

# Haitao Li

## List of Publications by Year in Descending Order

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**Version:** 2024-04-27

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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.

133  
papers

10,318  
citations

47  
h-index

101  
g-index

142  
ext. papers

12,320  
ext. citations

15.4  
avg, IF

5.99  
L-index

#	Paper	IF	Citations
133	Residential green and blue space associated with lower risk of adult-onset inflammatory bowel disease: Findings from a large prospective cohort study.. <i>Environment International</i> , <b>2022</b> , 160, 107084	12.9	1
132	Risk/benefit tradeoff of habitual physical activity and air pollution on chronic pulmonary obstructive disease: findings from a large prospective cohort study.. <i>BMC Medicine</i> , <b>2022</b> , 20, 70	11.4	1
131	The PAF1 complex promotes 3Qprocessing of pervasive transcripts.. <i>Cell Reports</i> , <b>2022</b> , 38, 110519	10.6	0
130	Histidine methyltransferase SETD3 methylates structurally diverse histidine mimics in actin.. <i>Protein Science</i> , <b>2022</b> , 31, e4305	6.3	0
129	Chromatin profiling in human neurons reveals aberrant roles for histone acetylation and BET family proteins in schizophrenia.. <i>Nature Communications</i> , <b>2022</b> , 13, 2195	17.4	3
128	Cryo-EM structure of R-loop monoclonal antibody S9.6 in recognizing RNA:DNA hybrids.. <i>Journal of Genetics and Genomics</i> , <b>2022</b> ,	4	0
127	Understanding the phase separation characteristics of nucleocapsid protein provides a new therapeutic opportunity against SARS-CoV-2. <i>Protein and Cell</i> , <b>2021</b> , 12, 734-740	7.2	9
126	Histone Modifications and Chondrocyte Fate: Regulation and Therapeutic Implications. <i>Frontiers in Cell and Developmental Biology</i> , <b>2021</b> , 9, 626708	5.7	1
125	An integrative drug repositioning framework discovered a potential therapeutic agent targeting COVID-19. <i>Signal Transduction and Targeted Therapy</i> , <b>2021</b> , 6, 165	21	40
124	A conserved BAH module within mammalian BAHD1 connects H3K27me3 to Polycomb gene silencing. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, 4441-4455	20.1	5
123	PRMT1-mediated H4R3me2a recruits SMARCA4 to promote colorectal cancer progression by enhancing EGFR signaling. <i>Genome Medicine</i> , <b>2021</b> , 13, 58	14.4	9
122	Maternal heterozygous mutation in CHEK1 leads to mitotic arrest in human zygotes. <i>Protein and Cell</i> , <b>2021</b> , 1	7.2	1
121	Histone benzoylation serves as an epigenetic mark for DPF and YEATS family proteins. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, 114-126	20.1	12
120	Hypertension and Comorbidities in Rural and Urban Chinese Older People: An Epidemiological Subanalysis From the SAGE Study. <i>American Journal of Hypertension</i> , <b>2021</b> , 34, 183-189	2.3	0
119	Missing for partnership: understanding nucleosomal de novo DNA cytosine methylation by a spliced DNMT3 complex. <i>Signal Transduction and Targeted Therapy</i> , <b>2021</b> , 6, 36	21	0
118	Histone H3Q5 serotonylation stabilizes H3K4 methylation and potentiates its readout. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2021</b> , 118,	11.5	8
117	Comparison of the management and control of hypertension by public and private primary care providers in Shenzhen, China. <i>Heliyon</i> , <b>2021</b> , 7, e06280	3.6	0

116	Molecular basis for bipartite recognition of histone H3 by the PZP domain of PHF14. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, 8961-8973	20.1	1
115	Merging PROTAC and molecular glue for degrading BTK and GSPT1 proteins concurrently. <i>Cell Research</i> , <b>2021</b> , 31, 1315-1318	24.7	5
114	Investigate Natural Product Indolmycin and the Synthetically Improved Analogue Toward Antimycobacterial Agents.. <i>ACS Chemical Biology</i> , <b>2021</b> ,	4.9	1
113	Selective Targeting of AF9 YEATS Domain by Cyclopeptide Inhibitors with Preorganized Conformation. <i>Journal of the American Chemical Society</i> , <b>2020</b> , 142, 21450-21459	16.4	9
112	Mammalian ALKBH1 serves as an N-mA demethylase of unpairing DNA. <i>Cell Research</i> , <b>2020</b> , 30, 197-210	24.7	38
111	Molecular basis for histidine N3-specific methylation of actin H73 by SETD3. <i>Cell Discovery</i> , <b>2020</b> , 6, 3	22.3	8
110	Impaired cell fate through gain-of-function mutations in a chromatin reader. <i>Nature</i> , <b>2020</b> , 577, 121-126	50.4	36
109	T9. EPIGENETIC PROFILING IN SCHIZOPHRENIA DERIVED HUMAN INDUCED PLURIPOTENT STEM CELLS (HIPSCS) AND NEURONS. <i>Schizophrenia Bulletin</i> , <b>2020</b> , 46, S234-S234	1.3	78
108	Histone H3.3 phosphorylation amplifies stimulation-induced transcription. <i>Nature</i> , <b>2020</b> , 583, 852-857	50.4	43
107	N-methyladenine in DNA antagonizes SATB1 in early development. <i>Nature</i> , <b>2020</b> , 583, 625-630	50.4	23
106	Molecular basis for histone H3 "K4me3-K9me3/2" methylation pattern readout by Spindlin1. <i>Journal of Biological Chemistry</i> , <b>2020</b> , 295, 16877-16887	5.4	9
105	Social capital, depressive symptoms, and perceived quality of care among hypertensive patients in primary care. <i>Health and Quality of Life Outcomes</i> , <b>2020</b> , 18, 378	3	0
104	The histone mark H3K36me2 recruits DNMT3A and shapes the intergenic DNA methylation landscape. <i>Nature</i> , <b>2019</b> , 573, 281-286	50.4	161
103	PTEN Suppresses Glycolysis by Dephosphorylating and Inhibiting Autophosphorylated PGK1. <i>Molecular Cell</i> , <b>2019</b> , 76, 516-527.e7	17.6	55
102	Histone Modifications Regulate Chromatin Compartmentalization by Contributing to a Phase Separation Mechanism. <i>Molecular Cell</i> , <b>2019</b> , 76, 646-659.e6	17.6	135
101	Architecture of SAGA complex. <i>Cell Discovery</i> , <b>2019</b> , 5, 25	22.3	11
100	Identification and characterization of Readers for novel histone modifications. <i>Current Opinion in Chemical Biology</i> , <b>2019</b> , 51, 57-65	9.7	13
99	Nitrooxidoreductase Rv2466c-Dependent Fluorescent Probe for Mycobacterium tuberculosis Diagnosis and Drug Susceptibility Testing. <i>ACS Infectious Diseases</i> , <b>2019</b> , 5, 949-961	5.5	3

98	Histone serotonylation is a permissive modification that enhances TFIID binding to H3K4me3. <i>Nature</i> , <b>2019</b> , 567, 535-539	50.4	166
97	Cross-Sectional Study on the Management and Control of Hypertension Among Migrants in Primary Care: What Is the Impact of Segmented Health Insurance Schemes?. <i>Journal of the American Heart Association</i> , <b>2019</b> , 8, e012674	6	2
96	Molecular basis for hierarchical histone de-hydroxybutyrylation by SIRT3. <i>Cell Discovery</i> , <b>2019</b> , 5, 35	22.3	31
95	Inequalities in Structural Social Capital and Health between Migrant and Local Hypertensive Patients. <i>Annals of Global Health</i> , <b>2019</b> , 85,	3.3	4
94	Impact of local health insurance schemes on primary care management and control of hypertension: a cross-sectional study in Shenzhen, China. <i>BMJ Open</i> , <b>2019</b> , 9, e031098	3	5
93	Interplay between the bacterial protein deacetylase CobB and the second messenger c-di-GMP. <i>EMBO Journal</i> , <b>2019</b> , 38, e100948	13	15
92	Plasticity in designing PROTACs for selective and potent degradation of HDAC6. <i>Chemical Communications</i> , <b>2019</b> , 55, 14848-14851	5.8	36
91	CXCL17 promotes cell metastasis and inhibits autophagy via the LKB1-AMPK pathway in hepatocellular carcinoma. <i>Gene</i> , <b>2019</b> , 690, 129-136	3.8	21
90	Plant HP1 protein ADCP1 links multivalent H3K9 methylation readout to heterochromatin formation. <i>Cell Research</i> , <b>2019</b> , 29, 54-66	24.7	51
89	Inhibition of CPAP-tubulin interaction prevents proliferation of centrosome-amplified cancer cells. <i>EMBO Journal</i> , <b>2019</b> , 38,	13	12
88	Molecular basis for histidine N1 position-specific methylation by CARNMT1. <i>Cell Research</i> , <b>2018</b> , 28, 494-496	4.7	5
87	Identification of a Mycothiol-Dependent Nitroreductase from <i>Mycobacterium tuberculosis</i> . <i>ACS Infectious Diseases</i> , <b>2018</b> , 4, 771-787	5.5	12
86	Expanding RNA binding specificity and affinity of engineered PUF domains. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, 4771-4782	20.1	25
85	Systematic Profiling of Histone Readers in <i>Arabidopsis thaliana</i> . <i>Cell Reports</i> , <b>2018</b> , 22, 1090-1102	10.6	35
84	Recognition of histone acetylation by the GAS41 YEATS domain promotes H2A.Z deposition in non-small cell lung cancer. <i>Genes and Development</i> , <b>2018</b> , 32, 58-69	12.6	47
83	Social capital and depression among migrant hypertensive patients in primary care. <i>Journal of the American Society of Hypertension</i> , <b>2018</b> , 12, 621-626		7
82	Gas41 links histone acetylation to H2A.Z deposition and maintenance of embryonic stem cell identity. <i>Cell Discovery</i> , <b>2018</b> , 4, 28	22.3	27
81	Structure-Based Design of 6-Chloro-4-aminoquinazoline-2-carboxamide Derivatives as Potent and Selective p21-Activated Kinase 4 (PAK4) Inhibitors. <i>Journal of Medicinal Chemistry</i> , <b>2018</b> , 61, 265-285	8.3	28

80	Beyond histone acetylation-writing and erasing histone acylations. <i>Current Opinion in Structural Biology</i> , <b>2018</b> , 53, 169-177	8.1	75
79	Nucleolar localization signal and histone methylation reader function is required for SPIN1 to promote rRNA gene expression. <i>Biochemical and Biophysical Research Communications</i> , <b>2018</b> , 505, 325-332	3.4	5
78	Structural insights into the H3K36me3 stacking mechanism and DNA-binding activity of the YEATS domain. <i>Nature Communications</i> , <b>2018</b> , 9, 4574	17.4	24
77	Structure-guided development of YEATS domain inhibitors by targeting H3K36me3 stacking. <i>Nature Chemical Biology</i> , <b>2018</b> , 14, 1140-1149	11.7	48
76	Cancer-driving H3G34V/R/D mutations block H3K36 methylation and H3K36me3-MutS $\beta$ interaction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, 9598-9603	11.5	54
75	Design, synthesis, structure-activity relationships study and X-ray crystallography of 3-substituted-indolin-2-one-5-carboxamide derivatives as PAK4 inhibitors. <i>European Journal of Medicinal Chemistry</i> , <b>2018</b> , 155, 197-209	6.8	12
74	Histone H1 defect in escort cells triggers germline tumor in <i>Drosophila</i> ovary. <i>Developmental Biology</i> , <b>2017</b> , 424, 40-49	3.1	7
73	ENL links histone acetylation to oncogenic gene expression in acute myeloid leukaemia. <i>Nature</i> , <b>2017</b> , 543, 265-269	50.4	124
72	Developing Spindlin1 small-molecule inhibitors by using protein microarrays. <i>Nature Chemical Biology</i> , <b>2017</b> , 13, 750-756	11.7	35
71	A cross-sectional study of the prevalence, awareness, treatment and control of hypertension in Shenzhen, China. <i>BMJ Open</i> , <b>2017</b> , 7, e015206	3	8
70	YEATS Domain-A Histone Acylation Reader in Health and Disease. <i>Journal of Molecular Biology</i> , <b>2017</b> , 429, 1994-2002	6.5	50
69	YEATS2 links histone acetylation to tumorigenesis of non-small cell lung cancer. <i>Nature Communications</i> , <b>2017</b> , 8, 1088	17.4	56
68	Diabetes in Shenzhen, China: epidemiological investigation and health care challenges. <i>Journal of Global Health</i> , <b>2017</b> , 7, 011102	4.3	4
67	Kinetic and high-throughput profiling of epigenetic interactions by 3D-carbene chip-based surface plasmon resonance imaging technology. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2017</b> , 114, E7245-E7254	11.5	37
66	Structural and biochemical characterization of DAXX-ATRAX interaction. <i>Protein and Cell</i> , <b>2017</b> , 8, 762-766	7.2	2
65	YEATS domain: Linking histone crotonylation to gene regulation. <i>Transcription</i> , <b>2017</b> , 8, 9-14	4.8	26
64	Changes of the perceived quality of care for older patients with hypertension by community health centers in shanghai. <i>BMC Family Practice</i> , <b>2017</b> , 18, 114	2.6	1
63	Assessing the impact of general practitioner team service on perceived quality of care among patients with non-communicable diseases in China: a natural experimental study. <i>International Journal for Quality in Health Care</i> , <b>2016</b> , 28, 554-560	1.9	12

62	Selective recognition of histone crotonylation by double PHD fingers of MOZ and DPF2. <i>Nature Chemical Biology</i> , <b>2016</b> , 12, 1111-1118	11.7	90
61	Molecular basis for CPAP-tubulin interaction in controlling centriolar and ciliary length. <i>Nature Communications</i> , <b>2016</b> , 7, 11874	17.4	45
60	Can integrated health services delivery have an impact on hypertension management? A cross-sectional study in two cities of China. <i>International Journal for Equity in Health</i> , <b>2016</b> , 15, 193	4.6	1
59	Divergent lncRNAs Regulate Gene Expression and Lineage Differentiation in Pluripotent Cells. <i>Cell Stem Cell</i> , <b>2016</b> , 18, 637-52	18	255
58	The BAH domain of BAHD1 is a histone H3K27me3 reader. <i>Protein and Cell</i> , <b>2016</b> , 7, 222-6	7.2	28
57	Reading between the Lines: "ADD"-ing Histone and DNA Methylation Marks toward a New Epigenetic "Sum". <i>ACS Chemical Biology</i> , <b>2016</b> , 11, 554-63	4.9	17
56	JMJD5 (Jumonji Domain-containing 5) Associates with Spindle Microtubules and Is Required for Proper Mitosis. <i>Journal of Biological Chemistry</i> , <b>2016</b> , 291, 4684-97	5.4	12
55	YEATS2 is a selective histone crotonylation reader. <i>Cell Research</i> , <b>2016</b> , 26, 629-32	24.7	101
54	Molecular Coupling of Histone Crotonylation and Active Transcription by AF9 YEATS Domain. <i>Molecular Cell</i> , <b>2016</b> , 62, 181-193	17.6	184
53	Multifaceted Histone H3 Methylation and Phosphorylation Readout by the Plant Homeodomain Finger of Human Nuclear Antigen Sp100C. <i>Journal of Biological Chemistry</i> , <b>2016</b> , 291, 12786-12798	5.4	13
52	ZMYND8 Reads the Dual Histone Mark H3K4me1-H3K14ac to Antagonize the Expression of Metastasis-Linked Genes. <i>Molecular Cell</i> , <b>2016</b> , 63, 470-84	17.6	72
51	Molecular basis for oncohistone H3 recognition by SETD2 methyltransferase. <i>Genes and Development</i> , <b>2016</b> , 30, 1611-6	12.6	75
50	Changes in the perceived quality of primary care in Shanghai and Shenzhen, China: a difference-in-difference analysis. <i>Bulletin of the World Health Organization</i> , <b>2015</b> , 93, 407-16	8.2	29
49	A comparison of the quality of hypertension management in primary care between Shanghai and Shenzhen: a cohort study of 3196 patients. <i>Medicine (United States)</i> , <b>2015</b> , 94, e455	1.8	22
48	Molecular basis for histone N-terminal methylation by NRMT1. <i>Genes and Development</i> , <b>2015</b> , 29, 2337-42	22.6	18
47	Chemical basis for the recognition of trimethyllysine by epigenetic reader proteins. <i>Nature Communications</i> , <b>2015</b> , 6, 8911	17.4	57
46	Epidemiological investigation of suspected autism in children and implications for healthcare system: a mainstream kindergarten-based population study in Longhua District, Shenzhen. <i>BMC Pediatrics</i> , <b>2015</b> , 15, 207	2.6	9
45	What are the similarities and differences in structure and function among the three main models of community health centers in China: a systematic review. <i>BMC Health Services Research</i> , <b>2015</b> , 15, 504	2.9	11

44	Comparing quality of public primary care between Hong Kong and Shanghai using validated patient assessment tools. <i>PLoS ONE</i> , <b>2015</b> , 10, e0121269	3.7	30
43	Cross-talk between PRMT1-mediated methylation and ubiquitylation on RBM15 controls RNA splicing. <i>ELife</i> , <b>2015</b> , 4,	8.9	69
42	Engineering of a Histone-Recognition Domain in Dnmt3a Alters the Epigenetic Landscape and Phenotypic Features of Mouse ESCs. <i>Molecular Cell</i> , <b>2015</b> , 59, 89-103	17.6	56
41	Histone Recognition by Tandem Modules and Modulation by Multiple PTMs <b>2015</b> , 149-172		3
40	Crystallography-Based Mechanistic Insights into Epigenetic Regulation <b>2015</b> , 125-147		
39	A Cross-Sectional Comparison of Perceived Quality of Primary Care by Hypertensive Patients in Shanghai and Shenzhen, China. <i>Medicine (United States)</i> , <b>2015</b> , 94, e1388	1.8	11
38	ATRX tolerates activity-dependent histone H3 methyl/phos switching to maintain repetitive element silencing in neurons. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2015</b> , 112, 6820-7	11.5	38
37	ZMYND11 links histone H3.3K36me3 to transcription elongation and tumour suppression. <i>Nature</i> , <b>2014</b> , 508, 263-8	50.4	202
36	Molecular basis underlying histone H3 lysine-arginine methylation pattern readout by Spin/Ssty repeats of Spindlin1. <i>Genes and Development</i> , <b>2014</b> , 28, 622-36	12.6	78
35	Targeting epigenetic regulators for cancer therapy. <i>Annals of the New York Academy of Sciences</i> , <b>2014</b> , 1309, 30-6	6.5	64
34	AF9 YEATS domain links histone acetylation to DOT1L-mediated H3K79 methylation. <i>Cell</i> , <b>2014</b> , 159, 558-71	56.2	229
33	Comparison of perceived quality amongst migrant and local patients using primary health care delivered by community health centres in Shenzhen, China. <i>BMC Family Practice</i> , <b>2014</b> , 15, 76	2.6	33
32	One-pot native chemical ligation of peptide hydrazides enables total synthesis of modified histones. <i>Organic and Biomolecular Chemistry</i> , <b>2014</b> , 12, 5435-41	3.9	88
31	Conserved TCP domain of Sas-4/CPAP is essential for pericentriolar material tethering during centrosome biogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, E354-63	11.5	50
30	Acetylation-mediated proteasomal degradation of core histones during DNA repair and spermatogenesis. <i>Cell</i> , <b>2013</b> , 153, 1012-24	56.2	203
29	Elderly healthcare service at the community health centers in the Pearl River Delta region, China. <i>Family Medicine and Community Health</i> , <b>2013</b> , 1, 30-36	4.7	
28	ATRX-mediated chromatin association of histone variant macroH2A1 regulates $\beta$ globin expression. <i>Genes and Development</i> , <b>2012</b> , 26, 433-8	12.6	89
27	Many keys to push: diversifying the readership of plant homeodomain fingers. <i>Acta Biochimica Et Biophysica Sinica</i> , <b>2012</b> , 44, 28-39	2.8	47

26	Anterior Segment Optical Coherence Tomography and its Clinical Applications in Glaucoma. <i>Journal of Current Glaucoma Practice</i> , <b>2012</b> , 6, 68-74	1.1	9
25	ATRX ADD domain links an atypical histone methylation recognition mechanism to human mental-retardation syndrome. <i>Nature Structural and Molecular Biology</i> , <b>2011</b> , 18, 769-76	17.6	193
24	Recognition of a mononucleosomal histone modification pattern by BPTF via multivalent interactions. <i>Cell</i> , <b>2011</b> , 145, 692-706	56.2	261
23	PHD finger recognition of unmodified histone H3R2 links UHRF1 to regulation of euchromatic gene expression. <i>Molecular Cell</i> , <b>2011</b> , 43, 275-284	17.6	143
22	Structural and functional insights into 5Qppp RNA pattern recognition by the innate immune receptor RIG-I. <i>Nature Structural and Molecular Biology</i> , <b>2010</b> , 17, 781-7	17.6	196
21	Crystal structures of SULT1A2 and SULT1A1 *3: insights into the substrate inhibition and the role of Tyr149 in SULT1A2. <i>Biochemical and Biophysical Research Communications</i> , <b>2010</b> , 396, 429-34	3.4	15
20	Pro isomerization in MLL1 PHD3-bromo cassette connects H3K4me readout to Cyp33 and HDAC-mediated repression. <i>Cell</i> , <b>2010</b> , 141, 1183-94	56.2	145
19	WSTF regulates the H2A.X DNA damage response via a novel tyrosine kinase activity. <i>Nature</i> , <b>2009</b> , 457, 57-62	50.4	314
18	Haematopoietic malignancies caused by dysregulation of a chromatin-binding PHD finger. <i>Nature</i> , <b>2009</b> , 459, 847-51	50.4	330
17	Nucleation, propagation and cleavage of target RNAs in Ago silencing complexes. <i>Nature</i> , <b>2009</b> , 461, 754-61	50.4	388
16	PRMT5-mediated methylation of histone H4R3 recruits DNMT3A, coupling histone and DNA methylation in gene silencing. <i>Nature Structural and Molecular Biology</i> , <b>2009</b> , 16, 304-311	17.6	385
15	An alpha motif at Tas3 C terminus mediates RITS cis spreading and promotes heterochromatic gene silencing. <i>Molecular Cell</i> , <b>2009</b> , 34, 155-67	17.6	29
14	Structure of an argonaute silencing complex with a seed-containing guide DNA and target RNA duplex. <i>Nature</i> , <b>2008</b> , 456, 921-6	50.4	428
13	Multivalent engagement of chromatin modifications by linked binding modules. <i>Nature Reviews Molecular Cell Biology</i> , <b>2007</b> , 8, 983-94	48.7	823
12	How chromatin-binding modules interpret histone modifications: lessons from professional pocket pickers. <i>Nature Structural and Molecular Biology</i> , <b>2007</b> , 14, 1025-1040	17.6	1141
11	Repeatability and reproducibility of anterior chamber angle measurement with anterior segment optical coherence tomography. <i>British Journal of Ophthalmology</i> , <b>2007</b> , 91, 1490-2	5.5	84
10	Structural basis for lower lysine methylation state-specific readout by MBT repeats of L3MBTL1 and an engineered PHD finger. <i>Molecular Cell</i> , <b>2007</b> , 28, 677-91	17.6	163
9	Yng1 PHD finger binding to H3 trimethylated at K4 promotes NuA3 HAT activity at K14 of H3 and transcription at a subset of targeted ORFs. <i>Molecular Cell</i> , <b>2006</b> , 24, 785-796	17.6	255



8	Histone H3 recognition and presentation by the WDR5 module of the MLL1 complex. <i>Nature Structural and Molecular Biology</i> , <b>2006</b> , 13, 704-12	17.6	191
7	Molecular basis for site-specific read-out of histone H3K4me3 by the BPTF PHD finger of NURF. <i>Nature</i> , <b>2006</b> , 442, 91-5	50.4	600
6	Crystal structure of human sulfotransferase SULT1A3 in complex with dopamine and 3 $\alpha$ -phosphoadenosine 5 $\alpha$ -phosphate. <i>Biochemical and Biophysical Research Communications</i> , <b>2005</b> , 335, 417-23	3.4	46
5	Crystal structure of C-terminal desundecapeptide nitrite reductase from <i>Achromobacter cycloclastes</i> . <i>Biochemical and Biophysical Research Communications</i> , <b>2005</b> , 338, 1935-42	3.4	3
4	pH-profile crystal structure studies of C-terminal despentapeptide nitrite reductase from <i>Achromobacter cycloclastes</i> . <i>Biochemical and Biophysical Research Communications</i> , <b>2004</b> , 316, 107-13	3.4	2
3	Preliminary crystallographic studies of two C-terminally truncated copper-containing nitrite reductases from <i>Achromobacter cycloclastes</i> : changed crystallizing behaviors caused by residue deletion. <i>Biochemical and Biophysical Research Communications</i> , <b>2002</b> , 299, 173-6	3.4	2
2	Phosphorylation of the ancestral histone variant H3.3 amplifies stimulation-induced transcription		2
1	Molecular basis for SPINDOC-Spindlin1 engagement and its role in transcriptional inhibition		1