

# Tom W Muir

## List of Publications by Year in descending order

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233  
papers

26,422  
citations

5248

83  
h-index

6979

154  
g-index

255  
all docs

255  
docs citations

255  
times ranked

20922  
citing authors

#	ARTICLE	IF	CITATIONS
1	Synthesis of Oriented Hexasomes and Asymmetric Nucleosomes Using a Template Editing Process. <i>Journal of the American Chemical Society</i> , 2022, 144, 2284-2291.	6.6	5
2	A Genetically Encoded Approach for Breaking Chromatin Symmetry. <i>ACS Central Science</i> , 2022, 8, 176-183.	5.3	4
3	Histone H3Q5 seronylation stabilizes H3K4 methylation and potentiates its readout. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	27
4	Oncohistone mutations enhance chromatin remodeling and alter cell fates. <i>Nature Chemical Biology</i> , 2021, 17, 403-411.	3.9	50
5	Engineering of a Peptide Î±â€Nâ€Methyltransferase to Methylate Nonâ€Proteinogenic Amino Acids. <i>Angewandte Chemie</i> , 2021, 133, 14440-14444.	1.6	0
6	Engineering of a Peptide Î±â€Nâ€Methyltransferase to Methylate Nonâ€Proteinogenic Amino Acids. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 14319-14323.	7.2	10
7	Molecular Epigenetics: Chemical Biology Tools Come of Age. <i>Annual Review of Biochemistry</i> , 2021, 90, 287-320.	5.0	9
8	DOT1L complex regulates transcriptional initiation in human erythroleukemic cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	26
9	Janus Bioparticles: Asymmetric Nucleosomes and Their Preparation Using Chemical Biology Approaches. <i>Accounts of Chemical Research</i> , 2021, 54, 3215-3227.	7.6	8
10	Chromatin landscape signals differentially dictate the activities of mSWI/SNF family complexes. <i>Science</i> , 2021, 373, 306-315.	6.0	64
11	Synthesis of ADP-Ribosylated Histones Reveals Site-Specific Impacts on Chromatin Structure and Function. <i>Journal of the American Chemical Society</i> , 2021, 143, 10847-10852.	6.6	26
12	Reactive intermediates for interactome mapping. <i>Chemical Society Reviews</i> , 2021, 50, 2911-2926.	18.7	35
13	Chemical biology approaches to study histone interactors. <i>Biochemical Society Transactions</i> , 2021, 49, 2431-2441.	1.6	2
14	Impaired cell fate through gain-of-function mutations in a chromatin reader. <i>Nature</i> , 2020, 577, 121-126.	13.7	84
15	The Chaperone FACT and Histone H2B Ubiquitination Maintain S.Âpombe Genome Architecture through Genic and Subtelomeric Functions. <i>Molecular Cell</i> , 2020, 77, 501-513.e7.	4.5	32
16	A basic motif anchoring ISWI to nucleosome acidic patch regulates nucleosome spacing. <i>Nature Chemical Biology</i> , 2020, 16, 134-142.	3.9	51
17	Chemoenzymatic Semisynthesis of Proteins. <i>Chemical Reviews</i> , 2020, 120, 3051-3126.	23.0	142
18	Bisphosphoglycerate Mutase Deficiency Protects against Cerebral Malaria and Severe Malaria-Induced Anemia. <i>Cell Reports</i> , 2020, 32, 108170.	2.9	7

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19	Single-molecule and in silico dissection of the interaction between Polycomb repressive complex 2 and chromatin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 30465-30475.	3.3	41
20	The nucleosome acidic patch and H2A ubiquitination underlie mSWI/SNF recruitment in synovial sarcoma. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 836-845.	3.6	32
21	Discovery of quorum quenchers targeting the membrane-embedded sensor domain of the <i>Staphylococcus aureus</i> receptor histidine kinase, AgrC. <i>Chemical Communications</i> , 2020, 56, 11223-11226.	2.2	19
22	H2B ubiquitylation enhances H3K4 methylation activities of human KMT2 family complexes. <i>Nucleic Acids Research</i> , 2020, 48, 5442-5456.	6.5	29
23	Live-cell protein engineering with an ultra-short split intein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 12041-12049.	3.3	29
24	In situ chromatin interactomics using a chemical bait and trap approach. <i>Nature Chemistry</i> , 2020, 12, 520-527.	6.6	53
25	Chromatin as a key consumer in the metabolite economy. <i>Nature Chemical Biology</i> , 2020, 16, 620-629.	3.9	50
26	Proximity Induced Splicing Utilizing Caged Split Inteins. <i>Journal of the American Chemical Society</i> , 2019, 141, 13708-13712.	6.6	16
27	Determining the Spheroid Geometry of Individual Metallic Nanoparticles by Two-Dimensional Single-Particle Dynamic Light Scattering. <i>Journal of Physical Chemistry C</i> , 2019, 123, 18565-18572.	1.5	2
28	Protein engineering through tandem transamidation. <i>Nature Chemistry</i> , 2019, 11, 737-743.	6.6	32
29	Single-Particle Dynamic Light Scattering: Shapes of Individual Nanoparticles. <i>Nano Letters</i> , 2019, 19, 5530-5536.	4.5	21
30	PRC2 engages a bivalent H3K27M-H3K27me3 dinucleosome inhibitor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 22152-22157.	3.3	33
31	Nucleation and Propagation of Heterochromatin by the Histone Methyltransferase PRC2: Geometric Constraints and Impact of the Regulatory Subunit JARID2. <i>Journal of the American Chemical Society</i> , 2019, 141, 15029-15039.	6.6	16
32	PFA ependymoma-associated protein EZHIP inhibits PRC2 activity through a H3 K27M-like mechanism. <i>Nature Communications</i> , 2019, 10, 2146.	5.8	136
33	Identification of a DNA N6-Adenine Methyltransferase Complex and Its Impact on Chromatin Organization. <i>Cell</i> , 2019, 177, 1781-1796.e25.	13.5	81
34	The expanding landscape of ãconohistoneã™ mutations in human cancers. <i>Nature</i> , 2019, 567, 473-478.	13.7	271
35	Histone seronylation is a permissive modification that enhances TFIID binding to H3K4me3. <i>Nature</i> , 2019, 567, 535-539.	13.7	292
36	Barcoding mit Nukleinsãuren: Anwendung der DNAãSequenzierung als molekulares Zãhlwerk. <i>Angewandte Chemie</i> , 2019, 131, 4188-4207.	1.6	2

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37	Histone H3 tail binds a unique sensing pocket in EZH2 to activate the PRC2 methyltransferase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 8295-8300.	3.3	71
38	Nucleic Acid Barcoding Technologies: Converting DNA Sequencing into a Broad Spectrum Molecular Counter. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 4144-4162.	7.2	22
39	Identification of a Molecular Latch that Regulates Staphylococcal Virulence. <i>Cell Chemical Biology</i> , 2019, 26, 548-558.e4.	2.5	18
40	Recurrent SMARCB1 Mutations Reveal a Nucleosome Acidic Patch Interaction Site That Potentiates mSWI/SNF Complex Chromatin Remodeling. <i>Cell</i> , 2019, 179, 1342-1356.e23.	13.5	72
41	JASPer controls interphase histone H3S10 phosphorylation by chromosomal kinase JIL-1 in <i>Drosophila</i> . <i>Nature Communications</i> , 2019, 10, 5343.	5.8	18
42	Functional crosstalk between histone H2B ubiquitylation and H2A modifications and variants. <i>Nature Communications</i> , 2018, 9, 1394.	5.8	59
43	How many human proteoforms are there?. <i>Nature Chemical Biology</i> , 2018, 14, 206-214.	3.9	580
44	HELLS and CDCA7 comprise a bipartite nucleosome remodeling complex defective in ICF syndrome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E876-E885.	3.3	88
45	A Chemical Probe for Protein Crotonylation. <i>Journal of the American Chemical Society</i> , 2018, 140, 4757-4760.	6.6	44
46	Improved protein splicing using embedded split inteins. <i>Protein Science</i> , 2018, 27, 614-619.	3.1	15
47	Crosstalk among Set1 complex subunits involved in H2B ubiquitylation-dependent H3K4 methylation. <i>Nucleic Acids Research</i> , 2018, 46, 11129-11143.	6.5	19
48	An Atypical Mechanism of Split Intein Molecular Recognition and Folding. <i>Journal of the American Chemical Society</i> , 2018, 140, 11791-11799.	6.6	15
49	Acetylation blocks DNA damage-induced chromatin ADP-ribosylation. <i>Nature Chemical Biology</i> , 2018, 14, 837-840.	3.9	66
50	Functional Plasticity of the AgrC Receptor Histidine Kinase Required for Staphylococcal Virulence. <i>Cell Chemical Biology</i> , 2017, 24, 76-86.	2.5	25
51	Genomic targeting of epigenetic probes using a chemically tailored Cas9 system. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 681-686.	3.3	53
52	A molecular engineering toolbox for the structural biologist. <i>Quarterly Reviews of Biophysics</i> , 2017, 50, e7.	2.4	42
53	Emerging Chemistry Strategies for Engineering Native Chromatin. <i>Journal of the American Chemical Society</i> , 2017, 139, 9090-9096.	6.6	25
54	Intein Zymogens: Conditional Assembly and Splicing of Split Inteins via Targeted Proteolysis. <i>Journal of the American Chemical Society</i> , 2017, 139, 8074-8077.	6.6	39

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55	Surface-attached molecules control <i>Staphylococcus aureus</i> quorum sensing and biofilm development. <i>Nature Microbiology</i> , 2017, 2, 17080.	5.9	95
56	Molecular analysis of PRC2 recruitment to DNA in chromatin and its inhibition by RNA. <i>Nature Structural and Molecular Biology</i> , 2017, 24, 1028-1038.	3.6	186
57	A promiscuous split intein with expanded protein engineering applications. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 8538-8543.	3.3	102
58	ISWI chromatin remodellers sense nucleosome modifications to determine substrate preference. <i>Nature</i> , 2017, 548, 607-611.	13.7	148
59	Long-term hepatitis B infection in a scalable hepatic co-culture system. <i>Nature Communications</i> , 2017, 8, 125.	5.8	58
60	Bisphosphoglycerate mutase controls serine pathway flux via 3-phosphoglycerate. <i>Nature Chemical Biology</i> , 2017, 13, 1081-1087.	3.9	47
61	Ubiquitin utilizes an acidic surface patch to alter chromatin structure. <i>Nature Chemical Biology</i> , 2017, 13, 105-110.	3.9	79
62	Evidence that ubiquitylated H2B corrals hDot1L on the nucleosomal surface to induce H3K79 methylation. <i>Nature Communications</i> , 2016, 7, 10589.	5.8	59
63	Histone H3K36 mutations promote sarcomagenesis through altered histone methylation landscape. <i>Science</i> , 2016, 352, 844-849.	6.0	327
64	A two-state activation mechanism controls the histone methyltransferase Suv39h1. <i>Nature Chemical Biology</i> , 2016, 12, 188-193.	3.9	90
65	Design of a Split Intein with Exceptional Protein Splicing Activity. <i>Journal of the American Chemical Society</i> , 2016, 138, 2162-2165.	6.6	133
66	Regulation of Virulence in <i>Staphylococcus aureus</i> : Molecular Mechanisms and Remaining Puzzles. <i>Cell Chemical Biology</i> , 2016, 23, 214-224.	2.5	166
67	Targeted Histone Peptides: Insights into the Spatial Regulation of the Methyltransferase PRC2 by using a Surrogate of Heterotypic Chromatin. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 6457-6461.	7.2	16
68	A Second-Generation Phosphohistidine Analog for Production of Phosphohistidine Antibodies. <i>Organic Letters</i> , 2015, 17, 187-189.	2.4	42
69	Increasing AIP Macrocyclic Size Reveals Key Features of <i>agr</i> Activation in <i>Staphylococcus aureus</i> . <i>ChemBioChem</i> , 2015, 16, 1093-1100.	1.3	32
70	Application of the Protein Semisynthesis Strategy to the Generation of Modified Chromatin. <i>Annual Review of Biochemistry</i> , 2015, 84, 265-290.	5.0	60
71	Chemical tagging and customizing of cellular chromatin states using ultrafast trans-splicing inteins. <i>Nature Chemistry</i> , 2015, 7, 394-402.	6.6	133
72	Key driving forces in the biosynthesis of autoinducing peptides required for staphylococcal virulence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 10679-10684.	3.3	34

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73	DNA-guided establishment of nucleosome patterns within coding regions of a eukaryotic genome. <i>Genome Research</i> , 2015, 25, 1727-1738.	2.4	26
74	Identification of a functional hotspot on ubiquitin required for stimulation of methyltransferase activity on chromatin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 10365-10370.	3.3	44
75	Histones: At the Crossroads of Peptide and Protein Chemistry. <i>Chemical Reviews</i> , 2015, 115, 2296-2349.	23.0	188
76	Structural and dynamical features of inteins and implications on protein splicing. <i>Journal of Biological Chemistry</i> , 2014, 289, 19278.	1.6	1
77	Inteins: nature's gift to protein chemists. <i>Chemical Science</i> , 2014, 5, 446-461.	3.7	310
78	A Phosphohistidine Proteomics Strategy Based on Elucidation of a Unique Gas-Phase Phosphopeptide Fragmentation Mechanism. <i>Journal of the American Chemical Society</i> , 2014, 136, 12899-12911.	6.6	64
79	Structural and Dynamical Features of Inteins and Implications on Protein Splicing. <i>Journal of Biological Chemistry</i> , 2014, 289, 14506-14511.	1.6	55
80	Strategy for "Detoxification" of a Cancer-Derived Histone Mutant Based on Mapping Its Interaction with the Methyltransferase PRC2. <i>Journal of the American Chemical Society</i> , 2014, 136, 13498-13501.	6.6	95
81	Accelerated chromatin biochemistry using DNA-barcoded nucleosome libraries. <i>Nature Methods</i> , 2014, 11, 834-840.	9.0	129
82	Activation and Inhibition of the Receptor Histidine Kinase AgrC Occurs through Opposite Helical Transduction Motions. <i>Molecular Cell</i> , 2014, 53, 929-940.	4.5	110
83	Structure of the branched intermediate in protein splicing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 8422-8427.	3.3	26
84	EXPLORING CHROMATIN BIOLOGY USING PROTEIN CHEMISTRY. , 2014, , .		0
85	SET1 and p300 Act Synergistically, through Coupled Histone Modifications, in Transcriptional Activation by p53. <i>Cell</i> , 2013, 154, 297-310.	13.5	147
86	Inhibition of PRC2 Activity by a Gain-of-Function H3 Mutation Found in Pediatric Glioblastoma. <i>Science</i> , 2013, 340, 857-861.	6.0	1,074
87	H3R42me2a is a histone modification with positive transcriptional effects. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 14894-14899.	3.3	115
88	The n-SET Domain of Set1 Regulates H2B Ubiquitylation-Dependent H3K4 Methylation. <i>Molecular Cell</i> , 2013, 49, 1121-1133.	4.5	119
89	Extein Residues Play an Intimate Role in the Rate-Limiting Step of Protein <i>Trans</i> -Splicing. <i>Journal of the American Chemical Society</i> , 2013, 135, 5839-5847.	6.6	63
90	ASH2L Regulates Ubiquitylation Signaling to MLL: trans-Regulation of H3 K4 Methylation in Higher Eukaryotes. <i>Molecular Cell</i> , 2013, 49, 1108-1120.	4.5	69

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91	Streamlined Expressed Protein Ligation Using Split Inteins. <i>Journal of the American Chemical Society</i> , 2013, 135, 286-292.	6.6	90
92	Induction of innate and adaptive immunity by delivery of poly dA:dT to dendritic cells. <i>Nature Chemical Biology</i> , 2013, 9, 250-256.	3.9	30
93	A pan-specific antibody for direct detection of protein histidine phosphorylation. <i>Nature Chemical Biology</i> , 2013, 9, 416-421.	3.9	119
94	Naturally Split Inteins Assemble through a "Capture and Collapse" Mechanism. <i>Journal of the American Chemical Society</i> , 2013, 135, 18673-18681.	6.6	63
95	Histone H3K27 Trimethylation Inhibits H3 Binding and Function of SET1-Like H3K4 Methyltransferase Complexes. <i>Molecular and Cellular Biology</i> , 2013, 33, 4936-4946.	1.1	61
96	From EPOthilone to EPO: A challenge for natural product synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 7134-7135.	3.3	3
97	Identification of ligand specificity determinants in AgrC, the <i>Staphylococcus aureus</i> quorum-sensing receptor.. <i>Journal of Biological Chemistry</i> , 2012, 287, 18588.	1.6	1
98	Histone Monoubiquitylation Position Determines Specificity and Direction of Enzymatic Cross-talk with Histone Methyltransferases Dot1L and PRC2. <i>Journal of Biological Chemistry</i> , 2012, 287, 23718-23725.	1.6	32
99	Stability of Nucleosomes Containing Homogenously Ubiquitylated H2A and H2B Prepared Using Semisynthesis. <i>Journal of the American Chemical Society</i> , 2012, 134, 19548-19551.	6.6	83
100	Chasing Phosphohistidine, an Elusive Sibling in the Phosphoamino Acid Family. <i>ACS Chemical Biology</i> , 2012, 7, 44-51.	1.6	107
101	Ultrafast Protein Splicing is Common among Cyanobacterial Split Inteins: Implications for Protein Engineering. <i>Journal of the American Chemical Society</i> , 2012, 134, 11338-11341.	6.6	122
102	Chromatin as an expansive canvas for chemical biology. <i>Nature Chemical Biology</i> , 2012, 8, 417-427.	3.9	109
103	Direct Interaction between an Allosteric Agonist Pepducin and the Chemokine Receptor CXCR4. <i>Journal of the American Chemical Society</i> , 2011, 133, 15878-15881.	6.6	64
104	Genetically Encoded 1,2-Aminothiols Facilitate Rapid and Site-Specific Protein Labeling via a Bio-orthogonal Cyanobenzothiazole Condensation. <i>Journal of the American Chemical Society</i> , 2011, 133, 11418-11421.	6.6	144
105	Recognition of a Mononucleosomal Histone Modification Pattern by BPTF via Multivalent Interactions. <i>Cell</i> , 2011, 145, 692-706.	13.5	300
106	Histone H2B ubiquitylation disrupts local and higher-order chromatin compaction. <i>Nature Chemical Biology</i> , 2011, 7, 113-119.	3.9	392
107	Tuning protein autoinhibition by domain destabilization. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 550-555.	3.6	30
108	Split Inteins: Nature's Protein Ligases. <i>Israel Journal of Chemistry</i> , 2011, 51, 854-861.	1.0	31

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109	Kinetic Control of Oneâ€Pot Transâ€Splicing Reactions by Using a Wildâ€Type and Designed Split Intein. <i>Angewandte Chemie - International Edition</i> , 2011, 50, 6511-6515.	7.2	61
110	Spreading Chromatin into Chemical Biology. <i>ChemBioChem</i> , 2011, 12, 264-279.	1.3	67
111	A Semisynthetic Eph Receptor Tyrosine Kinase Provides Insight into Ligand- Induced Kinase Activation. <i>Chemistry and Biology</i> , 2011, 18, 361-371.	6.2	30
112	Histone H2A deubiquitinase activity of the Polycomb repressive complex PR-DUB. <i>Nature</i> , 2010, 465, 243-247.	13.7	674
113	Disulfide-directed histone ubiquitylation reveals plasticity in hDot1L activation. <i>Nature Chemical Biology</i> , 2010, 6, 267-269.	3.9	227
114	Branched intermediate formation stimulates peptide bond cleavage in protein splicing. <i>Nature Chemical Biology</i> , 2010, 6, 527-533.	3.9	62
115	Chemical Approaches for Studying Histone Modifications. <i>Journal of Biological Chemistry</i> , 2010, 285, 11045-11050.	1.6	85
116	Development of Stable Phosphohistidine Analogues. <i>Journal of the American Chemical Society</i> , 2010, 132, 14327-14329.	6.6	126
117	Biological Applications of Protein Splicing. <i>Cell</i> , 2010, 143, 191-200.	13.5	152
118	Method for the Synthesis of Mono-ADP-ribose Conjugated Peptides. <i>Journal of the American Chemical Society</i> , 2010, 132, 15878-15880.	6.6	52
119	Traceless protein splicing utilizing evolved split inteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 10999-11004.	3.3	100
120	<i>agr</i> receptor mutants reveal distinct modes of inhibition by staphylococcal autoinducing peptides. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 1216-1221.	3.3	129
121	Chapter 11 Semisynthesis of Ubiquitylated Proteins. <i>Methods in Enzymology</i> , 2009, 462, 225-243.	0.4	24
122	Preface. <i>Methods in Enzymology</i> , 2009, 462, xiii-xv.	0.4	1
123	A Semisynthetic Strategy to Generate Phosphorylated and Acetylated Histone H2B. <i>ChemBioChem</i> , 2009, 10, 2182-2187.	1.3	59
124	Symmetric signalling within asymmetric dimers of the <i>Staphylococcus aureus</i> receptor histidine kinase AgrC. <i>Molecular Microbiology</i> , 2009, 74, 44-57.	1.2	60
125	Direct Measurement of Cathepsin B Activity in the Cytosol of Apoptotic Cells by an Activity-Based Probe. <i>Chemistry and Biology</i> , 2009, 16, 1001-1012.	6.2	36
126	Chemical Biology in a Time of Transition. <i>ACS Chemical Biology</i> , 2009, 4, 241-243.	1.6	3



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127	Structure-Activity Analysis of Semisynthetic Nucleosomes: Mechanistic Insights into the Stimulation of Dot1L by Ubiquitylated Histone H2B. <i>ACS Chemical Biology</i> , 2009, 4, 958-968.	1.6	109
128	RAD6-Mediated Transcription-Coupled H2B Ubiquitylation Directly Stimulates H3K4 Methylation in Human Cells. <i>Cell</i> , 2009, 137, 459-471.	13.5	453
129	PET Imaging of Leptin Biodistribution and Metabolism in Rodents and Primates. <i>Cell Metabolism</i> , 2009, 10, 148-159.	7.2	52
130	Expressed Protein Ligation (EPL) in the Study of Signal Transduction, Ion Conduction, And Chromatin Biology. <i>Accounts of Chemical Research</i> , 2009, 42, 107-116.	7.6	110
131	Anthrax Lethal Toxin Induced Lysosomal Membrane Permeabilization and Cytosolic Cathepsin Release Is Nlrp1b/Nalp1b-Dependent. <i>PLoS ONE</i> , 2009, 4, e7913.	1.1	53
132	An amalgamation of solid phase peptide synthesis and ribosomal peptide synthesis. <i>Biopolymers</i> , 2008, 90, 406-414.	1.2	14
133	Studying protein structure and function using semisynthesis. <i>Biopolymers</i> , 2008, 90, 743-750.	1.2	22
134	Activation of Protein Splicing by Protease- or Light-Triggered O to N Acyl Migration. <i>Angewandte Chemie - International Edition</i> , 2008, 47, 7764-7767.	7.2	69
135	A Full-Length Group 1 Bacterial Sigma Factor Adopts a Compact Structure Incompatible with DNA Binding. <i>Chemistry and Biology</i> , 2008, 15, 1091-1103.	6.2	51
136	Chemically ubiquitylated histone H2B stimulates hDot1L-mediated intranucleosomal methylation. <i>Nature</i> , 2008, 453, 812-816.	13.7	494
137	Activation of protein splicing with light in yeast. <i>Nature Methods</i> , 2008, 5, 303-305.	9.0	140
138	Toward the semisynthesis of multidomain transmembrane receptors: Modification of Eph tyrosine kinases. <i>Protein Science</i> , 2008, 17, 1740-1747.	3.1	13
139	Cyclic Peptide Inhibitors of Staphylococcal Virulence Prepared by Fmoc-Based Thiolactone Peptide Synthesis. <i>Journal of the American Chemical Society</i> , 2008, 130, 4914-4924.	6.6	127
140	Site-Specific <sup>18</sup> F-Labeling of the Protein Hormone Leptin Using a General Two-Step Ligation Procedure. <i>Journal of the American Chemical Society</i> , 2008, 130, 9106-9112.	6.6	67
141	Identification of Ligand Specificity Determinants in AgrC, the <i>Staphylococcus aureus</i> Quorum-sensing Receptor. <i>Journal of Biological Chemistry</i> , 2008, 283, 8930-8938.	1.6	88
142	Systems- and Molecular-Level Elucidation of Signaling Processes Through Chemistry. <i>Science Signaling</i> , 2008, 1, pe45.	1.6	2
143	Small-molecule-mediated rescue of protein function by an inducible proteolytic shunt. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 11209-11214.	3.3	88
144	Covalent Capture of Phospho-Dependent Protein Oligomerization by Site-Specific Incorporation of a Diazirine Photo-Cross-Linker. <i>Journal of the American Chemical Society</i> , 2007, 129, 8068-8069.	6.6	47

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145	Molecular Mechanisms of $\gamma$ Quorum Sensing in Virulent Staphylococci. <i>ChemBioChem</i> , 2007, 8, 847-855.	1.3	136
146	Tunable Photoactivation of a Post-translationally Modified Signaling Protein and its Unmodified Counterpart in Live Cells. <i>ChemBioChem</i> , 2007, 8, 2100-2105.	1.3	21
147	Auxiliary-Mediated Site-Specific Peptide Ubiquitylation. <i>Angewandte Chemie - International Edition</i> , 2007, 46, 2814-2818.	7.2	168
148	Post-translational enzyme activation in an animal via optimized conditional protein splicing. , 2007, 3, 50-54.		79
149	Ion Selectivity in a Semisynthetic K <sup>+</sup> Channel Locked in the Conductive Conformation. <i>Science</i> , 2006, 314, 1004-1007.	6.0	124
150	Structural and Functional Consequences of an Amide-to-Ester Substitution in the Selectivity Filter of a Potassium Channel. <i>Journal of the American Chemical Society</i> , 2006, 128, 11591-11599.	6.6	53
151	Solution Structure and Folding Characteristics of the C-Terminal SH3 Domain of c-Crk-II,. <i>Biochemistry</i> , 2006, 45, 8874-8884.	1.2	40
152	Cutting out the middle man. <i>Nature</i> , 2006, 442, 517-518.	13.7	4
153	Protein ligation: an enabling technology for the biophysical analysis of proteins. <i>Nature Methods</i> , 2006, 3, 429-438.	9.0	351
154	Semisynthetic proteins in mechanistic studies: using chemistry to go where nature can't. <i>Current Opinion in Chemical Biology</i> , 2006, 10, 487-491.	2.8	51
155	Biopolymers. <i>Current Opinion in Chemical Biology</i> , 2005, 9, 545-547.	2.8	0
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