Walter J Chazin

List of Publications by Year in Descending Order

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

12,652 63 108 187 h-index g-index citations papers 6.46 9.6 14,598 239 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
187	Neutrophil-associated responses to infection in a natural host model <i>Infection and Immunity</i> , 2022 , iai0	094662	. 1 0
186	TdfH selectively binds metal-loaded tetrameric calprotectin for zinc import <i>Communications Biology</i> , 2022 , 5, 103	6.7	O
185	Adherence enables to overcome zinc limitation imposed by nutritional immunity proteins <i>Infection and Immunity</i> , 2022 , iai0000922	3.7	O
184	Zn-regulated GTPase metalloprotein activator 1 modulates vertebrate zinc homeostasis <i>Cell</i> , 2022	56.2	5
183	Gain-of-Function Mutations in RPA1 Cause a Syndrome with Short Telomeres and Somatic Genetic Rescue. <i>Blood</i> , 2021 ,	2.2	2
182	Siderophore-mediated zinc acquisition enhances enterobacterial colonization of the inflamed gut. <i>Nature Communications</i> , 2021 , 12, 7016	17.4	3
181	RADX controls RAD51 filament dynamics to regulate replication fork stability. <i>Molecular Cell</i> , 2021 , 81, 1074-1083.e5	17.6	6
180	EXO5-DNA structure and BLM interactions direct DNA resection critical for ATR-dependent replication restart. <i>Molecular Cell</i> , 2021 , 81, 2989-3006.e9	17.6	8
179	A fragment-based approach to discovery of Receptor for Advanced Glycation End products inhibitors. <i>Proteins: Structure, Function and Bioinformatics</i> , 2021 , 89, 1399-1412	4.2	2
178	Neutrophil extracellular traps enhance macrophage killing of bacterial pathogens. <i>Science Advances</i> , 2021 , 7, eabj2101	14.3	9
177	DNA Recognition/Processing DNA Polymerase Alpha-Primase: Biochemical and Structural Mechanisms 2021 , 431-444		O
176	Molecular Insight into TdfH-Mediated Zinc Piracy from Human Calprotectin by Neisseria gonorrhoeae. <i>MBio</i> , 2020 , 11,	7.8	6
175	ZupT Facilitates Clostridioides difficile Resistance to Host-Mediated Nutritional Immunity. <i>MSphere</i> , 2020 , 5,	5	11
174	Calmodulin Mutations Associated with Heart Arrhythmia: A Status Report. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	15
173	A slipped-CAG DNA-binding small molecule induces trinucleotide-repeat contractions in vivo. Nature Genetics, 2020 , 52, 146-159	36.3	54
172	Identification of ubiquitin Ser57 kinases regulating the oxidative stress response in yeast. <i>ELife</i> , 2020 , 9,	8.9	2
171	A key interaction with RPA orients XPA in NER complexes. <i>Nucleic Acids Research</i> , 2020 , 48, 2173-2188	20.1	13

(2018-2020)

170	Envisioning how the prototypic molecular machine TFIIH functions in transcription initiation and DNA repair. <i>DNA Repair</i> , 2020 , 96, 102972	4.3	16
169	The anti-parasitic agent suramin and several of its analogues are inhibitors of the DNA binding protein Mcm10. <i>Open Biology</i> , 2019 , 9, 190117	7	10
168	Genetic Mosaicism in Calmodulinopathy. Circulation Genomic and Precision Medicine, 2019, 12, 375-385	5.2	20
167	Dynamics and selective remodeling of the DNA-binding domains of RPA. <i>Nature Structural and Molecular Biology</i> , 2019 , 26, 129-136	17.6	35
166	A new approach to discovery of S100 protein heterodimers. FEBS Journal, 2019, 286, 1838-1840	5.7	
165	Multi-metal Restriction by Calprotectin Impacts De Novo Flavin Biosynthesis in Acinetobacter baumannii. <i>Cell Chemical Biology</i> , 2019 , 26, 745-755.e7	8.2	35
164	The Innate Immune Protein S100A9 Protects from T-Helper Cell Type 2-mediated Allergic Airway Inflammation. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2019 , 61, 459-468	5.7	14
163	The novel interaction between Neisseria gonorrhoeae TdfJ and human S100A7 allows gonococci to subvert host zinc restriction. <i>PLoS Pathogens</i> , 2019 , 15, e1007937	7.6	11
162	S100 Proteins in the Innate Immune Response to Pathogens. <i>Methods in Molecular Biology</i> , 2019 , 1929, 275-290	1.4	25
161	An Acinetobacter baumannii, Zinc-Regulated Peptidase Maintains Cell Wall Integrity during Immune-Mediated Nutrient Sequestration. <i>Cell Reports</i> , 2019 , 26, 2009-2018.e6	10.6	36
160	The Immune Protein Calprotectin Impacts Clostridioides difficile Metabolism through Zinc Limitation. <i>MBio</i> , 2019 , 10,	7.8	7
159	Arachidonic Acid Kills Staphylococcus aureus through a Lipid Peroxidation Mechanism. <i>MBio</i> , 2019 , 10,	7.8	2 0
158	Identifying the substrate proteins of U-box E3s E4B and CHIP by orthogonal ubiquitin transfer. <i>Science Advances</i> , 2018 , 4, e1701393	14.3	19
157	A Mechanism of Calmodulin Modulation of the Human Cardiac Sodium Channel. <i>Structure</i> , 2018 , 26, 683	3- <u>6.9</u> 4.€	:332
156	Stress-induced acidification may contribute to formation of unusual structures in C9orf72-repeats. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2018 , 1862, 1482-1491	4	6
155	Calprotectin protects against experimental colonic inflammation in mice. <i>British Journal of Pharmacology</i> , 2018 , 175, 3797-3812	8.6	13
154	Single-Molecule Analysis of Replication Protein A-DNA Interactions. <i>Methods in Enzymology</i> , 2018 , 600, 439-461	1.7	3
153	Role of Calprotectin in Withholding Zinc and Copper from Candida albicans. <i>Infection and Immunity</i> , 2018 , 86,	3.7	65

152	Substrate Binding Regulates Redox Signaling in Human DNA Primase. <i>Journal of the American Chemical Society</i> , 2018 , 140, 17153-17162	16.4	6
151	Functional and structural similarity of human DNA primase [4Fe4S] cluster domain constructs. <i>PLoS ONE</i> , 2018 , 13, e0209345	3.7	4
150	Yeast require redox switching in DNA primase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, 13186-13191	11.5	10
149	Characterization and development of SAPP as a specific peptidic inhibitor that targets Porphyromonas gingivalis. <i>Molecular Oral Microbiology</i> , 2018 , 33, 430-439	4.6	7
148	Disrupted structure and aberrant function of CHIP mediates the loss of motor and cognitive function in preclinical models of SCAR16. <i>PLoS Genetics</i> , 2018 , 14, e1007664	6	19
147	A minimal threshold of FANCJ helicase activity is required for its response to replication stress or double-strand break repair. <i>Nucleic Acids Research</i> , 2018 , 46, 6238-6256	20.1	15
146	RPA Interacts with HIRA and Regulates H3.3 Deposition at Gene Regulatory Elements in Mammalian Cells. <i>Molecular Cell</i> , 2017 , 65, 272-284	17.6	51
145	The [4Fe4S] cluster of human DNA primase functions as a redox switch using DNA charge transport. <i>Science</i> , 2017 , 355,	33.3	82
144	Prp40 Homolog A Is a Novel Centrin Target. <i>Biophysical Journal</i> , 2017 , 112, 2529-2539	2.9	13
143	Molecular basis for PrimPol recruitment to replication forks by RPA. <i>Nature Communications</i> , 2017 , 8, 15222	17.4	47
142	A Polymerase With Potential: The Fe-S Cluster in Human DNA Primase. <i>Methods in Enzymology</i> , 2017 , 595, 361-390	1.7	5
141	Dietary Manganese Promotes Staphylococcal Infection of the Heart. <i>Cell Host and Microbe</i> , 2017 , 22, 531-542.e8	23.4	37
140	Response to Comments on "The [4Fe4S] cluster of human DNA primase functions as a redox switch using DNA charge transport". <i>Science</i> , 2017 , 357,	33.3	4
139	Analysis of DNA binding by human factor xeroderma pigmentosum complementation group A (XPA) provides insight into its interactions with nucleotide excision repair substrates. <i>Journal of Biological Chemistry</i> , 2017 , 292, 16847-16857	5.4	11
138	RADX Promotes Genome Stability and Modulates Chemosensitivity by Regulating RAD51 at Replication Forks. <i>Molecular Cell</i> , 2017 , 67, 374-386.e5	17.6	98
137	Analysis of Functional Dynamics of Modular Multidomain Proteins by SAXS and NMR. <i>Methods in Enzymology</i> , 2017 , 592, 49-76	1.7	5
136	Ubiquitin turnover and endocytic trafficking in yeast are regulated by Ser57 phosphorylation of ubiquitin. <i>ELife</i> , 2017 , 6,	8.9	19
135	Novel CPVT-Associated Calmodulin Mutation in CALM3 (CALM3-A103V) Activates Arrhythmogenic Ca Waves and Sparks. <i>Circulation: Arrhythmia and Electrophysiology</i> , 2016 , 9,	6.4	59

(2016-2016)

134	Novel calmodulin mutations associated with congenital long QT syndrome affect calcium current in human cardiomyocytes. <i>Heart Rhythm</i> , 2016 , 13, 2012-9	6.7	41
133	The innate immune protein calprotectin promotes Pseudomonas aeruginosa and Staphylococcus aureus interaction. <i>Nature Communications</i> , 2016 , 7, 11951	17.4	70
132	XPA: A key scaffold for human nucleotide excision repair. DNA Repair, 2016, 44, 123-135	4.3	68
131	Data publication with the structural biology data grid supports live analysis. <i>Nature Communications</i> , 2016 , 7, 10882	17.4	78
130	The Response of Acinetobacter baumannii to Zinc Starvation. <i>Cell Host and Microbe</i> , 2016 , 19, 826-36	23.4	79
129	Salmonella Mitigates Oxidative Stress and Thrives in the Inflamed Gut by Evading Calprotectin-Mediated Manganese Sequestration. <i>Cell Host and Microbe</i> , 2016 , 19, 814-25	23.4	74
128	Molecular Basis for the Interaction Between AP4 4 and its Accessory Protein, Tepsin. <i>Traffic</i> , 2016 , 17, 400-15	5.7	16
127	Single-molecule imaging reveals the mechanism of Exo1 regulation by single-stranded DNA binding proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E1	176:5	65
126	Spectrum and Prevalence of CALM1-, CALM2-, and CALM3-Encoded Calmodulin Variants in Long QT Syndrome and Functional Characterization of a Novel Long QT Syndrome-Associated Calmodulin Missense Variant, E141G. <i>Circulation: Cardiovascular Genetics</i> , 2016 , 9, 136-146		77
125	The pattern recognition reagents RAGE VC1 and peptide p5 share common binding sites and exhibit specific reactivity with AA amyloid in mice. <i>Amyloid: the International Journal of Experimental and Clinical Investigation: the Official Journal of the International Society of Amyloidosis</i>	2.7	3
124	CacyBP/SIPStructure and variety of functions. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2016 , 1860, 79-85	4	30
123	Zinc and Manganese Chelation by Neutrophil S100A8/A9 (Calprotectin) Limits Extracellular Aspergillus fumigatus Hyphal Growth and Corneal Infection. <i>Journal of Immunology</i> , 2016 , 196, 336-44	5.3	91
122	Identification and Optimization of Anthranilic Acid Based Inhibitors of Replication Protein A. <i>ChemMedChem</i> , 2016 , 11, 893-9	3.7	9
121	Dynamic binding of replication protein a is required for DNA repair. <i>Nucleic Acids Research</i> , 2016 , 44, 5758-72	20.1	45
120	Dietary zinc alters the microbiota and decreases resistance to Clostridium difficile infection. <i>Nature Medicine</i> , 2016 , 22, 1330-1334	50.5	136
119	Binding of transition metals to S100 proteins. <i>Science China Life Sciences</i> , 2016 , 59, 792-801	8.5	40
118	Acinetobacter baumannii Coordinates Urea Metabolism with Metal Import To Resist Host-Mediated Metal Limitation. <i>MBio</i> , 2016 , 7,	7.8	40
117	ETAA1 acts at stalled replication forks to maintain genome integrity. <i>Nature Cell Biology</i> , 2016 , 18, 1185	5-2131.295	132

116	Nutritional Immunity: S100 Proteins at the Host-Pathogen Interface. <i>Journal of Biological Chemistry</i> , 2015 , 290, 18991-8	5.4	136
115	Functional dynamics in replication protein A DNA binding and protein recruitment domains. <i>Structure</i> , 2015 , 23, 1028-38	5.2	21
114	Promotion of BRCA2-Dependent Homologous Recombination by DSS1 via RPA Targeting and DNA Mimicry. <i>Molecular Cell</i> , 2015 , 59, 176-87	17.6	97
113	Characteristics and concepts of dynamic hub proteins in DNA processing machinery from studies of RPA. <i>Progress in Biophysics and Molecular Biology</i> , 2015 , 117, 206-211	4.7	22
112	Zinc regulates a switch between primary and alternative S18 ribosomal proteins in Mycobacterium tuberculosis. <i>Molecular Microbiology</i> , 2015 , 97, 263-80	4.1	30
111	Arrhythmogenic Calmodulin Mutations Affect the Activation and Termination of Cardiac Ryanodine Receptor-mediated Ca2+ Release. <i>Journal of Biological Chemistry</i> , 2015 , 290, 26151-62	5.4	45
110	Diphenylpyrazoles as replication protein a inhibitors. ACS Medicinal Chemistry Letters, 2015, 6, 140-5	4.3	15
109	Helicobacter pylori Resists the Antimicrobial Activity of Calprotectin via Lipid A Modification and Associated Biofilm Formation. <i>MBio</i> , 2015 , 6, e01349-15	7.8	30
108	Mechanochemical regulations of RPAR binding to ssDNA. Scientific Reports, 2015, 5, 9296	4.9	26
107	Simian virus Large T antigen interacts with the N-terminal domain of the 70 kD subunit of Replication Protein A in the same mode as multiple DNA damage response factors. <i>PLoS ONE</i> , 2015 , 10, e0116093	3.7	4
106	Calprotectin Increases the Activity of the SaeRS Two Component System and Murine Mortality during Staphylococcus aureus Infections. <i>PLoS Pathogens</i> , 2015 , 11, e1005026	7.6	38
105	The Human Antimicrobial Protein Calgranulin C Participates in Control of Helicobacter pylori Growth and Regulation of Virulence. <i>Infection and Immunity</i> , 2015 , 83, 2944-56	3.7	47
104	Dimerization and phosphatase activity of calcyclin-binding protein/Siah-1 interacting protein: the influence of oxidative stress. <i>FASEB Journal</i> , 2015 , 29, 1711-24	0.9	16
103	Human PrimPol is a highly error-prone polymerase regulated by single-stranded DNA binding proteins. <i>Nucleic Acids Research</i> , 2015 , 43, 1056-68	20.1	63
102	Biochemical and Proteomic Analysis of Ubiquitination of Hsc70 and Hsp70 by the E3 Ligase CHIP. <i>PLoS ONE</i> , 2015 , 10, e0128240	3.7	19
101	Insights into eukaryotic primer synthesis from structures of the p48 subunit of human DNA primase. <i>Journal of Molecular Biology</i> , 2014 , 426, 558-69	6.5	20
100	Diffusion of human replication protein A along single-stranded DNA. <i>Journal of Molecular Biology</i> , 2014 , 426, 3246-3261	6.5	85
99	Redefining the DNA-binding domain of human XPA. <i>Journal of the American Chemical Society</i> , 2014 , 136, 10830-3	16.4	29

(2012-2014)

98	Structural analysis of replication protein A recruitment of the DNA damage response protein SMARCAL1. <i>Biochemistry</i> , 2014 , 53, 3052-61	3.2	20
97	Acinetobacter baumannii response to host-mediated zinc limitation requires the transcriptional regulator Zur. <i>Journal of Bacteriology</i> , 2014 , 196, 2616-26	3.5	67
96	Novel calmodulin mutations associated with congenital arrhythmia susceptibility. <i>Circulation: Cardiovascular Genetics</i> , 2014 , 7, 466-74		133
95	Discovery of a potent stapled helix peptide that binds to the 70N domain of replication protein A. <i>Journal of Medicinal Chemistry</i> , 2014 , 57, 2455-61	8.3	39
94	Novel function of the Fanconi anemia group J or RECQ1 helicase to disrupt protein-DNA complexes in a replication protein A-stimulated manner. <i>Journal of Biological Chemistry</i> , 2014 , 289, 19928-41	5.4	26
93	The host protein calprotectin modulates the Helicobacter pylori cag type IV secretion system via zinc sequestration. <i>PLoS Pathogens</i> , 2014 , 10, e1004450	7.6	65
92	Replication protein A: single-stranded DNAß first responder: dynamic DNA-interactions allow replication protein A to direct single-strand DNA intermediates into different pathways for synthesis or repair. <i>BioEssays</i> , 2014 , 36, 1156-61	4.1	139
91	Divergent regulation of ryanodine receptor 2 calcium release channels by arrhythmogenic human calmodulin missense mutants. <i>Circulation Research</i> , 2014 , 114, 1114-24	15.7	96
90	Surface reengineering of RPA70N enables cocrystallization with an inhibitor of the replication protein A interaction motif of ATR interacting protein. <i>Biochemistry</i> , 2013 , 52, 6515-24	3.2	14
89	Discovery of a potent inhibitor of replication protein a protein-protein interactions using a fragment-linking approach. <i>Journal of Medicinal Chemistry</i> , 2013 , 56, 9242-50	8.3	48
88	Xeroderma pigmentosum complementation group C protein (XPC) serves as a general sensor of damaged DNA. <i>DNA Repair</i> , 2013 , 12, 947-53	4.3	36
87	Calmodulin mutations associated with recurrent cardiac arrest in infants. Circulation, 2013, 127, 1009-1	716.7	262
86	Molecular basis for manganese sequestration by calprotectin and roles in the innate immune response to invading bacterial pathogens. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 3841-6	11.5	245
85	Activation of UbcH5c~Ub is the result of a shift in interdomain motions of the conjugate bound to U-box E3 ligase E4B. <i>Biochemistry</i> , 2013 , 52, 2991-9	3.2	35
84	Zinc piracy as a mechanism of Neisseria meningitidis for evasion of nutritional immunity. <i>PLoS Pathogens</i> , 2013 , 9, e1003733	7.6	57
83	A new structural framework for integrating replication protein A into DNA processing machinery. <i>Nucleic Acids Research</i> , 2013 , 41, 2313-27	20.1	62
82	MntABC and MntH contribute to systemic Staphylococcus aureus infection by competing with calprotectin for nutrient manganese. <i>Infection and Immunity</i> , 2013 , 81, 3395-405	3.7	143
81	Zinc sequestration by the neutrophil protein calprotectin enhances Salmonella growth in the inflamed gut. <i>Cell Host and Microbe</i> , 2012 , 11, 227-39	23.4	243

80	Structure of an E3:E2~Ub complex reveals an allosteric mechanism shared among RING/U-box ligases. <i>Molecular Cell</i> , 2012 , 47, 933-42	17.6	217
79	Repair-specific functions of replication protein A. <i>Journal of Biological Chemistry</i> , 2012 , 287, 3908-18	5.4	26
78	Human DNA helicase B (HDHB) binds to replication protein A and facilitates cellular recovery from replication stress. <i>Journal of Biological Chemistry</i> , 2012 , 287, 6469-81	5.4	30
77	A high-throughput fluorescence polarization anisotropy assay for the 70N domain of replication protein A. <i>Analytical Biochemistry</i> , 2012 , 421, 742-9	3.1	33
76	Identification of an Acinetobacter baumannii zinc acquisition system that facilitates resistance to calprotectin-mediated zinc sequestration. <i>PLoS Pathogens</i> , 2012 , 8, e1003068	7.6	184
75	Relating form and function of EF-hand calcium binding proteins. <i>Accounts of Chemical Research</i> , 2011 , 44, 171-9	24.3	90
74	Nutrient metal sequestration by calprotectin inhibits bacterial superoxide defense, enhancing neutrophil killing of Staphylococcus aureus. <i>Cell Host and Microbe</i> , 2011 , 10, 158-64	23.4	273
73	Solution NMR structure of Apo-calmodulin in complex with the IQ motif of human cardiac sodium channel NaV1.5. <i>Journal of Molecular Biology</i> , 2011 , 406, 106-19	6.5	92
72	E2 conjugating enzyme selectivity and requirements for function of the E3 ubiquitin ligase CHIP. Journal of Biological Chemistry, 2011 , 286, 21277-86	5.4	37
71	BID binds to replication protein A and stimulates ATR function following replicative stress. <i>Molecular and Cellular Biology</i> , 2011 , 31, 4298-309	4.8	22
70	A naturally occurring human RPA subunit homolog does not support DNA replication or cell-cycle progression. <i>Nucleic Acids Research</i> , 2010 , 38, 846-58	20.1	27
69	Insights into eukaryotic DNA priming from the structure and functional interactions of the 4Fe-4S cluster domain of human DNA primase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 13684-9	11.5	66
68	Functional characterization of a cancer causing mutation in human replication protein A. <i>Molecular Cancer Research</i> , 2010 , 8, 1017-26	6.6	23
67	Reconstitution of RPA-covered single-stranded DNA-activated ATR-Chk1 signaling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 13660-5	11.5	90
66	Biochemical and Structural Domain Analysis of Xeroderma Pigmentosum Complementation Group C Protein. <i>ACS Symposium Series</i> , 2010 , 59-71	0.4	
65	Structural dynamics and single-stranded DNA binding activity of the three N-terminal domains of the large subunit of replication protein A from small angle X-ray scattering. <i>Biochemistry</i> , 2010 , 49, 288	30 ³⁹²	34
64	Structural and functional characterization of the monomeric U-box domain from E4B. <i>Biochemistry</i> , 2010 , 49, 347-55	3.2	27
63	Structural basis for ligand recognition and activation of RAGE. <i>Structure</i> , 2010 , 18, 1342-52	5.2	164

(2006-2010)

62	Defining the modular protein intereactions that coordinate recruitment of DNA polymerase alpha to initiate SV40 DNA replication. <i>FASEB Journal</i> , 2010 , 24, 196.2	0.9	
61	Engineering a ubiquitin ligase reveals conformational flexibility required for ubiquitin transfer. Journal of Biological Chemistry, 2009 , 284, 26797-802	5.4	34
60	NMR analysis of the architecture and functional remodeling of a modular multidomain protein, RPA. <i>Journal of the American Chemical Society</i> , 2009 , 131, 6346-7	16.4	38
59	Experimental model of the interplay between TBL1-mediated activation and Siah-1-induced poly-ubiquitination of Eatenin. <i>FASEB Journal</i> , 2009 , 23, 849.2	0.9	
58	Evolution of the NIGMS Protein Structure Initiative. Structure, 2008, 16, 12-4	5.2	1
57	Metal chelation and inhibition of bacterial growth in tissue abscesses. <i>Science</i> , 2008 , 319, 962-5	33.3	627
56	Cellular functions of human RPA1. Multiple roles of domains in replication, repair, and checkpoints. <i>Journal of Biological Chemistry</i> , 2008 , 283, 19095-111	5.4	84
55	Regulatory functions of the N-terminal domain of the 70-kDa subunit of replication protein A (RPA). <i>Journal of Biological Chemistry</i> , 2008 , 283, 21559-70	5.4	28
54	The basic cleft of RPA70N binds multiple checkpoint proteins, including RAD9, to regulate ATR signaling. <i>Molecular and Cellular Biology</i> , 2008 , 28, 7345-53	4.8	137
53	S100A8/A9 at low concentration promotes tumor cell growth via RAGE ligation and MAP kinase-dependent pathway. <i>Journal of Leukocyte Biology</i> , 2008 , 83, 1484-92	6.5	221
52	Replication protein A prevents accumulation of single-stranded telomeric DNA in cells that use alternative lengthening of telomeres. <i>Nucleic Acids Research</i> , 2007 , 35, 7267-78	20.1	49
51	An iron-sulfur cluster in the C-terminal domain of the p58 subunit of human DNA primase. <i>Journal of Biological Chemistry</i> , 2007 , 282, 33444-33451	5.4	95
50	The impact of X-ray crystallography and NMR on intracellular calcium signal transduction by EF-hand proteins: crossing the threshold from structure to biology and medicine. <i>Sciencens STKE:</i> Signal Transduction Knowledge Environment, 2007, 2007, pe27		3
49	The extracellular region of the receptor for advanced glycation end products is composed of two independent structural units. <i>Biochemistry</i> , 2007 , 46, 6957-70	3.2	134
48	Biochemical and structural domain analysis of xeroderma pigmentosum complementation group C protein. <i>Biochemistry</i> , 2006 , 45, 14965-79	3.2	60
47	The biochemical effect of Ser167 phosphorylation on Chlamydomonas reinhardtii centrin. <i>Biochemical and Biophysical Research Communications</i> , 2006 , 342, 342-8	3.4	14
46	Structural mechanism of RPA loading on DNA during activation of a simple pre-replication complex. <i>EMBO Journal</i> , 2006 , 25, 5516-26	13	61
45	Calcium-Dependent Regulation of Ion Channels 2006 , 1, 203-212		9

44	Insights into hRPA32 C-terminal domainmediated assembly of the simian virus 40 replisome. <i>Nature Structural and Molecular Biology</i> , 2005 , 12, 332-9	17.6	71
43	Structural mechanisms of DNA replication, repair, and recombination. <i>Journal of Biological Chemistry</i> , 2004 , 279, 30915-8	5.4	64
42	Target selectivity in EF-hand calcium binding proteins. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2004 , 1742, 69-79	4.9	188
41	Physical interaction between replication protein A and Rad51 promotes exchange on single-stranded DNA. <i>Journal of Biological Chemistry</i> , 2004 , 279, 25638-45	5.4	82
40	Replication protein A phosphorylation and the cellular response to DNA damage. <i>DNA Repair</i> , 2004 , 3, 1015-24	4.3	232
39	Independent and coordinated functions of replication protein A tandem high affinity single-stranded DNA binding domains. <i>Journal of Biological Chemistry</i> , 2003 , 278, 41077-82	5.4	97
38	The phosphorylation domain of the 32-kDa subunit of replication protein A (RPA) modulates RPA-DNA interactions. Evidence for an intersubunit interaction. <i>Journal of Biological Chemistry</i> , 2003 , 278, 35584-91	5.4	77
37	Structural insights into the U-box, a domain associated with multi-ubiquitination. <i>Nature Structural and Molecular Biology</i> , 2003 , 10, 250-5	17.6	216
36	Replication protein A interactions with DNA: differential binding of the core domains and analysis of the DNA interaction surface. <i>Biochemistry</i> , 2003 , 42, 12909-18	3.2	64
35	Nascent structure in the kinase anchoring domain of microtubule-associated protein 2. <i>Biochemical and Biophysical Research Communications</i> , 2003 , 301, 136-42	3.4	6
34	The EF-hand domain: a globally cooperative structural unit. <i>Protein Science</i> , 2002 , 11, 198-205	6.3	80
33	Characterization of binding-induced changes in dynamics suggests a model for sequence-nonspecific binding of ssDNA by replication protein A. <i>Protein Science</i> , 2002 , 11, 2316-25	6.3	15
32	Engineering and design of ligand-induced conformational change in proteins. <i>Current Opinion in Structural Biology</i> , 2002 , 12, 459-63	8.1	34
31	1H, 15N and 13C assignments of the regulatory domains of calcium-dependent protein kinase (CDPK). <i>Journal of Biomolecular NMR</i> , 2002 , 23, 249-50	3	5
30	Molecular basis for recognition and binding of specific DNA sequences by calicheamicin and duocarmycin. <i>Advances in DNA Sequence-Specific Agents</i> , 2002 , 4, 47-73		1
29	Analysis of the human replication protein A:Rad52 complex: evidence for crosstalk between RPA32, RPA70, Rad52 and DNA. <i>Journal of Molecular Biology</i> , 2002 , 321, 133-48	6.5	57
28	1H, 15N and 13C resonance assignments for the C-terminal protein interaction region of the 32 kDa subunit of human replication protein A. <i>Journal of Biomolecular NMR</i> , 2000 , 17, 179-80	3	4
27	DNA replication but not nucleotide excision repair is required for UVC-induced replication protein A phosphorylation in mammalian cells. <i>Molecular and Cellular Biology</i> , 2000 , 20, 2696-705	4.8	34

(1995-2000)

26	Structural basis for the recognition of DNA repair proteins UNG2, XPA, and RAD52 by replication factor RPA. <i>Cell</i> , 2000 , 103, 449-56	56.2	191
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