

Ronald Hay

List of Publications by Year in descending order

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221
papers

25,671
citations

5896

81
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7160

153
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234
all docs

234
docs citations

234
times ranked

21160
citing authors

#	ARTICLE	IF	CITATIONS
1	SUMO. <i>Molecular Cell</i> , 2005, 18, 1-12.	9.7	1,468
2	SUMO-1 Modification of I κ B Inhibits NF- κ B Activation. <i>Molecular Cell</i> , 1998, 2, 233-239.	9.7	982
3	Thiordoxin regulates the DNA binding activity of NF- κ B by reduction of a disulphid bond involving cysteine 62. <i>Nucleic Acids Research</i> , 1992, 20, 3821-3830.	14.5	791
4	RNF4 is a poly-SUMO-specific E3 ubiquitin ligase required for arsenic-induced PML degradation. <i>Nature Cell Biology</i> , 2008, 10, 538-546.	10.3	746
5	Polymeric Chains of SUMO-2 and SUMO-3 Are Conjugated to Protein Substrates by SAE1/SAE2 and Ubc9. <i>Journal of Biological Chemistry</i> , 2001, 276, 35368-35374.	3.4	690
6	SUMO-1 Conjugation in Vivo Requires Both a Consensus Modification Motif and Nuclear Targeting. <i>Journal of Biological Chemistry</i> , 2001, 276, 12654-12659.	3.4	650
7	SUMO-1 modification activates the transcriptional response of p53. <i>EMBO Journal</i> , 1999, 18, 6455-6461.	7.8	602
8	Inhibition of NF- κ B DNA Binding by Nitric Oxide. <i>Nucleic Acids Research</i> , 1996, 24, 2236-2242.	14.5	500
9	OMERO: flexible, model-driven data management for experimental biology. <i>Nature Methods</i> , 2012, 9, 245-253.	19.0	478
10	Mdm2-Mediated NEDD8 Conjugation of p53 Inhibits Its Transcriptional Activity. <i>Cell</i> , 2004, 118, 83-97.	28.9	477
11	Structure of a RING E3 ligase and ubiquitin-loaded E2 primed for catalysis. <i>Nature</i> , 2012, 489, 115-120.	27.8	437
12	System-Wide Changes to SUMO Modifications in Response to Heat Shock. <i>Science Signaling</i> , 2009, 2, ra24.	3.6	415
13	p300 Transcriptional Repression Is Mediated by SUMO Modification. <i>Molecular Cell</i> , 2003, 11, 1043-1054.	9.7	406
14	Family-wide analysis of poly(ADP-ribose) polymerase activity. <i>Nature Communications</i> , 2014, 5, 4426.	12.8	386
15	Multiple C-Terminal Lysine Residues Target p53 for Ubiquitin-Proteasome-Mediated Degradation. <i>Molecular and Cellular Biology</i> , 2000, 20, 8458-8467.	2.3	337
16	Mutations in the I κ B α gene in Hodgkin's disease suggest a tumour suppressor role for I κ B α . <i>Oncogene</i> , 1999, 18, 3063-3070.	5.9	330
17	Ubc9 conjugates SUMO but not ubiquitin. <i>FEBS Letters</i> , 1997, 417, 297-300.	2.8	314
18	Identification of the Enzyme Required for Activation of the Small Ubiquitin-like Protein SUMO-1. <i>Journal of Biological Chemistry</i> , 1999, 274, 10618-10624.	3.4	306

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19	SIRT1 Deacetylation and Repression of p300 Involves Lysine Residues 1020/1024 within the Cell Cycle Regulatory Domain 1. <i>Journal of Biological Chemistry</i> , 2005, 280, 10264-10276.	3.4	301
20	Initiation of SV40 DNA replication in vivo: Location and structure of 5' ends of DNA synthesized in the ori region. <i>Cell</i> , 1982, 28, 767-779.	28.9	294
21	Distinct and Overlapping Sets of SUMO-1 and SUMO-2 Target Proteins Revealed by Quantitative Proteomics. <i>Molecular and Cellular Proteomics</i> , 2006, 5, 2298-2310.	3.8	274
22	Calpain 3 deficiency is associated with myonuclear apoptosis and profound perturbation of the I κ B/NF- κ B pathway in limb-girdle muscular dystrophy type 2A. <i>Nature Medicine</i> , 1999, 5, 503-511.	30.7	261
23	Posttranslational hydroxylation of ankyrin repeats in I κ B proteins by the hypoxia-inducible factor (HIF) asparaginyl hydroxylase, factor inhibiting HIF (FIH). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 14767-14772.	7.1	258
24	SUMO-specific proteases: a twist in the tail. <i>Trends in Cell Biology</i> , 2007, 17, 370-376.	7.9	253
25	In Vivo Identification of Human Small Ubiquitin-like Modifier Polymerization Sites by High Accuracy Mass Spectrometry and an in Vitro to in Vivo Strategy. <i>Molecular and Cellular Proteomics</i> , 2008, 7, 132-144.	3.8	251
26	An N-terminal p14ARF peptide blocks Mdm2-dependent ubiquitination in vitro and can activate p53 in vivo. <i>Oncogene</i> , 2000, 19, 2312-2323.	5.9	240
27	Dynamic Interplay of the SUMO and ERK Pathways in Regulating Elk-1 Transcriptional Activity. <i>Molecular Cell</i> , 2003, 12, 63-74.	9.7	231
28	An additional role for SUMO in ubiquitin-mediated proteolysis. <i>Nature Reviews Molecular Cell Biology</i> , 2009, 10, 564-568.	37.0	231
29	Nuclear Retention of I κ B α Protects It from Signal-induced Degradation and Inhibits Nuclear Factor κ B Transcriptional Activation. <i>Journal of Biological Chemistry</i> , 1999, 274, 9108-9115.	3.4	221
30	Single-stranded DNA-binding protein hSSB1 is critical for genomic stability. <i>Nature</i> , 2008, 453, 677-681.	27.8	220
31	zeta PKC induces phosphorylation and inactivation of I kappa B-alpha in vitro.. <i>EMBO Journal</i> , 1994, 13, 2842-2848.	7.8	216
32	Absolute dependence on kappa B responsive elements for initiation and Tat-mediated amplification of HIV transcription in blood CD4 T lymphocytes.. <i>EMBO Journal</i> , 1995, 14, 1552-1560.	7.8	214
33	SUMO-targeted ubiquitin E3 ligase RNF4 is required for the response of human cells to DNA damage. <i>Genes and Development</i> , 2012, 26, 1196-1208.	5.9	209
34	Role of I B α Ubiquitination in Signal-induced Activation of NF- B in Vivo. <i>Journal of Biological Chemistry</i> , 1996, 271, 7844-7850.	3.4	206
35	A Proteomic Study of SUMO-2 Target Proteins. <i>Journal of Biological Chemistry</i> , 2004, 279, 33791-33798.	3.4	197
36	Proteolytic degradation of MAD3 (I κ B α) and enhanced processing of the NF- κ B precursor p105 are obligatory steps in the activation of NF- κ B. <i>Nucleic Acids Research</i> , 1993, 21, 5059-5066.	14.5	196

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37	p75-Mediated NF- κ B Activation Enhances the Survival Response of Developing Sensory Neurons to Nerve Growth Factor. <i>Molecular and Cellular Neurosciences</i> , 1999, 14, 28-40.	2.2	177
38	The adenovirus protease is activated by a virus-coded disulphide-linked peptide. <i>Cell</i> , 1993, 72, 97-104.	28.9	175
39	Activation of NF- κ B by PM10 Occurs via an Iron-Mediated Mechanism in the Absence of κ B Degradation. <i>Toxicology and Applied Pharmacology</i> , 2000, 166, 101-110.	2.8	174
40	Proteome-Wide Identification of SUMO2 Modification Sites. <i>Science Signaling</i> , 2014, 7, rs2.	3.6	174
41	NEDP1, a Highly Conserved Cysteine Protease That deNEDDylates Cullins. <i>Journal of Biological Chemistry</i> , 2003, 278, 25637-25643.	3.4	170
42	A Mechanism for Inhibiting the SUMO Pathway. <i>Molecular Cell</i> , 2004, 16, 549-561.	9.7	164
43	SUMO and transcriptional regulation. <i>Seminars in Cell and Developmental Biology</i> , 2004, 15, 201-210.	5.0	158
44	Mechanism of ubiquitylation by dimeric RING ligase RNF4. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 1052-1059.	8.2	157
45	Androgen Receptor Acetylation Governs trans Activation and MEKK1-Induced Apoptosis without Affecting In Vitro Sumoylation and trans -Repression Function. <i>Molecular and Cellular Biology</i> , 2002, 22, 3373-3388.	2.3	155
46	Purification and identification of endogenous polySUMO conjugates. <i>EMBO Reports</i> , 2011, 12, 142-148.	4.5	155
47	Comparative Proteomic Analysis Identifies a Role for SUMO in Protein Quality Control. <i>Science Signaling</i> , 2011, 4, rs4.	3.6	153
48	SUMOylation of the GTPase Rac1 is required for optimal cell migration. <i>Nature Cell Biology</i> , 2010, 12, 1078-1085.	10.3	149
49	Role of cysteine62 in DNA recognition by the P50 subunit of NF- κ B. <i>Nucleic Acids Research</i> , 1993, 21, 1727-1734.	14.5	147
50	Ribosomal proteins are targets for the NEDD8 pathway. <i>EMBO Reports</i> , 2008, 9, 280-286.	4.5	147
51	Detection of protein SUMOylation in vivo. <i>Nature Protocols</i> , 2009, 4, 1363-1371.	12.0	144
52	HSP90 Protein Stabilizes Unloaded Argonaute Complexes and Microscopic P-bodies in Human Cells. <i>Molecular Biology of the Cell</i> , 2010, 21, 1462-1469.	2.1	143
53	Cytokine-Induced Nuclear Factor Kappa B Activation Promotes the Survival of Developing Neurons. <i>Journal of Cell Biology</i> , 2000, 148, 325-332.	5.2	141
54	Protein modification by SUMO. <i>Trends in Biochemical Sciences</i> , 2001, 26, 332-333.	7.5	140

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55	Persistent activation of nuclear factor- κ B in cultured rat hepatic stellate cells involves the induction of potentially novel rel-like factors and prolonged changes in the expression of I κ B family proteins. <i>Hepatology</i> , 1999, 30, 761-769.	7.3	131
56	Defective I κ B α in Hodgkin cell lines with constitutively active NF- κ B. <i>Oncogene</i> , 1998, 16, 2131-2139.	5.9	130
57	Unique binding interactions among Ubc9, SUMO and RanBP2 reveal a mechanism for SUMO paralog selection. <i>Nature Structural and Molecular Biology</i> , 2005, 12, 67-74.	8.2	125
58	The structure of SENP1 α SUMO-2 complex suggests a structural basis for discrimination between SUMO paralogues during processing. <i>Biochemical Journal</i> , 2006, 397, 279-288.	3.7	121
59	I κ B α -Mediated Proteolysis of NF- κ B1 p105 Requires Phosphorylation of p105 Serines 927 and 932. <i>Molecular and Cellular Biology</i> , 2003, 23, 402-413.	2.3	119
60	Modification of nuclear PML protein by SUMO-1 regulates Fas-induced apoptosis in rheumatoid arthritis synovial fibroblasts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 5073-5078.	7.1	119
61	Common Properties of Nuclear Body Protein SP100 and TIF1 α : Chromatin Factor: Role of SUMO Modification. <i>Molecular and Cellular Biology</i> , 2001, 21, 3314-3324.	2.3	118
62	Antipeptide antisera define neutralizing epitopes on the adenovirus hexon. <i>Journal of General Virology</i> , 1992, 73, 1429-1435.	2.9	113
63	The origin of adenovirus DNA replication: minimal DNA sequence requirement in vivo.. <i>EMBO Journal</i> , 1985, 4, 421-426.	7.8	110
64	SUMO protease SENP1 induces isomerization of the scissile peptide bond. <i>Nature Structural and Molecular Biology</i> , 2006, 13, 1069-1077.	8.2	110
65	Characterization of I κ B α Nuclear Import Pathway. <i>Journal of Biological Chemistry</i> , 1999, 274, 6804-6812.	3.4	109
66	NF- κ B Inhibits Apoptosis in Murine Mammary Epithelia. <i>Journal of Biological Chemistry</i> , 2000, 275, 12737-12742.	3.4	109
67	Stable-isotope labeling with amino acids in nematodes. <i>Nature Methods</i> , 2011, 8, 849-851.	19.0	108
68	Identification of a Substrate Recognition Site on Ubc9. <i>Journal of Biological Chemistry</i> , 2002, 277, 21740-21748.	3.4	107
69	Post-translational modification by SUMO. <i>Toxicology</i> , 2010, 278, 288-293.	4.2	105
70	Decoding the SUMO signal. <i>Biochemical Society Transactions</i> , 2013, 41, 463-473.	3.4	105
71	The Protein Stability and Transcriptional Activity of p63 α are Regulated by SUMO-1 Conjugation. <i>Cell Cycle</i> , 2005, 4, 183-190.	2.6	104
72	Structural basis of NEDD8 ubiquitin discrimination by the deNEDDylating enzyme NEDP1. <i>EMBO Journal</i> , 2005, 24, 1341-1351.	7.8	103

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73	SUMO-1 Modification Alters ADAR1 Editing Activity. <i>Molecular Biology of the Cell</i> , 2005, 16, 5115-5126.	2.1	102
74	Adenovirus DNA Replication. <i>Cold Spring Harbor Perspectives in Biology</i> , 2013, 5, a013003-a013003.	5.5	100
75	Expression of the genome of potato leafroll virus: readthrough of the coat protein termination codon in vivo. <i>Journal of General Virology</i> , 1990, 71, 2251-2256.	2.9	99
76	Structural basis for the RING-catalyzed synthesis of K63-linked ubiquitin chains. <i>Nature Structural and Molecular Biology</i> , 2015, 22, 597-602.	8.2	99
77	Functional interactions between ubiquitin E2 enzymes and TRIM proteins. <i>Biochemical Journal</i> , 2011, 434, 309-319.	3.7	93
78	Ube2W conjugates ubiquitin to Î±-amino groups of protein N-termini. <i>Biochemical Journal</i> , 2013, 453, 137-145.	3.7	90
79	Replication of adenovirus mini-chromosomes. <i>Journal of Molecular Biology</i> , 1984, 175, 493-510.	4.2	89
80	Role of an N-Terminal Site of Ubc9 in SUMO-1, -2, and -3 Binding and Conjugation. <i>Biochemistry</i> , 2003, 42, 9959-9969.	2.5	89
81	Regulation of transcription factors by protein degradation. <i>Cellular and Molecular Life Sciences</i> , 2000, 57, 1207-1219.	5.4	88
82	Characterization of SENP7, a SUMO-2/3-specific isopeptidase. <i>Biochemical Journal</i> , 2009, 421, 223-230.	3.7	88
83	Mutations of <i>NFKBIA</i> , encoding Î±, are a recurrent finding in classical Hodgkin lymphoma but are not a unifying feature of non-EBV-associated cases. <i>International Journal of Cancer</i> , 2009, 125, 1334-1342.	5.1	85
84	Dss1 Is a 26S Proteasome Ubiquitin Receptor. <i>Molecular Cell</i> , 2014, 56, 453-461.	9.7	81
85	Proteotoxic stress reprograms the chromatin landscape of SUMO modification. <i>Science Signaling</i> , 2015, 8, rs7.	3.6	81
86	Sequence specificity for the initiation of RNA-primed simian virus 40 DNA synthesis in vivo. <i>Journal of Molecular Biology</i> , 1984, 175, 131-157.	4.2	80
87	Properties of herpesvirus-induced "immediate early" polypeptides. <i>Virology</i> , 1980, 104, 230-234.	2.4	79
88	Origin of adenovirus DNA replication. <i>Journal of Molecular Biology</i> , 1985, 186, 129-136.	4.2	79
89	Control of NF-Î±B transcriptional activation by signal induced proteolysis of Î±. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 1999, 354, 1601-1609.	4.0	79
90	Reanalysis of phosphoproteomics data uncovers ADP-ribosylation sites. <i>Nature Methods</i> , 2012, 9, 771-772.	19.0	79

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91	Global Reprogramming of Host SUMOylation during Influenza Virus Infection. <i>Cell Reports</i> , 2015, 13, 1467-1480.	6.4	79
92	Recognition mechanisms in the synthesis of animal virus DNA. <i>Biochemical Journal</i> , 1989, 258, 3-16.	3.7	78
93	P14ARF promotes accumulation of SUMO-1 conjugated (H)Mdm2. <i>FEBS Letters</i> , 2002, 528, 207-211.	2.8	78
94	An influenza virus-triggered SUMO switch orchestrates co-opted endogenous retroviruses to stimulate host antiviral immunity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 17399-17408.	7.1	78
95	Identification and purification of EBP1: a HeLa cell protein that binds to a region overlapping the 'core' of the SV40 enhancer.. <i>Genes and Development</i> , 1988, 2, 991-1002.	5.9	76
96	NF-kappa B-dependent induction of the NF-kappa B p50 subunit gene promoter underlies self-perpetuation of human immunodeficiency virus transcription in monocytic cells.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1992, 89, 7826-7830.	7.1	74
97	Recognition of the adenovirus type 2 origin of DNA replication by the virally encoded DNA polymerase and preterminal proteins.. <i>EMBO Journal</i> , 1992, 11, 761-768.	7.8	71
98	SUMO modification of the DEAD box protein p68 modulates its transcriptional activity and promotes its interaction with HDAC1. <i>Oncogene</i> , 2007, 26, 5866-5876.	5.9	69
99	A SUMO-Dependent Protein Network Regulates Chromosome Congression during Oocyte Meiosis. <i>Molecular Cell</i> , 2017, 65, 66-77.	9.7	69
100	The Adenovirus Type 40 Hexon: Sequence, Predicted Structure and Relationship to Other Adenovirus Hexons. <i>Journal of General Virology</i> , 1989, 70, 3203-3214.	2.9	69
101	Expression of a foreign epitope on the surface of the adenovirus hexon. <i>Journal of General Virology</i> , 1994, 75, 133-139.	2.9	68
102	SUMO Chain-Induced Dimerization Activates RNF4. <i>Molecular Cell</i> , 2014, 53, 880-892.	9.7	68
103	Regulation of the DNA binding activity of NF- κ B. <i>International Journal of Biochemistry and Cell Biology</i> , 1995, 27, 865-879.	2.8	67
104	Analysis of the SUMO2 Proteome during HSV-1 Infection. <i>PLoS Pathogens</i> , 2015, 11, e1005059.	4.7	66
105	Co-operative interactions between NFI and the adenovirus DNA binding protein at the adenovirus origin of replication.. <i>EMBO Journal</i> , 1989, 8, 1841-1848.	7.8	65
106	The SUMO protease SENP6 is a direct regulator of PML nuclear bodies. <i>Molecular Biology of the Cell</i> , 2011, 22, 78-90.	2.1	64
107	Arsenic-Induced SUMO-Dependent Recruitment of RNF4 into PML Nuclear Bodies. <i>Molecular Biology of the Cell</i> , 2010, 21, 4227-4239.	2.1	63
108	Conjugation of Human Topoisomerase α with Small Ubiquitin-like Modifiers 2/3 in Response to Topoisomerase Inhibitors: Cell Cycle Stage and Chromosome Domain Specificity. <i>Cancer Research</i> , 2008, 68, 2409-2418.	0.9	61

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109	E3 Ubiquitin Ligase HOIP Attenuates Apoptotic Cell Death Induced by Cisplatin. <i>Cancer Research</i> , 2014, 74, 2246-2257.	0.9	61
110	Human Immunodeficiency Virus Type 1 (HIV-1) Viral Protein R (Vpr) Interacts with Lys-tRNA Synthetase: Implications for Priming of HIV-1 Reverse Transcription. <i>Journal of Virology</i> , 1998, 72, 3037-3044.	3.4	60
111	DNA Sequence of the Adenovirus Type 41 Hexon Gene and Predicted Structure of the Protein. <i>Journal of General Virology</i> , 1988, 69, 2291-2301.	2.9	59
112	Modulation of A ¹² generation by small ubiquitin-like modifiers does not require conjugation to target proteins. <i>Biochemical Journal</i> , 2007, 404, 309-316.	3.7	59
113	Aphid Acquisition and Cellular Transport of Potato leafroll virus-like Particles Lacking P5 Readthrough Protein. <i>Phytopathology</i> , 2000, 90, 1153-1161.	2.2	58
114	Inhibition of nuclear factor- κ B activation un-masks the ability of TNF- α to induce human eosinophil apoptosis. <i>European Journal of Immunology</i> , 2002, 32, 457-466.	2.9	58
115	Proteome-wide identification of SUMO modification sites by mass spectrometry. <i>Nature Protocols</i> , 2015, 10, 1374-1388.	12.0	56
116	Interaction between hnRNPA1 and I κ B α Is Required for Maximal Activation of NF- κ B-Dependent Transcription. <i>Molecular and Cellular Biology</i> , 2001, 21, 3482-3490.	2.3	55
117	NF- κ B is a critical regulator of the survival of rodent and human hepatic myofibroblasts. <i>Journal of Hepatology</i> , 2008, 48, 589-597.	3.7	55
118	Nuclear Factor- κ B Activation via Tyrosine Phosphorylation of Inhibitor κ B- α Is Crucial for Ciliary Neurotrophic Factor-Promoted Neurite Growth from Developing Neurons. <i>Journal of Neuroscience</i> , 2007, 27, 9664-9669.	3.6	53
119	Fourier Transform Ion Cyclotron Resonance Mass Spectrometry for the Analysis of Small Ubiquitin-like Modifier (SUMO) Modification: A Identification of Lysines in RanBP2 and SUMO Targeted for Modification during the E3 AutoSUMOylation Reaction. <i>Analytical Chemistry</i> , 2005, 77, 6310-6319.	6.5	51
120	The P-body component USP52/PAN2 is a novel regulator of <i>HIF1A</i> mRNA stability. <i>Biochemical Journal</i> , 2013, 451, 185-194.	3.7	51
121	Dynamic SUMO modification regulates mitotic chromosome assembly and cell cycle progression in <i>Caenorhabditis elegans</i> . <i>Nature Communications</i> , 2014, 5, 5485.	12.8	51
122	Identification of Sites of Ubiquitination in Proteins: A Fourier Transform Ion Cyclotron Resonance Mass Spectrometry Approach. <i>Analytical Chemistry</i> , 2004, 76, 6982-6988.	6.5	50
123	Regulation of Homeodomain-interacting Protein Kinase 2 (HIPK2) Effector Function through Dynamic Small Ubiquitin-related Modifier-1 (SUMO-1) Modification. <i>Journal of Biological Chemistry</i> , 2005, 280, 29224-29232.	3.4	50
124	Quantitative analysis of multi-protein interactions using FRET: Application to the SUMO pathway. <i>Protein Science</i> , 2008, 17, 777-784.	7.6	50
125	Structural Basis of BRCC36 Function in DNA Repair and Immune Regulation. <i>Molecular Cell</i> , 2019, 75, 483-497.e9.	9.7	50
126	Extra-pair paternity in the shag, <i>Phalacrocorax aristotelis</i> determined by DNA fingerprinting. <i>Journal of Zoology</i> , 1992, 226, 399-408.	1.7	49

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127	Adenovirus DNA binding protein: helix destabilising properties. <i>Nucleic Acids Research</i> , 1994, 22, 742-748.	14.5	49
128	Autophagosomes cooperate in the degradation of intracellular C-terminal fragments of the amyloid precursor protein <i>via</i> the MVB/lysosomal pathway. <i>FASEB Journal</i> , 2017, 31, 2446-2459.	0.5	47
129	Role of Two Residues Proximal to the Active Site of Ubc9 in Substrate Recognition by the Ubc9-SUMO-1 Thioester Complex. <i>Biochemistry</i> , 2003, 42, 3168-3179.	2.5	46
130	Screen for multi-SUMO-binding proteins reveals a multi-SIM-binding mechanism for recruitment of the transcriptional regulator ZMYM2 to chromatin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E4854-63.	7.1	46
131	Sequence requirement for specific interaction of an enhancer binding protein (EBP1) with DNA. <i>Nucleic Acids Research</i> , 1989, 17, 499-516.	14.5	45
132	Medea SUMOylation restricts the signaling range of the Dpp morphogen in the <i>Drosophila</i> embryo. <i>Genes and Development</i> , 2008, 22, 2578-2590.	5.9	45
133	Sumoylation controls host anti-bacterial response to the gut invasive pathogen <i>Shigella flexneri</i> . <i>EMBO Reports</i> , 2014, 15, 965-972.	4.5	45
134	A Proteomic Approach to Analyze the Aspirin-mediated Lysine Acetylome. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 310-326.	3.8	45
135	SUMOylation of HNF4 β regulates protein stability and hepatocyte function. <i>Journal of Cell Science</i> , 2012, 125, 3630-3635.	2.0	43
136	Viable Viruses with Deletions in the Left Inverted Terminal Repeat Define the Adenovirus Origin of DNA Replication. <i>Journal of General Virology</i> , 1986, 67, 321-332.	2.9	41
137	SUMO Modification of the Ets-related Transcription Factor ERM Inhibits Its Transcriptional Activity. <i>Journal of Biological Chemistry</i> , 2005, 280, 24330-24338.	3.4	41
138	SUMO Modification Regulates MafB-Driven Macrophage Differentiation by Enabling Myb-Dependent Transcriptional Repression. <i>Molecular and Cellular Biology</i> , 2007, 27, 5554-5564.	2.3	41
139	Sp100 Isoform-Specific Regulation of Human Adenovirus 5 Gene Expression. <i>Journal of Virology</i> , 2014, 88, 6076-6092.	3.4	41
140	Antibody RING-Mediated Destruction of Endogenous Proteins. <i>Molecular Cell</i> , 2020, 79, 155-166.e9.	9.7	40
141	Absolute SILAC-Compatible Expression Strain Allows Sumo-2 Copy Number Determination in Clinical Samples. <i>Journal of Proteome Research</i> , 2011, 10, 4869-4875.	3.7	39
142	A role for paralog-specific sumoylation in histone deacetylase 1 stability. <i>Journal of Molecular Cell Biology</i> , 2013, 5, 416-427.	3.3	38
143	PML isoforms in response to arsenic: high resolution analysis of PML body structure and degradation characteristics. <i>Journal of Cell Science</i> , 2014, 127, 365-75.	2.0	38
144	Ubiquitin C-terminal hydrolases cleave isopeptide- and peptide-linked ubiquitin from structured proteins but do not edit ubiquitin homopolymers. <i>Biochemical Journal</i> , 2015, 466, 489-498.	3.7	38

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145	Rapid generation of endogenously driven transcriptional reporters in cells through CRISPR/Cas9. <i>Scientific Reports</i> , 2015, 5, 9811.	3.3	38
146	Ufd1-Npl4 Recruit Cdc48 for Disassembly of Ubiquitylated CMG Helicase at the End of Chromosome Replication. <i>Cell Reports</i> , 2017, 18, 3033-3042.	6.4	38
147	DNA sequences required for the initiation of adenovirus type 4 DNA replication in vitro. <i>Journal of Molecular Biology</i> , 1988, 201, 57-67.	4.2	37
148	Structural insight into SUMO chain recognition and manipulation by the ubiquitin ligase RNF4. <i>Nature Communications</i> , 2014, 5, 4217.	12.8	37
149	Interaction of the C-terminal region of p105 with the nuclear localisation signal of p50 is required for inhibition of NF- κ B binding activity. <i>Nucleic Acids Research</i> , 1993, 21, 4516-4523.	14.5	36
150	The carboxy-terminus of I κ B α determines susceptibility to degradation by the catalytic core of the proteasome. <i>Oncogene</i> , 1997, 15, 1841-1850.	5.9	35
151	SUMO-1 Modification of Human Transcription Factor (TF) IID Complex Subunits. <i>Journal of Biological Chemistry</i> , 2005, 280, 9937-9945.	3.4	35
152	Inhibition of NF- κ B by a cell permeable form of I κ B α induces apoptosis in eosinophils. <i>Biochemical and Biophysical Research Communications</i> , 2005, 326, 632-637.	2.1	35
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