

Dylan J Taatjes

List of Publications by Year in descending order

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56
papers

8,947
citations

126907

33
h-index

168389

53
g-index

61
all docs

61
docs citations

61
times ranked

11236
citing authors

#	ARTICLE	IF	CITATIONS
1	Mediator and cohesin connect gene expression and chromatin architecture. <i>Nature</i> , 2010, 467, 430-435.	27.8	1,707
2	Transcription Factors Activate Genes through the Phase-Separation Capacity of Their Activation Domains. <i>Cell</i> , 2018, 175, 1842-1855.e16.	28.9	1,195
3	Activating RNAs associate with Mediator to enhance chromatin architecture and transcription. <i>Nature</i> , 2013, 494, 497-501.	27.8	759
4	The Mediator complex: a central integrator of transcription. <i>Nature Reviews Molecular Cell Biology</i> , 2015, 16, 155-166.	37.0	707
5	Pol ^{II} phosphorylation regulates a switch between transcriptional and splicing condensates. <i>Nature</i> , 2019, 572, 543-548.	27.8	457
6	Mediator kinase inhibition further activates super-enhancer-associated genes in AML. <i>Nature</i> , 2015, 526, 273-276.	27.8	307
7	CDK8 is a positive regulator of transcriptional elongation within the serum response network. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 194-201.	8.2	303
8	The human CDK8 subcomplex is a molecular switch that controls Mediator coactivator function. <i>Genes and Development</i> , 2009, 23, 439-451.	5.9	290
9	The human Mediator complex: a versatile, genome-wide regulator of transcription. <i>Trends in Biochemical Sciences</i> , 2010, 35, 315-322.	7.5	281
10	Partitioning of cancer therapeutics in nuclear condensates. <i>Science</i> , 2020, 368, 1386-1392.	12.6	281
11	CDK8 Kinase Phosphorylates Transcription Factor STAT1 to Selectively Regulate the Interferon Response. <i>Immunity</i> , 2013, 38, 250-262.	14.3	220
12	The Human CDK8 Subcomplex Is a Histone Kinase That Requires Med12 for Activity and Can Function Independently of Mediator. <i>Molecular and Cellular Biology</i> , 2009, 29, 650-661.	2.3	193
13	Mediator Condensates Localize Signaling Factors to Key Cell Identity Genes. <i>Molecular Cell</i> , 2019, 76, 753-766.e6.	9.7	188
14	Structure and mechanism of the RNA polymerase II transcription machinery. <i>Genes and Development</i> , 2020, 34, 465-488.	5.9	167
15	Regulatory diversity among metazoan co-activator complexes. <i>Nature Reviews Molecular Cell Biology</i> , 2004, 5, 403-410.	37.0	137
16	TRIM28 regulates RNA polymerase II promoter-proximal pausing and pause release. <i>Nature Structural and Molecular Biology</i> , 2014, 21, 876-883.	8.2	125
17	Human TFIID Kinase CDK7 Regulates Transcription-Associated Chromatin Modifications. <i>Cell Reports</i> , 2017, 20, 1173-1186.	6.4	123
18	Identification of Mediator Kinase Substrates in Human Cells using Cortistatin A and Quantitative Phosphoproteomics. <i>Cell Reports</i> , 2016, 15, 436-450.	6.4	117

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19	Redox Pathway Leading to the Alkylation of DNA by the Anthracycline, Antitumor Drugs Adriamycin and Daunomycin. <i>Journal of Medicinal Chemistry</i> , 1997, 40, 1276-1286.	6.4	114
20	The SCF ^{FBW7} ubiquitin ligase degrades MED13 and MED13L and regulates CDK8 module association with Mediator. <i>Genes and Development</i> , 2013, 27, 151-156.	5.9	96
21	Activator-Mediator binding regulates Mediator-cofactor interactions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 11283-11288.	7.1	92
22	Architecture of the Human and Yeast General Transcription and DNA Repair Factor TFIIH. <i>Molecular Cell</i> , 2015, 59, 794-806.	9.7	91
23	The essential and multifunctional TFIIH complex. <i>Protein Science</i> , 2018, 27, 1018-1037.	7.6	88
24	Regulatory functions of the Mediator kinases CDK8 and CDK19. <i>Transcription</i> , 2019, 10, 76-90.	3.1	79
25	Backtracked and paused transcription initiation intermediate of <i>Escherichia coli</i> RNA polymerase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E6562-E6571.	7.1	78
26	Doxoform and Daunoform: Anthracycline ⁺ Formaldehyde Conjugates Toxic to Resistant Tumor Cells. <i>Journal of Medicinal Chemistry</i> , 1997, 40, 2452-2461.	6.4	71
27	The Mediator complex as a master regulator of transcription by RNA polymerase II. <i>Nature Reviews Molecular Cell Biology</i> , 2022, 23, 732-749.	37.0	71
28	The complex structure and function of Mediator. <i>Journal of Biological Chemistry</i> , 2018, 293, 13778-13785.	3.4	65
29	A Kinase-Independent Role for Cyclin-Dependent Kinase 19 in p53 Response. <i>Molecular and Cellular Biology</i> , 2017, 37, .	2.3	57
30	TFIID Enables RNA Polymerase II Promoter-Proximal Pausing. <i>Molecular Cell</i> , 2020, 78, 785-793.e8.	9.7	55
31	Transcriptional Responses to IFN- β Require Mediator Kinase-Dependent Pause Release and Mechanistically Distinct CDK8 and CDK19 Functions. <i>Molecular Cell</i> , 2019, 76, 485-499.e8.	9.7	52
32	Selective inhibition of CDK7 reveals high-confidence targets and new models for TFIIH function in transcription. <i>Genes and Development</i> , 2020, 34, 1452-1473.	5.9	47
33	Merging Established Mechanisms with New Insights: Condensates, Hubs, and the Regulation of RNA Polymerase II Transcription. <i>Journal of Molecular Biology</i> , 2022, 434, 167216.	4.2	44
34	The nuclear interactome of DYRK1A reveals a functional role in DNA damage repair. <i>Scientific Reports</i> , 2019, 9, 6539.	3.3	42
35	Nuclear Targeting and Nuclear Retention of Anthracycline ⁺ Formaldehyde Conjugates Implicates DNA Covalent Bonding in the Cytotoxic Mechanism of Anthracyclines. <i>Chemical Research in Toxicology</i> , 1999, 12, 588-596.	3.3	37
36	The Continuing SAGA of TFIID and RNA Polymerase II Transcription. <i>Molecular Cell</i> , 2017, 68, 1-2.	9.7	31

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37	The Mediator kinase module: an interface between cell signaling and transcription. Trends in Biochemical Sciences, 2022, 47, 314-327.	7.5	26
38	Epidoxoform: A Hydrolytically More Stable Anthracycline-Formaldehyde Conjugate Toxic to Resistant Tumor Cells. Journal of Medicinal Chemistry, 1998, 41, 1306-1314.	6.4	22
39	Transcription factor enrichment analysis (TFEA) quantifies the activity of multiple transcription factors from a single experiment. Communications Biology, 2021, 4, 661.	4.4	21
40	The human β 53 isoform triggers metabolic and gene expression changes that activate mTOR and alter mitochondrial function. Aging Cell, 2013, 12, 863-872.	6.7	14
41	The β 40p53 isoform inhibits p53-dependent eRNA transcription and enables regulation by signal-specific transcription factors during p53 activation. PLoS Biology, 2021, 19, e3001364.	5.6	14
42	Studying transcription initiation by RNA polymerase with diffusion-based single-molecule fluorescence. Protein Science, 2017, 26, 1278-1290.	7.6	13
43	Everything at once: cryo-EM yields remarkable insights into human RNA polymerase II transcription. Nature Structural and Molecular Biology, 2021, 28, 540-543.	8.2	8
44	The Role of XPB/Xpd dsDNA Translocase Processivity in Transcription Start-site Scanning. Journal of Molecular Biology, 2021, 433, 166813.	4.2	8
45	Small-Molecule Probes to Target the Human Mediator Complex. Israel Journal of Chemistry, 2013, 53, 588-595.	2.3	7
46	Chemical Synthesis of the Multiply Phosphorylated and Biotinylated N-Terminal Transactivation Domain of Human p53 (p53TAD). Synlett, 2017, 28, 1917-1922.	1.8	6
47	Suppression of p53 response by targeting p53-Mediator binding with a stapled peptide. Cell Reports, 2022, 39, 110630.	6.4	5
48	Mediator redefines itself. Cell Research, 2014, 24, 775-776.	12.0	3
49	All in the Family: A Portrait of a Nuclear Receptor Co-Activator Complex. Molecular Cell, 2015, 57, 952-954.	9.7	3
50	Mediating transcription and RNA export. Nature, 2015, 526, 199-200.	27.8	3
51	Transcription Factor-Mediator Interfaces: Multiple and Multi-Valent. Journal of Molecular Biology, 2017, 429, 2996-2998.	4.2	3
52	RNA Polymerase II Transcription. Journal of Molecular Biology, 2021, 433, 167037.	4.2	3
53	Macromolecular Complexes in Transcription and Co-Transcriptional RNA Processing. Journal of Molecular Biology, 2016, 428, 2539-2541.	4.2	1
54	Mediator co-activator function is controlled by activator-induced structural shifts. FASEB Journal, 2010, 24, 679.4.	0.5	0

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55	CDK8 is a positive regulator of transcriptional elongation within the serum response network.. FASEB Journal, 2010, 24, 456.5.	0.5	0
56	Structure and Mechanism of the human Transcription Initiation Machinery. FASEB Journal, 2012, 26, 227.1.	0.5	0