophilus thermoluteolus. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 395-398.	0.7	0
165	Crystallization and preliminary X-ray crystallographic analysis of a conserved domain in plants and prokaryotes fromPyrococcus horikoshiiOT3. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 414-416.	0.7	6
166	Collagen-like triple helix formation of synthetic (Pro-Pro-Gly)10 analogues: (4(S)-hydroxyprolyl-4(R)-hydroxyprolyl-Gly)10, (4(R)-hydroxyprolyl-4(R)-hydroxyprolyl-Gly)10 and (4(S)-fluoroprolyl-4(R)-fluoroprolyl-Gly)10. Journal of Peptide Science, 2005, 11, 609-616.	1.4	31
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