

# Wen-Ming Yang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5206121/publications.pdf>

Version: 2024-02-01

40  
papers

6,117  
citations

186265

28  
h-index

302126

39  
g-index

40  
all docs

40  
docs citations

40  
times ranked

5782  
citing authors

#	ARTICLE	IF	CITATIONS
1	A complex containing N-CoR, mSin3 and histone deacetylase mediates transcriptional repression. <i>Nature</i> , 1997, 387, 43-48.	27.8	1,204
2	Histone Deacetylases Associated with the mSin3 Corepressor Mediate Mad Transcriptional Repression. <i>Cell</i> , 1997, 89, 349-356.	28.9	929
3	Transcriptional repression by YY1 is mediated by interaction with a mammalian homolog of the yeast global regulator RPD3. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 12845-12850.	7.1	521
4	Isolation and Characterization of cDNAs Corresponding to an Additional Member of the Human Histone Deacetylase Gene Family. <i>Journal of Biological Chemistry</i> , 1997, 272, 28001-28007.	3.4	396
5	Regulation of Transcription Factor YY1 by Acetylation and Deacetylation. <i>Molecular and Cellular Biology</i> , 2001, 21, 5979-5991.	2.3	385
6	Histone Deacetylases Specifically Down-regulate p53-dependent Gene Activation. <i>Journal of Biological Chemistry</i> , 2000, 275, 20436-20443.	3.4	363
7	The histone deacetylase-3 complex contains nuclear receptor corepressors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 7202-7207.	7.1	321
8	Functional Domains of Histone Deacetylase-3. <i>Journal of Biological Chemistry</i> , 2002, 277, 9447-9454.	3.4	201
9	Histone deacetylase interacts directly with DNA topoisomerase II. <i>Nature Genetics</i> , 2000, 26, 349-353.	21.4	159
10	RBP1 Recruits Both Histone Deacetylase-Dependent and -Independent Repression Activities to Retinoblastoma Family Proteins. <i>Molecular and Cellular Biology</i> , 1999, 19, 6632-6641.	2.3	156
11	SUMO5, a Novel Poly-SUMO Isoform, Regulates PML Nuclear Bodies. <i>Scientific Reports</i> , 2016, 6, 26509.	3.3	149
12	The Growth Suppressor PML Represses Transcription by Functionally and Physically Interacting with Histone Deacetylases. <i>Molecular and Cellular Biology</i> , 2001, 21, 2259-2268.	2.3	138
13	The Metastasis-associated Proteins 1 and 2 Form Distinct Protein Complexes with Histone Deacetylase Activity. <i>Journal of Biological Chemistry</i> , 2003, 278, 42560-42568.	3.4	123
14	Ligand-induced recruitment of a histone deacetylase in the negative-feedback regulation of the thyrotropin beta gene. <i>EMBO Journal</i> , 1999, 18, 5389-5398.	7.8	109
15	The FK506-binding protein 25 functionally associates with histone deacetylases and with transcription factor YY1. <i>EMBO Journal</i> , 2001, 20, 4814-4825.	7.8	103
16	Cyclophilin A and FKBP12 Interact with YY1 and Alter Its Transcriptional Activity. <i>Journal of Biological Chemistry</i> , 1995, 270, 15187-15193.	3.4	94
17	Differential Effects of Nuclear Receptor Corepressor (N-CoR) Expression Levels on Retinoic Acid Receptor-Mediated Repression Support the Existence of Dynamically Regulated Corepressor Complexes. <i>Molecular Endocrinology</i> , 1997, 11, 682-692.	3.7	83
18	Beyond Histone and Deacetylase: An Overview of Cytoplasmic Histone Deacetylases and Their Nonhistone Substrates. <i>Journal of Biomedicine and Biotechnology</i> , 2011, 2011, 1-15.	3.0	82

#	ARTICLE	IF	CITATIONS
19	The transcriptional factor YY1 upregulates the novel invasion suppressor HLJ1 expression and inhibits cancer cell invasion. <i>Oncogene</i> , 2005, 24, 4081-4093.	5.9	81
20	The Hepatitis B Virus X-associated Protein, XAP3, Is a Protein Kinase C-binding Protein. <i>Journal of Biological Chemistry</i> , 1997, 272, 16482-16489.	3.4	59
21	HDAC1/HDAC3 modulates PPARG2 transcription through the sumoylated CEBPD in hepatic lipogenesis. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2008, 1783, 1803-1814.	4.1	47
22	Yin-Yang 1 Activates Interleukin-4 Gene Expression in T Cells. <i>Journal of Biological Chemistry</i> , 2001, 276, 48871-48878.	3.4	44
23	Histone Deacetylase Activity Represses Gamma Interferon-Inducible HLA-DR Gene Expression following the Establishment of a DNase I-Hypersensitive Chromatin Conformation. <i>Molecular and Cellular Biology</i> , 2001, 21, 6495-6506.	2.3	39
24	Histone Deacetylase 10 Relieves Repression on the Melanogenic Program by Maintaining the Deacetylation Status of Repressors. <i>Journal of Biological Chemistry</i> , 2010, 285, 7187-7196.	3.4	37
25	Sumoylation of LAP1 is involved in the HDAC4-mediated repression of COX-2 transcription. <i>Nucleic Acids Research</i> , 2008, 36, 6066-6079.	14.5	36
26	Loss of ZBRK1 Contributes to the Increase of KAP1 and Promotes KAP1-Mediated Metastasis and Invasion in Cervical Cancer. <i>PLoS ONE</i> , 2013, 8, e73033.	2.5	35
27	FKBPs in chromatin modification and cancer. <i>Current Opinion in Pharmacology</i> , 2011, 11, 301-307.	3.5	32
28	Loading of PAX3 to Mitotic Chromosomes Is Mediated by Arginine Methylation and Associated with Waardenburg Syndrome. <i>Journal of Biological Chemistry</i> , 2015, 290, 20556-20564.	3.4	29
29	Differential Effects of Nuclear Receptor Corepressor (N-CoR) Expression Levels on Retinoic Acid Receptor-Mediated Repression Support the Existence of Dynamically Regulated Corepressor Complexes. <i>Molecular Endocrinology</i> , 1997, 11, 682-692.	3.7	29
30	Transcriptional and subcellular regulation of the TRIP-Br family. <i>Gene</i> , 2007, 388, 102-109.	2.2	25
31	Cloning and Characterization of the Mouse Histone Deacetylase-2 Gene. <i>Journal of Biological Chemistry</i> , 1998, 273, 28921-28930.	3.4	23
32	Transcriptional repression activity of PAX3 is modulated by competition between corepressor KAP1 and heterochromatin protein 1. <i>Biochemical and Biophysical Research Communications</i> , 2006, 349, 573-581.	2.1	20
33	Nuclear Proteins: Promising Targets for Cancer Drugs. <i>Current Cancer Drug Targets</i> , 2005, 5, 595-610.	1.6	17
34	PARP-2 regulates cell cycle-related genes through histone deacetylation and methylation independently of poly(ADP-ribosyl)ation. <i>Biochemical and Biophysical Research Communications</i> , 2013, 431, 58-64.	2.1	17
35	NAPP2, a Peroxisomal Membrane Protein, Is Also a Transcriptional Corepressor. <i>Genomics</i> , 2002, 79, 423-431.	2.9	14
36	Histone Deacetylases: Purification of the Enzymes, Substrates, and Assay Conditions. <i>Methods in Enzymology</i> , 2003, 377, 167-179.	1.0	11

#	ARTICLE	IF	CITATIONS
37	The transcriptional repression activity of STAF65 <sup>Δ3</sup> is facilitated by promoter tethering and nuclear import of class IIa histone deacetylases. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2014, 1839, 579-591.	1.9	4
38	Ocular disease-associated mutations diminish the mitotic chromosome retention ability of PAX6. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2021, 1864, 194751.	1.9	1
39	PAX3 loads onto pericentromeric heterochromatin during S phase through PARP1. <i>Anticancer Research</i> , 2014, 34, 4717-22.	1.1	1
40	M33 condenses chromatin through nuclear body formation and methylation of both histone H3 lysine 9 and lysine 27. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2021, 1868, 119100.	4.1	0