Joan Weliky Conaway

List of Publications by Year
in descending order

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152 papers	15,465 citations	20036 63 h-index	20023 121 g-index
154 all docs	154 docs citations	154 times ranked	17110 citing authors

#	Article	IF	Citations
1	A High-throughput Automated ELISA Assay for Detection of IgG Antibodies to the SARS-CoV-2 Spike Protein. Bio-protocol, 2022, 12, e4301.	0.2	0
2	Genome-wide analysis of cis-regulatory changes underlying metabolic adaptation of cavefish. Nature Genetics, 2022, 54, 684-693.	9.4	14
3	UBAP2/UBAP2L regulate UV-induced ubiquitylation of RNA polymerase II and are the human orthologues of yeast Def1. DNA Repair, 2022, 115, 103343.	1.3	6
4	The 3â€² Pol II pausing at replication-dependent histone genes is regulated by Mediator through Cajal bodiesâ€™ association with histone locus bodies. Nature Communications, 2022, 13, .	5.8	9
5	A role for the Cockayne Syndrome B (CSB)-Elongin ubiquitin ligase complex in signal-dependent RNA polymerase II transcription. Journal of Biological Chemistry, 2021, 297, 100862.	1.6	3
6	Multiple roles for PARP1 in ALC1-dependent nucleosome remodeling. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	3
7	Elongin functions as a loading factor for Mediator at ATF6Îą-regulated ER stress response genes. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, e2108751118.	3.3	2
8	A six-amino-acid motif is a major determinant in functional evolution of HOX1 proteins. Genes and Development, 2020, 34, 1680-1696.	2.7	16
9	NRBP1-Containing CRL2/CRL4A Regulates Amyloid Î² Production by Targeting BRI2 and BRI3 for Degradation. Cell Reports, 2020, 30, 3478-3491.e6.	2.9	20
10	Regulation of the RNAPII Pool Is Integral to the DNA Damage Response. Cell, 2020, 180, 1245-1261.e21.	13.5	116
11	The role of Mediator and Little Elongation Complex in transcription termination. Nature Communications, 2020, 11, 1063.	5.8	21
12	Roles of Mediator subunit MED26 in Regulation of Postâ€initiation Events in RNA Pol II Transcription. FASEB Journal, 2020, 34, 1-1.	0.2	0
13	The hunt for RNA polymerase II elongation factors: a historical perspective. Nature Structural and Molecular Biology, 2019, 26, 771-776.	3.6	15
14	A Role for FACT in RNA Polymerase II Promoter-Proximal Pausing. Cell Reports, 2019, 27, 3770-3779.e7.	2.9	41
15	Artificial RNA Polymerase II Elongation Complexes for Dissecting Co-transcriptional RNA Processing Events. Journal of Visualized Experiments, 2019, , .	0.2	1
16	DNA-dependent protein kinase catalytic subunit (DNA-PKcs) contributes to incorporation of histone variant H2A.Z into nucleosomes. Protein and Cell, 2019, 10, 694-699.	4.8	1
17	Imaging-based assays for investigating functions of the RNA polymerase II elongation factor Elongin and the Elongin ubiquitin ligase. Methods, 2019, 159-160, 157-164.	1.9	6
18	Frozen in Transcription: Cryo-EM Structures of Pol II Transcribing through a Nucleosome. Molecular Cell, 2018, 72, 802-804.	4.5	0

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19	Schizosaccharomyces pombePol II transcription elongation factor ELL functions as part of a rudimentary super elongation complex. Nucleic Acids Research, 2018, 46, 10095-10105.	6.5	7
20	CTD-dependent and -independent mechanisms govern co-transcriptional capping of Pol II transcripts. Nature Communications, 2018, 9, 3392.	5.8	21
21	Mediator structure and rearrangements required for holoenzyme formation. Nature, 2017, 544, 196-201.	13.7	120
22	O-Linked N-acetylglucosamine transferase 1 regulates global histone H4 acetylation via stabilization of the nonspecific lethal protein NSL3. Journal of Biological Chemistry, 2017, 292, 10014-10025.	1.6	13
23	Cockayne syndrome B protein regulates recruitment of the Elongin A ubiquitin ligase to sites of DNA damage. Journal of Biological Chemistry, 2017, 292, 6431-6437.	1.6	16
24	Role for the MED21-MED7 Hinge in Assembly of the Mediator-RNA Polymerase II Holoenzyme. Journal of Biological Chemistry, 2016, 291, 26886-26898.	1.6	19
25	Multiomic Analysis of the UV-Induced DNA Damage Response. Cell Reports, 2016, 15, 1597-1610.	2.9	162
26	Conserved abundance and topological features in chromatinâ€remodeling protein interaction networks. EMBO Reports, 2015, 16, 116-126.	2.0	17
27	Orchestrating transcription with the pol II CTD. Nature Reviews Molecular Cell Biology, 2015, 16, 128-128.	16.1	6
28	MED26 regulates the transcription of snRNA genes through the recruitment of little elongation complex. Nature Communications, 2015, 6, 5941.	5.8	42
29	TRIM29 regulates the assembly of DNA repair proteins into damaged chromatin. Nature Communications, 2015, 6, 7299.	5.8	45
30	Assembly of the Elongin A Ubiquitin Ligase Is Regulated by Genotoxic and Other Stresses. Journal of Biological Chemistry, 2015, 290, 15030-15041.	1.6	24
31	Regulation of the Rhp26 ^{ERCC6/CSB} chromatin remodeler by a novel conserved leucine latch motif. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 18566-18571.	3.3	22
32	Generation and Purification of Human INO80 Chromatin Remodeling Complexes and Subcomplexes. Journal of Visualized Experiments, 2014, , e51720.	0.2	8
33	Biochemical Assays for Analyzing Activities of ATP-dependent Chromatin Remodeling Enzymes. Journal of Visualized Experiments, 2014, , e51721.	0.2	3
34	Subunit Architecture and Functional Modular Rearrangements of the Transcriptional Mediator Complex. Cell, 2014, 157, 1430-1444.	13.5	179
35	The Mediator complex and transcription elongation. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2013, 1829, 69-75.	0.9	110
36	Multiple modes of regulation of the human Ino80 SNF2 ATPase by subunits of the INO80 chromatin-remodeling complex. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 20497-20502.	3.3	44

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37	Immunoaffinity Purification of Protein Complexes from Mammalian Cells. Methods in Molecular Biology, 2013, 977, 273-287.	0.4	14
38	Role for Human Mediator Subunit MED25 in Recruitment of Mediator to Promoters by Endoplasmic Reticulum Stress-responsive Transcription Factor ATF6Î±. Journal of Biological Chemistry, 2013, 288, 26179-26187.	1.6	33
39	Crosstalk between NSL Histone Acetyltransferase and MLL/SET Complexes: NSL Complex Functions in Promoting Histone H3K4 Di-Methylation Activity by MLL/SET Complexes. PLoS Genetics, 2013, 9, e1003940.	1.5	44
40	Transcriptional Properties of Mammalian Elongin A and Its Role in Stress Response. Journal of Biological Chemistry, 2013, 288, 24302-24315.	1.6	22
41	A conserved Mediatorâ€“CDK8 kinase module association regulates Mediatorâ€“RNA polymerase II interaction. Nature Structural and Molecular Biology, 2013, 20, 611-619.	3.6	184
42	Activation of the SNF2 Family ATPase ALC1 by Poly(ADP-ribose) in a Stable ALC1Â·PARP1Â·Nucleosome Intermediate. Journal of Biological Chemistry, 2012, 287, 43527-43532.	1.6	52
43	Endoplasmic Reticulum Stress-responsive Transcription Factor ATF6Î± Directs Recruitment of the Mediator of RNA Polymerase II Transcription and Multiple Histone Acetyltransferase Complexes. Journal of Biological Chemistry, 2012, 287, 23035-23045.	1.6	22
44	Transcriptional Elongation Factor Elongin A Regulates Retinoic Acid-Induced Gene Expression during Neuronal Differentiation. Cell Reports, 2012, 2, 1129-1136.	2.9	16
45	The Human EKC/KEOPS Complex Is Recruited to Cullin2 Ubiquitin Ligases by the Human Tumour Antigen PRAME. PLoS ONE, 2012, 7, e42822.	1.1	41
46	Introduction to Theme â€œChromatin, Epigenetics, and Transcriptionâ€• Annual Review of Biochemistry, 2012, 81, 61-64.	5.0	34
47	O-GlcNAc Transferase Catalyzes Site-Specific Proteolysis of HCF-1. Cell, 2011, 144, 376-388.	13.5	199
48	Human Mediator Subunit MED26 Functions as a Docking Site for Transcription Elongation Factors. Cell, 2011, 146, 92-104.	13.5	293
49	Function and regulation of the Mediator complex. Current Opinion in Genetics and Development, 2011, 21, 225-230.	1.5	258
50	Origins and activity of the Mediator complex. Seminars in Cell and Developmental Biology, 2011, 22, 729-734.	2.3	102
51	Subunit Organization of the Human INO80 Chromatin Remodeling Complex. Journal of Biological Chemistry, 2011, 286, 11283-11289.	1.6	93
52	Thematic Minireview Series on Computational Systems Biology. Journal of Biological Chemistry, 2011, 286, 23621-23622.	1.6	0
53	The tumour antigen PRAME is a subunit of a Cul2 ubiquitin ligase and associates with active NFY promoters. EMBO Journal, 2011, 30, 3786-3798.	3.5	59
54	Subunit Composition and Substrate Specificity of a MOF-containing Histone Acetyltransferase Distinct from the Male-specific Lethal (MSL) Complex. Journal of Biological Chemistry, 2010, 285, 4268-4272.	1.6	211

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55	AFF4, a Component of the ELL/P-TEFb Elongation Complex and a Shared Subunit of MLL Chimeras, Can Link Transcription Elongation to Leukemia. Molecular Cell, 2010, 37, 429-437.	4.5	504
56	Poly(ADP-ribosyl)ation directs recruitment and activation of an ATP-dependent chromatin remodeler. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 13770-13774.	3.3	322
57	Proteomics Reveals a Physical and Functional Link between Hepatocyte Nuclear Factor 4Î± and Transcription Factor IID. Journal of Biological Chemistry, 2009, 284, 32405-32412.	1.6	9
58	Distinct ubiquitin ligases act sequentially for RNA polymerase II polyubiquitylation. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 20705-20710.	3.3	144
59	Direct Inhibition of RNA Polymerase II Transcription by RECQL5. Journal of Biological Chemistry, 2009, 284, 23197-23203.	1.6	52
60	Mediator Comes Out from the Shadows. Structure, 2009, 17, 485-486.	1.6	2
61	The INO80 chromatin remodeling complex in transcription, replication and repair. Trends in Biochemical Sciences, 2009, 34, 71-77.	3.7	157
62	MED19 and MED26 Are Synergistic Functional Targets of the RE1 Silencing Transcription Factor in Epigenetic Silencing of Neuronal Gene Expression. Journal of Biological Chemistry, 2009, 284, 2648-2656.	1.6	64
63	Interaction of Hepatocyte nuclear factor 4alpha (HNF4alpha) with the TATA box binding protein (TBP) contributes to TFIID recruitment and HNF4alpha dependent transcription. FASEB Journal, 2009, 23, 660.12.	0.2	0
64	Functions of the Uch37 deubiquitinating enzyme in the proteasome and the INO80 chromatin remodeling complex. FASEB Journal, 2009, 23, 669.1.	0.2	0
65	ALC1: A Chromatin Remodeling Enzyme Activated by Poly(ADPâ€Ribose) Polymerase (PARP) and NAD. FASEB Journal, 2009, 23, 488.2.	0.2	0
66	Mammalian Elongin A complex mediates DNAâ€damageâ€induced ubiquitylation and degradation of Rpb1. FASEB Journal, 2009, 23, 495.2.	0.2	1
67	When transcription meets recombination: a lesson from the human RECQ protein complexes. F1000 Biology Reports, 2009, 1, 76.	4.0	0
68	Mammalian Elongin A complex mediates DNA-damage-induced ubiquitylation and degradation of Rpb1. EMBO Journal, 2008, 27, 3256-3266.	3.5	88
69	New clues to actin function in chromatin regulation. Nature Structural and Molecular Biology, 2008, 15, 432-433.	3.6	4
70	Distinct Modes of Regulation of the Uch37 Deubiquitinating Enzyme in the Proteasome and in the Ino80 Chromatin-Remodeling Complex. Molecular Cell, 2008, 31, 909-917.	4.5	132
71	Probabilistic assembly of human protein interaction networks from label-free quantitative proteomics. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 1454-1459.	3.3	220
72	Characterization of Cullin-box Sequences That Direct Recruitment of Cul2-Rbx1 and Cul5-Rbx2 Modules to Elongin BC-based Ubiquitin Ligases. Journal of Biological Chemistry, 2008, 283, 8005-8013.	1.6	147

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73	Neuralized-like 1 (Neurl1) Targeted to the Plasma Membrane by N-Myristoylation Regulates the Notch Ligand Jagged1. Journal of Biological Chemistry, 2008, 283, 3846-3853.	1.6	69
74	Identification and Characterization of a Schizosaccharomyces pombe RNA Polymerase II Elongation Factor with Similarity to the Metazoan Transcription Factor ELL. Journal of Biological Chemistry, 2007, 282, 5761-5769.	1.6	25
75	YY1 functions with INO80 to activate transcription. Nature Structural and Molecular Biology, 2007, 14, 872-874.	3.6	178
76	Purification of a Human SRCAP Complex That Remodels Chromatin by Incorporating the Histone Variant H2A.Z into Nucleosomes. Biochemistry, 2006, 45, 5671-5677.	1.2	211
77	RNA Polymerase II: A â€œNobelâ€•Enzyme Demystified. Molecular Cell, 2006, 24, 637-642.	4.5	3
78	Purification and assay of the human INO80 and SRCAP chromatin remodeling complexes. Methods, 2006, 40, 312-317.	1.9	31
79	Proteasome recruitment and activation of the Uch37 deubiquitinating enzyme by Adrm1. Nature Cell Biology, 2006, 8, 994-1002.	4.6	282
80	RNA polymerase II bypass of oxidative DNA damage is regulated by transcription elongation factors. EMBO Journal, 2006, 25, 5481-5491.	3.5	160
81	Quantitative proteomic analysis of distinct mammalian Mediator complexes using normalized spectral abundance factors. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 18928-18933.	3.3	495
82	The mammalian Mediator complex and its role in transcriptional regulation. Trends in Biochemical Sciences, 2005, 30, 250-255.	3.7	267
83	In and out: histone variant exchange in chromatin. Trends in Biochemical Sciences, 2005, 30, 680-687.	3.7	134
84	ASB2 Is an Elongin BC-interacting Protein That Can Assemble with Cullin 5 and Rbx1 to Reconstitute an E3 Ubiquitin Ligase Complex. Journal of Biological Chemistry, 2005, 280, 5468-5474.	1.6	64
85	A Mammalian Chromatin Remodeling Complex with Similarities to the Yeast INO80 Complex. Journal of Biological Chemistry, 2005, 280, 41207-41212.	1.6	211
86	Regulation of Heat Shock Gene Expression by RNA Polymerase II Elongation Factor, Elongin A. Journal of Biological Chemistry, 2005, 280, 4017-4020.	1.6	34
87	ELL-associated factors 1 and 2 are positive regulators of RNA polymerase II elongation factor ELL. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 10094-10098.	3.3	81
88	The Mammalian YL1 Protein Is a Shared Subunit of the TRRAP/TIP60 Histone Acetyltransferase and SRCAP Complexes. Journal of Biological Chemistry, 2005, 280, 13665-13670.	1.6	185
89	The mammalian Mediator complex. FEBS Letters, 2005, 579, 904-908.	1.3	43
90	A Mammalian Mediator Subunit that Shares Properties with Saccharomyces cerevisiae Mediator Subunit Cse2. Journal of Biological Chemistry, 2004, 279, 5846-5851.	1.6	24

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91	Both BC-Box Motifs of Adenovirus Protein E4orf6 Are Required To Efficiently Assemble an E3 Ligase Complex That Degrades p53. Molecular and Cellular Biology, 2004, 24, 9619-9629.	1.1	91
92	In Vivo Requirement of the RNA Polymerase II Elongation Factor Elongin A for Proper Gene Expression and Development. Molecular and Cellular Biology, 2004, 24, 9911-9919.	1.1	33
93	Identification of Elongin C and Skp1 Sequences That Determine Cullin Selection. Journal of Biological Chemistry, 2004, 279, 43019-43026.	1.6	10
94	VHL-box and SOCS-box domains determine binding specificity for Cul2-Rbx1 and Cul5-Rbx2 modules of ubiquitin ligases. Genes and Development, 2004, 18, 3055-3065.	2.7	405
95	A Set of Consensus Mammalian Mediator Subunits Identified by Multidimensional Protein Identification Technology. Molecular Cell, 2004, 14, 685-691.	4.5	270
96	A Unified Nomenclature for Protein Subunits of Mediator Complexes Linking Transcriptional Regulators to RNA Polymerase II. Molecular Cell, 2004, 14, 553-557.	4.5	230
97	Ozz-E3, A Muscle-Specific Ubiquitin Ligase, Regulates Î²-Catenin Degradation during Myogenesis. Developmental Cell, 2004, 6, 269-282.	3.1	85
98	Von Hippel-Lindau (VHL) Protein. , 2004, , 416-418.		0
99	RNA Polymerase II and Basal Transcription Factors in Eukaryotes. , 2004, , 763-765.		0
100	The RNA Polymerase II Elongation Complex. Annual Review of Biochemistry, 2003, 72, 693-715.	5.0	212
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102	Identification of Mammalian Mediator Subunits with Similarities to Yeast Mediator Subunits Srb5, Srb6, Med11, and Rox3. Journal of Biological Chemistry, 2003, 278, 15123-15127.	1.6	46
103	Identification of New Subunits of the Multiprotein Mammalian TRRAP/TIP60-containing Histone Acetyltransferase Complex. Journal of Biological Chemistry, 2003, 278, 42733-42736.	1.6	196
104	A Mammalian Homolog of Drosophila melanogaster Transcriptional Coactivator Intersex Is a Subunit of the Mammalian Mediator Complex. Journal of Biological Chemistry, 2003, 278, 49671-49674.	1.6	44
105	von Hippel-Lindau protein binds hyperphosphorylated large subunit of RNA polymerase II through a proline hydroxylation motif and targets it for ubiquitination. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 2706-2711.	3.3	206
106	Preparation and Assay of RNA Polymerase II Elongation Factors Elongin and ELL. Methods in Enzymology, 2003, 371, 276-283.	0.4	2
107	Multiple Splice Variants of the Human HIF-3Î± Locus Are Targets of the von Hippel-Lindau E3 Ubiquitin Ligase Complex. Journal of Biological Chemistry, 2003, 278, 11032-11040.	1.6	238
108	Assays for Investigating the Mechanism of Promoter Escape by RNA Polymerase II. Methods in Enzymology, 2003, 370, 733-740.	0.4	1

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109	A Molecular Basis for Stabilization of the von Hippel-Lindau (VHL) Tumor Suppressor Protein by Components of the VHL Ubiquitin Ligase. Journal of Biological Chemistry, 2002, 277, 30388-30393.	1.6	51
110	Emerging Roles of Ubiquitin in Transcription Regulation. Science, 2002, 296, 1254-1258.	6.0	375
111	Mammalian mediator subunit mMED8 is an Elongin BC-interacting protein that can assemble with Cul2 and Rbx1 to reconstitute a ubiquitin ligase. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 10353-10358.	3.3	76
112	The Von Hippelâ€"Lindau tumor suppressor complex and regulation of hypoxia-inducible transcription. Advances in Cancer Research, 2002, 85, 1-12.	1.9	15
113	Structure of the Cul1â€"Rbx1â€"Skp1â€"F boxSkp2 SCF ubiquitin ligase complex. Nature, 2002, 416, 703-709.	13.7	1,322
114	Biochemical purification and pharmacological inhibition of a mammalian prolyl hydroxylase acting on hypoxia-inducible factor. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 13459-13464.	3.3	520
115	Roles of SCF and VHL Ubiquitin Ligases in Regulation of Cell Growth. Progress in Molecular and Subcellular Biology, 2002, 29, 1-15.	0.9	8
116	Mechanism of transcription initiation and promoter escape by RNA polymerase II. Current Opinion in Genetics and Development, 2001, 11, 209-214.	1.5	123
117	Defective Interplay of Activators and Repressors with TFIIH in Xeroderma Pigmentosum. Cell, 2001, 104, 353-363.	13.5	117
118	Cloning and Characterization of ELL-associated Proteins EAP45 and EAP20. Journal of Biological Chemistry, 2001, 276, 16528-16533.	1.6	50
119	MUF1, A Novel Elongin BC-interacting Leucine-rich Repeat Protein That Can Assemble with Cul5 and Rbx1 to Reconstitute a Ubiquitin Ligase. Journal of Biological Chemistry, 2001, 276, 29748-29753.	1.6	135
120	Transcription Factors TFIIF, ELL, and Elongin Negatively Regulate SII-induced Nascent Transcript Cleavage by Non-arrested RNA Polymerase II Elongation Intermediates. Journal of Biological Chemistry, 2001, 276, 23109-23114.	1.6	35
121	Degradation of p53 by adenovirus E4orf6 and E1B55K proteins occurs via a novel mechanism involving a Cullin-containing complex. Genes and Development, 2001, 15, 3104-3117.	2.7	418
122	Control of elongation by RNA polymerase II. Trends in Biochemical Sciences, 2000, 25, 375-380.	3.7	194
123	Elongin from Saccharomyces cerevisiae. Journal of Biological Chemistry, 2000, 275, 11174-11180.	1.6	21
124	The Elongin B Ubiquitin Homology Domain. Journal of Biological Chemistry, 1999, 274, 13629-13636.	1.6	15
125	A Role for the TFIIH XPB DNA Helicase in Promoter Escape by RNA Polymerase II. Journal of Biological Chemistry, 1999, 274, 22127-22130.	1.6	77
126	Dual Roles for Transcription Factor IIF in Promoter Escape by RNA Polymerase II. Journal of Biological Chemistry, 1999, 274, 35668-35675.	1.6	65

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128	Transcription Elongation and Human Disease. Annual Review of Biochemistry, 1999, 68, 301-319.	5.0	92
129	Reconstitution of G1 Cyclin Ubiquitination with Complexes Containing SCFGrr1 and Rbx1. Science, 1999, 284, 662-665.	6.0	367
130	Synthetic peptides define critical contacts between elongin C, elongin B, and the von Hippel-Lindau protein. Journal of Clinical Investigation, 1999, 104, 1583-1591.	3.9	89
131	The Elongin BC complex and the von Hippelâ€“Lindau tumor suppressor protein. Biochimica Et Biophysica Acta: Reviews on Cancer, 1998, 1377, M49-M54.	3.3	20
132	Characterization of the Residues Phosphorylated in Vitro by Different C-terminal Domain Kinases. Journal of Biological Chemistry, 1998, 273, 6769-6775.	1.6	106
133	Mechanism of Action of RNA Polymerase II Elongation Factor Elongin. Journal of Biological Chemistry, 1998, 273, 26610-26617.	1.6	15
134	Regulation of Hypoxia-Inducible mRNAs by the von Hippel-Lindau Tumor Suppressor Protein Requires Binding to Complexes Containing Elongins B/C and Cul2. Molecular and Cellular Biology, 1998, 18, 732-741.	1.1	349
135	Structure and Function of RNA Polymerase II Elongation Factor ELL. Journal of Biological Chemistry, 1997, 272, 22355-22363.	1.6	71
136	Identification of Elongin C Sequences Required for Interaction with the von Hippel-Lindau Tumor Suppressor Protein. Journal of Biological Chemistry, 1997, 272, 27444-27449.	1.6	16
137	Promoter Escape by RNA Polymerase II. Journal of Biological Chemistry, 1997, 272, 28175-28178.	1.6	39
138	General Transcription Factors for RNA Polymerase II1. Progress in Molecular Biology and Translational Science, 1997, 56, 327-346.	1.9	54
139	Assays for Investigating Transcription by RNA Polymerase IIin Vitro. Methods, 1997, 12, 192-202.	1.9	8
140	Mechanism and regulation of transcriptional elongation and termination by RNA polymerase II. Current Opinion in Genetics and Development, 1997, 7, 199-204.	1.5	41
141	A human cDNA encoding the 110-kDa A subunit of RNA polymerase II transcription factor elongin. Gene, 1996, 168, 277-278.	1.0	11
142	[18] Purification of RNA polymerase II general transcription factors from rat liver. Methods in Enzymology, 1996, 273, 194-207.	0.4	36
143	Interaction of von Hippel-Lindau tumor suppressor gene product with elongin. Methods in Enzymology, 1996, 274, 436-441.	0.4	7
144	Promoter Escape by RNA Polymerase II A ROLE FOR AN ATP COFACTOR IN SUPPRESSION OF ARREST BY POLYMERASE AT PROMOTER-PROXIMAL SITES. Journal of Biological Chemistry, 1996, 271, 23352-23356.	1.6	55

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146	A Role for ATP and TFIIH in Activation of the RNA Polymerase II Preinitiation Complex Prior to Transcription Initiation. Journal of Biological Chemistry, 1996, 271, 7245-7248.	1.6	76
147	Association of Cdk-activating kinase subunits with transcription factor TFIIH. Nature, 1995, 374, 280-282.	13.7	366
148	A Novel Activity Associated with RNA Polymerase II Elongation Factor SIII. Journal of Biological Chemistry, 1995, 270, 24300-24305.	1.6	18
149	A human cDNA encoding the small subunit of RNA polymerase II transcription factor SIII. Gene, 1994, 150, 413-414.	1.0	8
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151	General Initiation Factors for RNA Polymerase II. Annual Review of Biochemistry, 1993, 62, 161-190.	5.0	420
152	RNA polymerase II initiation factor Î± from rat liver is almost identical to human TFIIB. Nucleic Acids Research, 1992, 20, 3250-3250.	6.5	51