## Kyeong Kyu Kim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8385613/publications.pdf

Version: 2024-02-01

136950 155660 4,176 165 32 55 citations h-index g-index papers 171 171 171 5302 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Analysis of Novel Drug-Resistant Human Cytomegalovirus DNA Polymerase Mutations Reveals the Role of a DNA-Binding Loop in Phosphonoformic Acid Resistance. Frontiers in Microbiology, 2022, 13, 771978.	3.5	7
2	Dual functional roles of a novel bifunctional $\hat{l}^2$ -lactamase/esterase from Lactococcus garvieae. International Journal of Biological Macromolecules, 2022, 206, 203-212.	7.5	3
3	Interaction of Prion Peptides with DNA Structures. ACS Omega, 2022, 7, 176-186.	3.5	1
4	Sequence Analysis and Preliminary X-ray Crystallographic Analysis of an Acetylesterase (LgEstl) from Lactococcus garvieae. Crystals, 2022, 12, 46.	2.2	3
5	Identification, Characterization, and Preliminary X-ray Diffraction Analysis of a Novel Esterase (ScEst) from Staphylococcus chromogenes. Crystals, 2022, 12, 546.	2.2	1
6	Functional and Structural Changes in the Membrane-Bound O-Acyltransferase Family Member 7 (MBOAT7) Protein: The Pathomechanism of a Novel MBOAT7 Variant in Patients With Intellectual Disability. Frontiers in Neurology, 2022, 13, 836954.	2.4	7
7	Generation of a Pure Culture of Neuron-like Cells with a Glutamatergic Phenotype from Mouse Astrocytes. Biomedicines, 2022, 10, 928.	3.2	5
8	Targeted epigenetic modulation using a DNAâ€based histone deacetylase inhibitor enhances cardiomyogenesis in mouse embryonic stem cells. Journal of Cellular Physiology, 2021, 236, 3946-3962.	4.1	3
9	Sphingosylphosphorylcholine blocks ovariectomyâ€induced bone loss by suppressing Ca 2+ /calmodulinâ€mediated osteoclast differentiation. Journal of Cellular and Molecular Medicine, 2021, 25, 473-483.	3.6	4
10	Purification and Crystallographic Analysis of a Novel Cold-Active Esterase (HaEst1) from Halocynthiibacter arcticus. Crystals, 2021, 11, 170.	2.2	2
11	An Antibacterial Nanorobotic Approach for the Specific Targeting and Removal of Multiple Drugâ€Resistant <i>Staphylococcus aureus</i> ). Small, 2021, 17, e2100257.	10.0	20
12	The effect of hairpin loop on the structure and gene expression activity of the long-loop G-quadruplex. Nucleic Acids Research, 2021, 49, 10689-10706.	14.5	12
13	AC-motif: a DNA motif containing adenine and cytosine repeat plays a role in gene regulation. Nucleic Acids Research, 2021, 49, 10150-10165.	14.5	14
14	Ftsh Sensitizes Methicillin-Resistant Staphylococcus aureus to $\hat{l}^2$ -Lactam Antibiotics by Degrading YpfP, a Lipoteichoic Acid Synthesis Enzyme. Antibiotics, 2021, 10, 1198.	3.7	8
15	MD-TSPC4: Computational Method for Predicting the Thermal Stability of I-Motif. International Journal of Molecular Sciences, 2021, 22, 61.	4.1	2
16	Roles of Two-Component Systems in Pseudomonas aeruginosa Virulence. International Journal of Molecular Sciences, 2021, 22, 12152.	4.1	51
17	Characterization and Immobilization of a Novel SGNH Family Esterase (LaSGNH1) from Lactobacillus acidophilus NCFM. International Journal of Molecular Sciences, 2020, 21, 91.	4.1	11
18	Identification, characterization, and immobilization of a novel YbfF esterase from Halomonas elongata. International Journal of Biological Macromolecules, 2020, 165, 1139-1148.	7.5	8

#	Article	IF	CITATIONS
19	Improved differentiation of human adipose stem cells to insulin-producing $\hat{l}^2$ -like cells using PDFGR kinase inhibitor Tyrphostin9. Biochemical and Biophysical Research Communications, 2020, 533, 132-138.	2.1	5
20	Proline Hinged Amphipathic α-Helical Peptide Sensitizes Gram-Negative Bacteria to Various Gram-Positive Antibiotics. Journal of Medicinal Chemistry, 2020, 63, 14937-14950.	6.4	27
21	Exosome-Mediated Differentiation of Mouse Embryonic Fibroblasts and Exocrine Cells into $\hat{l}^2$ -Like Cells and the Identification of Key miRNAs for Differentiation. Biomedicines, 2020, 8, 485.	3.2	6
22	Pontin arginine methylation by CARM1 is crucial for epigenetic regulation of autophagy. Nature Communications, 2020, 11, 6297.	12.8	36
23	Molecular Packing Interaction in DNA Crystals. Crystals, 2020, 10, 1093.	2.2	3
24	Draft Genome Sequences of Lysostaphin-Resistant (K07-204) and Lysostaphin-Susceptible (K07-561) Staphylococcus aureus Sequence Type 72 Strains Isolated from Patients in South Korea. Microbiology Resource Announcements, 2020, 9, .	0.6	2
25	Functional Identification of Serine Hydroxymethyltransferase as a Key Gene Involved in Lysostaphin Resistance and Virulence Potential of Staphylococcus aureus Strains. International Journal of Molecular Sciences, 2020, 21, 9135.	4.1	9
26	Crystallization and Preliminary X-ray Diffraction Study of a Novel Bacterial Homologue of Mammalian Hormone-Sensitive Lipase (halip1) from Halocynthiibacter arcticus. Crystals, 2020, 10, 963.	2.2	1
27	New screening system using Twist1 promoter activity identifies dihydrorotenone as a potent drug targeting cancer-associated fibroblasts. Scientific Reports, 2020, 10, 7058.	3.3	3
28	Biodiesel and flavor compound production using a novel promiscuous cold-adapted SGNH-type lipase (HaSGNH1) from the psychrophilic bacterium Halocynthiibacter arcticus. Biotechnology for Biofuels, 2020, 13, 55.	6.2	21
29	Crystallographic Studies of Enzymes. Crystals, 2020, 10, 6.	2.2	0
30	Ubiquitin-Specific Protease 21 Promotes Colorectal Cancer Metastasis by Acting as a Fra-1 Deubiquitinase. Cancers, 2020, 12, 207.	3.7	28
31	Genome-Wide Analysis of Staphylococcus aureus Sequence Type 72 Isolates Provides Insights Into Resistance Against Antimicrobial Agents and Virulence Potential. Frontiers in Microbiology, 2020, 11, 613800.	3.5	8
32	Targeting Mannitol Metabolism as an Alternative Antimicrobial Strategy Based on the Structure-Function Study of Mannitol-1-Phosphate Dehydrogenase in Staphylococcus aureus. MBio, 2019, 10, .	4.1	22
33	A novel enantioselective SGNH family esterase (NmSGNH1) from Neisseria meningitides: Characterization, mutational analysis, and ester synthesis. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2019, 1864, 1438-1448.	2.4	10
34	Identification of a new Zâ€DNA inducer using SYBR green 1 as a DNA conformation sensor. FEBS Letters, 2019, 593, 2628-2636.	2.8	3
35	Unraveling the Regulatory G-Quadruplex Puzzle: Lessons From Genome and Transcriptome-Wide Studies. Frontiers in Genetics, 2019, 10, 1002.	2.3	15
36	Contribution of SLC22A12 on hypouricemia and its clinical significance for screening purposes. Scientific Reports, 2019, 9, 14360.	3.3	13

3

#	Article	IF	Citations
37	Pneumococcal VncR Strain-Specifically Regulates Capsule Polysaccharide Synthesis. Frontiers in Microbiology, 2019, 10, 2279.	3.5	3
38	Structural and functional characterization of a novel cold-active S-formylglutathione hydrolase (SfSFGH) homolog from Shewanella frigidimarina, a psychrophilic bacterium. Microbial Cell Factories, 2019, 18, 140.	4.0	11
39	Z-DNA in the genome: from structure to disease. Biophysical Reviews, 2019, 11, 383-387.	3.2	65
40	Characterization, immobilization, and mutagenesis of a novel cold-active acetylesterase (EaAcE) from Exiguobacterium antarcticum B7. International Journal of Biological Macromolecules, 2019, 136, 1042-1051.	7.5	7
41	Introduction to the Korean Biophysical Society (KBPS). Biophysical Reviews, 2019, 11, 267-268.	3.2	2
42	Editorial: Special issue of Biophysical Reviews dedicated to the joint 10th Asian Biophysics Association Symposium and 42nd Australian Society for Biophysics Meeting, Melbourne, Australia, December 2–6, 2018. Biophysical Reviews, 2019, 11, 245-247.	3.2	4
43	Suppression of the Ubiquitin Pathway by Small Molecule Binding to Ubiquitin Enhances Doxorubicin Sensitivity of the Cancer Cells. Molecules, 2019, 24, 1073.	3.8	4
44	Characterization and mutation anaylsis of a cold-active bacterial hormone-sensitive lipase from Salinisphaera sp. P7-4. Archives of Biochemistry and Biophysics, 2019, 663, 132-142.	3.0	17
45	Alternative Enzyme Protection Assay To Overcome the Drawbacks of the Gentamicin Protection Assay for Measuring Entry and Intracellular Survival of Staphylococci. Infection and Immunity, 2019, 87, .	2.2	23
46	Molecular Characterization of a Novel Cold-Active Hormone-Sensitive Lipase (HaHSL) from Halocynthiibacter Arcticus. Biomolecules, 2019, 9, 704.	4.0	10
47	Carboxylic Ester Hydrolases in Bacteria: Active Site, Structure, Function and Application. Crystals, 2019, 9, 597.	2.2	24
48	Molecular Characterization of a Novel Family VIII Esterase with $\hat{l}^2$ -Lactamase Activity (PsEstA) from Paenibacillus sp Biomolecules, 2019, 9, 786.	4.0	10
49	Structural Basis for the Enantioselectivity of Esterase Est-Y29 toward ( <i>S</i> )-Ketoprofen. ACS Catalysis, 2019, 9, 755-767.	11.2	14
50	NMR Dynamics Study Reveals the $Z\hat{l}\pm$ Domain of Human ADAR1 Associates with and Dissociates from Z-RNA More Slowly than Z-DNA. ACS Chemical Biology, 2019, 14, 245-255.	3.4	20
51	Chemical-induced formation of BZ-junction with base extrusion. Biochemical and Biophysical Research Communications, 2019, 508, 1215-1220.	2.1	16
52	Structural and functional analysis of a dimeric fumarylacetoacetate hydrolase (EaFAH) from psychrophilic Exiguobacterium antarcticum. Biochemical and Biophysical Research Communications, 2019, 509, 773-778.	2.1	3
53	Discovery of Natural Compounds Promoting Cardiomyocyte Differentiation. Stem Cells and Development, 2019, 28, 13-27.	2.1	4
54	Chemical induced conversion of mouse fibroblasts and human adipose-derived stem cells into skeletal muscle-like cells. Biomaterials, 2019, 193, 30-46.	11.4	23

#	Article	IF	Citations
55	Modulating α-synuclein fibril formation using DNA tetrahedron nanostructures. Biochimica Et Biophysica Acta - General Subjects, 2019, 1863, 73-81.	2.4	3
56	Computational Approaches to Predict the Non-canonical DNAs. Current Bioinformatics, 2019, 14, 470-479.	1.5	7
57	A Selective Inhibitor of Ubiquitin-Specific Protease 4 Suppresses Colorectal Cancer Progression by Regulating $\hat{l}^2$ -Catenin Signaling., 2019, 53, 157-171.		32
58	The FDA-approved anti-cancer drugs, streptozotocin and floxuridine, reduce the virulence of Staphylococcus aureus. Scientific Reports, 2018, 8, 2521.	3.3	45
59	Identification, characterization, immobilization, and mutational analysis of a novel acetylesterase with industrial potential (La AcE) from Lactobacillus acidophilus. Biochimica Et Biophysica Acta - General Subjects, 2018, 1862, 197-210.	2.4	20
60	Crystal structure and functional characterization of a cold-active acetyl xylan esterase (PbAcE) from psychrophilic soil microbe Paenibacillus sp PLoS ONE, 2018, 13, e0206260.	2.5	21
61	Sequence preference and structural heterogeneity of BZ junctions. Nucleic Acids Research, 2018, 46, 10504-10513.	14.5	25
62	Genome-wide analysis of regulatory G-quadruplexes affecting gene expression in human cytomegalovirus. PLoS Pathogens, 2018, 14, e1007334.	4.7	41
63	Total Synthesis of Xanthoangelol B and Its Various Fragments: Toward Inhibition of Virulence Factor Production of <i>Staphylococcus aureus</i> ). Journal of Medicinal Chemistry, 2018, 61, 10473-10487.	6.4	11
64	Identification and Validation of an Antivirulence Agent Targeting HlyU-Regulated Virulence in Vibrio vulnificus. Frontiers in Cellular and Infection Microbiology, 2018, 8, 152.	3.9	24
65	Identification and Crystallographic Analysis of a New Carbohydrate Acetylesterase (SmAcE1) from Sinorhizobium meliloti. Crystals, 2018, 8, 12.	2.2	9
66	Identification of 2′,4′-Dihydroxychalcone as an Antivirulence Agent Targeting HlyU, a Master Virulence Regulator in Vibrio vulnificus. Molecules, 2018, 23, 1492.	3.8	6
67	Crystal structure of a junction between B-DNA and Z-DNA reveals two extruded bases. journal of hand surgery Asian-Pacific volume, The, 2018, , 207-210.	0.4	1
68	Ubiquitin-specific protease 4 (USP4) suppresses myoblast differentiation by down regulating MyoD activity in a catalytic-independent manner. Cellular Signalling, 2017, 35, 48-60.	3.6	10
69	Structural and functional study of ChuY from Escherichia coli strain CFT073. Biochemical and Biophysical Research Communications, 2017, 482, 1176-1182.	2.1	9
70	Characterization of a novel SGNH-type esterase from Lactobacillus plantarum. International Journal of Biological Macromolecules, 2017, 96, 560-568.	7.5	18
71	A synthetic DNA-binding inhibitor of SOX2 guides human induced pluripotent stem cells to differentiate into mesoderm. Nucleic Acids Research, 2017, 45, 9219-9228.	14.5	44
72	Small molecule-induced cellular conversion. Chemical Society Reviews, 2017, 46, 6241-6254.	38.1	29

#	Article	IF	CITATIONS
73	The mutational landscape of ocular marginal zone lymphoma identifies frequent alterations in <i>TNFAIP3</i> followed by mutations in <i>TBL1XR1</i> and <i>CREBBP</i> . Oncotarget, 2017, 8, 17038-17049.	1.8	55
74	Identification and Crystallization of Penicillin-Binding Protein/ $\hat{l}^2$ -Lactamase Homolog (Rp46) from Ruegeria Pomeroyi. Crystals, 2017, 7, 6.	2.2	4
75	Examining cooperative binding of Sox2 on DC5 regulatory element upon complex formation with Pax6 through excess electron transfer assay. Nucleic Acids Research, 2016, 44, e125-e125.	14.5	7
76	Structural and functional studies of a large winged Zâ€∢scp>DNAâ€binding domain of <i>Danio rerio</i> protein kinase <scp>PKZ</scp> . FEBS Letters, 2016, 590, 2275-2285.	2.8	20
77	Biochemical and Structural Analysis of a Novel Esterase from Caulobacter crescentus related to Penicillin-Binding Protein (PBP). Scientific Reports, 2016, 6, 37978.	3.3	16
78	Solution structure of the Z-DNA binding domain of PKR-like protein kinase from <i>Carassius auratus</i> auratus	14.5	25
79	Ubiquitin-specific protease 4 controls metastatic potential through $\hat{l}^2$ -catenin stabilization in brain metastatic lung adenocarcinoma. Scientific Reports, 2016, 6, 21596.	3.3	37
80	An ubiquitin-binding molecule can work as an inhibitor of ubiquitin processing enzymes and ubiquitin receptors. Biochemical and Biophysical Research Communications, 2016, 479, 33-39.	2.1	9
81	Structural Studies on the Extracellular Domain of Sensor Histidine Kinase YycG from Staphylococcus aureus and Its Functional Implications. Journal of Molecular Biology, 2016, 428, 3074-3089.	4.2	19
82	Bioinorganic Nanohybrid Catalyst for Multistep Synthesis of Acetaminophen, an Analgesic. ACS Applied Materials & Samp; Interfaces, 2016, 8, 30058-30065.	8.0	16
83	Coupling of radiofrequency with magnetic nanoparticles treatment as an alternative physical antibacterial strategy against multiple drug resistant bacteria. Scientific Reports, 2016, 6, 33662.	3.3	40
84	Structural and Biochemical Characterization of an Octameric Carbohydrate Acetylesterase from <i>Sinorhizobium meliloti</i> FEBS Letters, 2016, 590, 1242-1252.	2.8	15
85	Photocurrent enhancement of SiNW-FETs by integrating protein-shelled CdSe quantum dots. Nanoscale, 2016, 8, 1921-1925.	5 <b>.</b> 6	3
86	Design of a RANK-Mimetic Peptide Inhibitor of Osteoclastogenesis with Enhanced RANKL-Binding Affinity. Molecules and Cells, 2016, 39, 316-321.	2.6	6
87	Crystal structure analysis of c4763, a uropathogenicEscherichia coli-specific protein. Acta Crystallographica Section F, Structural Biology Communications, 2015, 71, 1042-1047.	0.8	1
88	Structural basis for the substrate selectivity of a HAD phosphatase from Thermococcus onnurineus NA1. Biochemical and Biophysical Research Communications, 2015, 461, 122-127.	2.1	2
89	Ubiquitin specific protease 4 positively regulates the WNT/βâ€catenin signaling in colorectal cancer. Molecular Oncology, 2015, 9, 1834-1851.	4.6	90
90	Preparation of cobalt nanoparticles from polymorphic bacterial templates: A novel platform for biocatalysis. International Journal of Biological Macromolecules, 2015, 81, 747-753.	<b>7.</b> 5	12

#	Article	IF	Citations
91	Combining Suppression of Stemness with Lineage-Specific Induction Leads to Conversion of Pluripotent Cells into Functional Neurons. Chemistry and Biology, 2015, 22, 1512-1520.	6.0	7
92	Structural and biochemical characterization of a carbohydrate acetylesterase from <i>Sinorhizobium meliloti 1021</i> . FEBS Letters, 2015, 589, 117-122.	2.8	18
93	Structural insights into the molecular mechanism of <i>Escherichia coli</i> SdiA, a quorum-sensing receptor. Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 694-707.	2.5	55
94	Combining Protein-Shelled Platinum Nanoparticles with Graphene to Build a Bionanohybrid Capacitor. ACS Nano, 2014, 8, 12120-12129.	14.6	14
95	Interaction of TIF-90 and filamin A in the regulation of rRNA synthesis in leukemic cells. Blood, 2014, 124, 579-589.	1.4	13
96	Inhibition of master transcription factors in pluripotent cells induces early stage differentiation. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1778-1783.	7.1	10
97	Crystallographic analysis and biochemical applications of a novel penicillin-binding protein $\hat{N}^2$ -lactamase homologue from a metagenomic library. Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 2455-2466.	2.5	32
98	Crystallization and preliminary X-ray analysis of a highly stable novel SGNH hydrolase (Est24) from <i>Sinorhizobium meliloti</i> . Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 193-195.	0.8	2
99	Crystallization and preliminary X-ray analysis of a novel type of lipolytic hydrolase fromBacillus licheniformis. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 473-475.	0.8	0
100	Structural and kinetic bases for the metal preference of the M18 aminopeptidase from Pseudomonas aeruginosa. Biochemical and Biophysical Research Communications, 2014, 447, 101-107.	2.1	13
101	Novel Interaction of the Z-DNA Binding Domain of Human ADAR1 with the Oncogenic c-Myc Promoter G-Quadruplex. Journal of Molecular Biology, 2014, 426, 2594-2604.	4.2	24
102	Single Molecule Visualization and Characterization of Sox2–Pax6 Complex Formation on a Regulatory DNA Element Using a DNA Origami Frame. Nano Letters, 2014, 14, 2286-2292.	9.1	38
103	Distinct Rayleigh Scattering from Hot Spot Mutant p53 Proteins Reveals Cancer Cells. Small, 2014, 10, 2954-2962.	10.0	5
104	Radiofrequency treatment enhances the catalytic function of an immobilized nanobiohybrid catalyst. Nanoscale, 2014, 6, 6009-6017.	5.6	6
105	Distinct Z-DNA binding mode of a PKR-like protein kinase containing a Z-DNA binding domain (PKZ). Nucleic Acids Research, 2014, 42, 5937-5948.	14.5	46
106	Structural and functional characterization of an Isd-type haem-degradation enzyme from <i>Listeria monocytogenes</i> . Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 615-626.	2.5	32
107	Investigation of the heating properties of platinum nanoparticles under a radiofrequency current. International Journal of Hyperthermia, 2013, 29, 99-105.	2.5	30
108	Energetics of Z-DNA Binding Protein-Mediated Helicity Reversals in DNA, RNA, and DNA–RNA Duplexes. Journal of Physical Chemistry B, 2013, 117, 13866-13871.	2.6	11

#	Article	IF	CITATIONS
109	Synthesis and electric characterization of protein-shelled CdSe quantum dots. Journal of Materials Chemistry C, 2013, 1, 2412.	5.5	8
110	One-step immobilization and purification of his-tagged enzyme using poly(2-acetamidoacrylic acid) hydrogel. Macromolecular Research, 2013, 21, 5-9.	2.4	12
111	Size-controlled synthesis and characterization of CoPt nanoparticles using protein shells. Journal of Materials Chemistry B, 2013, 1, 1453.	5.8	18
112	DNA Sensing-Independent Inhibition of Herpes Simplex Virus 1 Replication by DAI/ZBP1. Journal of Virology, 2013, 87, 3076-3086.	3.4	58
113	Structure-based elucidation of the regulatory mechanism for aminopeptidase activity. Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 1738-1747.	2.5	4
114	Structural and functional analyses of a bacterial homologue of hormone-sensitive lipase from a metagenomic library. Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 1726-1737.	2.5	33
115	Glutamyl Aminopeptidase (Lactococcus). , 2013, , 1631-1635.		0
116	The effect of protein shells on the antioxidant activity of protein-encapsulated platinum nanoparticles. Journal of Materials Chemistry, 2012, 22, 1774-1780.	6.7	20
117	EZH2 Generates a Methyl Degron that Is Recognized by the DCAF1/DDB1/CUL4 E3ÂUbiquitin Ligase Complex. Molecular Cell, 2012, 48, 572-586.	9.7	200
118	Characterization, amyloid formation, and immobilization of a novel SGNH hydrolase from Listeria innocua 11262. International Journal of Biological Macromolecules, 2012, 50, 103-111.	7.5	22
119	The crystal structure of the periplasmic domain of <i>Vibrio parahaemolyticus</i> CpxA. Protein Science, 2012, 21, 1334-1343.	7.6	13
120	Characterization, crystallization and preliminary X-ray diffraction analysis of an ( <i>S</i> )-specific esterase ( <i>pf</i> EstA) from <i>Pseudomonas fluorescens</i> KCTC 1767: enantioselectivity for potential industrial applications. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 1374-1377.	0.7	3
121	Structures of Staphylococcus aureuspeptide deformylase in complex with two classes of new inhibitors. Acta Crystallographica Section D: Biological Crystallography, 2012, 68, 784-793.	2.5	4
122	Identification, crystallization and preliminary X-ray diffraction analysis of esterase A fromCaulobacter crescentusCB15, a family VIII lipolytic enzyme. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 560-564.	0.7	6
123	Intrinsic Z-DNA Is Stabilized by the Conformational Selection Mechanism of Z-DNA-Binding Proteins. Journal of the American Chemical Society, 2011, 133, 668-671.	13.7	92
124	Crystal Structure of the LG3 Domain of Endorepellin, an Angiogenesis Inhibitor. Journal of Molecular Biology, 2011, 414, 231-242.	4.2	18
125	The $Z\hat{l}^2$ domain of human DAI binds to Z-DNA via a novel B-Z transition pathway. FEBS Letters, 2011, 585, 772-778.	2.8	19
126	Amyloid formation using 1-butyl-3-methyl-imidazolium-based ionic liquids. Analytical Biochemistry, 2011, 419, 354-356.	2.4	24

#	Article	IF	CITATIONS
127	Crystallization and diffraction analysis of Sm23: an SGNH-family arylesterase from <i>Sinorhizobium meliloti</i> 1021. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 572-574.	0.7	1
128	Platinum Nanoparticles Encapsulated by Aminopeptidase: A Multifunctional Bioinorganic Nanohybrid Catalyst. Angewandte Chemie - International Edition, 2011, 50, 11924-11929.	13.8	60
129	Solution structure of the $Z\hat{I}^2$ domain of human DNA-dependent activator of IFN-regulatory factors and its binding modes to B- and Z-DNAs. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 6921-6926.	7.1	36
130	Structural Basis for the Reaction Mechanism of UDP-Glucose Pyrophosphorylase. Molecules and Cells, 2010, 29, 397-406.	2.6	41
131	Crystal structure of <i>Streptococcus pneumoniae</i> Sp1610, a putative tRNA methyltransferase, in complex with <i>S</i> â€adenosylâ€ <scp>L</scp> â€methionine. Protein Science, 2010, 19, 617-624.	7.6	7
132	Structural basis for the negative regulation of bacterial stress response by RseB. Protein Science, 2010, 19, 1258-1263.	7.6	25
133	The crystal structure <i>Escherichia coli</i> Spy. Protein Science, 2010, 19, 2252-2259.	7.6	31
134	Structure-based development of a receptor activator of nuclear factor-κB ligand (RANKL) inhibitor peptide and molecular basis for osteopetrosis. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 20281-20286.	7.1	67
135	Z-DNA Binding Proteins as Targets for Structure-Based Virtual Screening. Current Drug Targets, 2010, 11, 335-344.	2.1	29
136	Transition between B-DNA and Z-DNA: Free Energy Landscape for the Bâ^'Z Junction Propagation. Journal of Physical Chemistry B, 2010, 114, 9872-9881.	2.6	26
137	Structural basis for the substrate specificity of PepA from Streptococcus pneumoniae, a dodecameric tetrahedral protease. Biochemical and Biophysical Research Communications, 2010, 391, 431-436.	2.1	41
138	Analyzing the Interaction of RseA and RseB, the Two Negative Regulators of the ÏfE Envelope Stress Response, Using a Combined Bioinformatic and Experimental Strategy. Journal of Biological Chemistry, 2009, 284, 5403-5413.	3.4	11
139	The structures of non-CG-repeat Z-DNAs co-crystallized with the Z-DNA-binding domain, hZα ADAR1. Nucleic Acids Research, 2009, 37, 629-637.	14.5	67
140	Base extrusion is found at helical junctions between right- and left-handed forms of DNA and RNA. Nucleic Acids Research, 2009, 37, 4353-4359.	14.5	36
141	Disulfide Bond as a Structural Determinant of Prion Protein Membrane Insertion. Molecules and Cells, 2009, 27, 673-680.	2.6	12
142	Crystallization and preliminary X-ray crystallographic studies of the Z-DNA-binding domain of a PKR-like kinase (PKZ) in complex with Z-DNA. Acta Crystallographica Section F: Structural Biology Communications, 2009, 65, 267-270.	0.7	14
143	Purification, crystallization and preliminary crystallographic analysis of Est-Y29: a novel oligomeric $\hat{l}^2$ -lactamase. Acta Crystallographica Section F: Structural Biology Communications, 2009, 65, 310-312.	0.7	7
144	Structural and functional characterization of soluble endoglin receptor. Biochemical and Biophysical Research Communications, 2009, 383, 386-391.	2.1	15

#	Article	IF	CITATIONS
145	Enhanced SUMOylation of proteins containing a SUMO-interacting motif by SUMO-Ubc9 fusion. Biochemical and Biophysical Research Communications, 2009, 388, 41-45.	2.1	12
146	NMR Spectroscopic Elucidation of the Bâ^2 Transition of a DNA Double Helix Induced by the $Z\hat{l}\pm$ Domain of Human ADAR1. Journal of the American Chemical Society, 2009, 131, 11485-11491.	13.7	67
147	Crystal structure of the mouse p53 core domain in zincâ€free state. Proteins: Structure, Function and Bioinformatics, 2008, 70, 280-283.	2.6	6
148	Solution structures of RseA and its complex with RseB. Journal of Synchrotron Radiation, 2008, 15, 219-222.	2.4	18
149	Crystallization and preliminary X-ray studies of SdiA from <i>Escherichia coli</i> . Acta Crystallographica Section F: Structural Biology Communications, 2008, 64, 19-21.	0.7	3
150	The Crystal Structure of Guamerin in Complex with Chymotrypsin and the Development of an Elastase-specific Inhibitor. Journal of Molecular Biology, 2008, 376, 184-192.	4.2	15
151	The Mechanism of Temperature-Induced Bacterial HtrA Activation. Journal of Molecular Biology, 2008, 377, 410-420.	4.2	19
152	The Structural Basis for the Activation and Peptide Recognition of Bacterial ClpP. Journal of Molecular Biology, 2008, 379, 760-771.	4.2	49
153	The crystal structure of the second Z-DNA binding domain of human DAI (ZBP1) in complex with Z-DNA reveals an unusual binding mode to Z-DNA. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 20671-20676.	7.1	99
154	Crystal structure of RseB and a model of its binding mode to RseA. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 8779-8784.	7.1	50
155	Characterization of DNA-binding activity of ZÂ domains from poxviruses and the importance of the -wing regions in converting B-DNA to Z-DNA. Nucleic Acids Research, 2007, 35, 7714-7720.	14.5	18
156	Crystallization and preliminary X-ray studies of TON_1713 fromThermococcus onnurineusNA1, a putative member of the haloacid dehalogenase superfamily. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 1048-1050.	0.7	0
157	Intracellular localization of human ZBP1: Differential regulation by the Z-DNA binding domain, $Z\hat{l}_{\pm}$ , in splice variants. Biochemical and Biophysical Research Communications, 2006, 348, 145-152.	2.1	25
158	Crystal Structure of Visfatin/Pre-B Cell Colony-enhancing Factor 1/Nicotinamide Phosphoribosyltransferase, Free and in Complex with the Anti-cancer Agent FK-866. Journal of Molecular Biology, 2006, 362, 66-77.	4.2	107
159	S2c2-5 Z-DNA and its binding proteins(S2-c2: "Structural biology reveals macromolecular) Tj ETQq1 1 0.784314	rgBT/Ove	rlogk 10 Tf 5
160	A peptide with alternating lysines can act as a highly specific Z-DNA binding domain. Nucleic Acids Research, 2006, 34, 4937-4942.	14.5	15
161	Crystal structure of a junction between B-DNA and Z-DNA reveals two extruded bases. Nature, 2005, 437, 1183-1186.	27.8	261
162	A poxvirus protein forms a complex with left-handed Z-DNA: Crystal structure of a Yatapoxvirus ZÂ bound to DNA. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 14367-14372.	7.1	106

#	Article	IF	CITATION
163	Evidence that vaccinia virulence factor E3L binds to Z-DNA in vivo: Implications for development of a therapy for poxvirus infection. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 1514-1518.	7.1	92
164	The crystal structure of a triacylglycerol lipase from Pseudomonas cepacia reveals a highly open conformation in the absence of a bound inhibitor. Structure, 1997, 5, 173-185.	3.3	301
165	Crystal structure of carboxylesterase from Pseudomonas fluorescens, an $\hat{l}\pm\hat{l}^2$ hydrolase with broad substrate specificity. Structure, 1997, 5, 1571-1584.	3.3	109