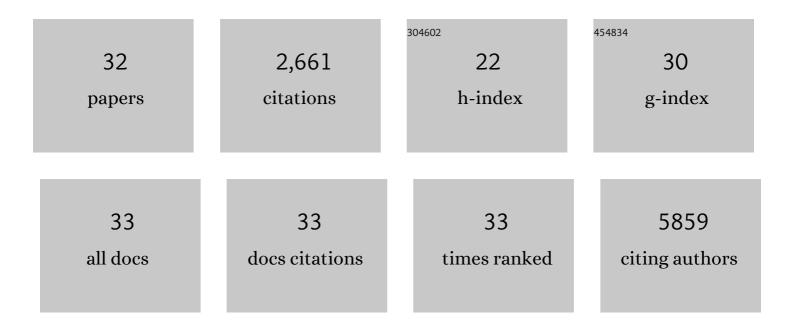
## James M Ward

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6945977/publications.pdf Version: 2024-02-01



IAMES M WADD

#	Article	IF	CITATIONS
1	Identification of miRNA Changes in Alzheimer's Disease Brain and CSF Yields Putative Biomarkers and Insights into Disease Pathways. Journal of Alzheimer's Disease, 2008, 14, 27-41.	1.2	835
2	Different Neuronal Activity Patterns Induce Different Gene Expression Programs. Neuron, 2018, 98, 530-546.e11.	3.8	262
3	MicroRNA 132 Regulates Nutritional Stress-Induced Chemokine Production through Repression of SirT1. Molecular Endocrinology, 2009, 23, 1876-1884.	3.7	223
4	INO80 Facilitates Pluripotency Gene Activation in Embryonic Stem Cell Self-Renewal, Reprogramming, and Blastocyst Development. Cell Stem Cell, 2014, 14, 575-591.	5.2	148
5	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 2062-2074.	1.5	147
6	Core and region-enriched networks of behaviorally regulated genes and the singing genome. Science, 2014, 346, 1256780.	6.0	97
7	The THO Complex Regulates Pluripotency Gene mRNA Export and Controls Embryonic Stem Cell Self-Renewal and Somatic Cell Reprogramming. Cell Stem Cell, 2013, 13, 676-690.	5.2	85
8	Genome-wide DNA methylation and long-term ambient air pollution exposure in Korean adults. Clinical Epigenetics, 2019, 11, 37.	1.8	76
9	High-coverage sequencing and annotated assemblies of the budgerigar genome. GigaScience, 2014, 3, 11.	3.3	75
10	Revealing a human p53 universe. Nucleic Acids Research, 2018, 46, 8153-8167.	6.5	75
11	Comparison of smoking-related DNA methylation between newborns from prenatal exposure and adults from personal smoking. Epigenomics, 2019, 11, 1487-1500.	1.0	64
12	Hippocampal Subregions Express Distinct Dendritic Transcriptomes that Reveal Differences in Mitochondrial Function in CA2. Cell Reports, 2019, 29, 522-539.e6.	2.9	61
13	SIRT1-Mediated Deacetylation of CRABPII Regulates Cellular Retinoic Acid Signaling and Modulates Embryonic Stem Cell Differentiation. Molecular Cell, 2014, 55, 843-855.	4.5	60
14	BRG1 governs glucocorticoid receptor interactions with chromatin and pioneer factors across the genome. ELife, 2018, 7, .	2.8	59
15	Convergent Differential Regulation of Parvalbumin in the Brains of Vocal Learners. PLoS ONE, 2012, 7, e29457.	1.1	45
16	Histone Hypervariants H2A.Z.1 and H2A.Z.2 Play Independent and Context-Specific Roles in Neuronal Activity-Induced Transcription of <i>Arc/Arg3.1</i> and Other Immediate Early Genes. ENeuro, 2017, 4, ENEURO.0040-17.2017.	0.9	43
17	Epigenome-wide association study of DNA methylation and adult asthma in the Agricultural Lung Health Study. European Respiratory Journal, 2020, 56, 2000217.	3.1	40
18	TNF is required for TLR ligand–mediated but not protease-mediated allergic airway inflammation. Journal of Clinical Investigation, 2017, 127, 3313-3326.	3.9	35

JAMES M WARD

#	Article	IF	CITATIONS
19	Analysis of Chromatin Dynamics during Glucocorticoid Receptor Activation. Molecular and Cellular Biology, 2012, 32, 1805-1817.	1.1	34
20	Tumor suppressor SMARCB1 suppresses super-enhancers to govern hESC lineage determination. ELife, 2019, 8, .	2.8	30
21	Microproteomics: Quantitative Proteomic Profiling of Small Numbers of Laser-Captured Cells. Cold Spring Harbor Protocols, 2011, 2011, pdb.prot5573.	0.2	27
22	MiRNA-Mediated Regulation of the SWI/SNF Chromatin Remodeling Complex Controls Pluripotency and Endodermal Differentiation in Human ESCs. Stem Cells, 2015, 33, 2925-2935.	1.4	23
23	Life without TTP: apparent absence of an important anti-inflammatory protein in birds. American Journal of Physiology - Regulatory Integrative and Comparative Physiology, 2013, 305, R689-R700.	0.9	20
24	Optimized Method for Robust Transcriptome Profiling of Minute Tissues Using Laser Capture Microdissection and Low-Input RNA-Seq. Frontiers in Molecular Neuroscience, 2017, 10, 185.	1.4	19
25	MyD88-dependent dendritic and epithelial cell crosstalk orchestrates immune responses to allergens. Mucosal Immunology, 2018, 11, 796-810.	2.7	18
26	Ancestry-dependent gene expression correlates with reprogramming to pluripotency and multiple dynamic biological processes. Science Advances, 2020, 6, .	4.7	16
27	Pulmonary Function and Blood DNA Methylation: A Multiancestry Epigenome-Wide Association Meta-analysis. American Journal of Respiratory and Critical Care Medicine, 2022, 206, 321-336.	2.5	15
28	Multimodal regulatory elements within a hormone-specific super enhancer control a heterogeneous transcriptional response. Molecular Cell, 2022, 82, 803-815.e5.	4.5	14
29	Effects of Combined Tristetraprolin/Tumor Necrosis Factor Receptor Deficiency on the Splenic Transcriptome. Molecular and Cellular Biology, 2016, 36, 1395-1411.	1.1	7
30	Lrp1 Regulation of Pulmonary Function. Follow-Up of Human GWAS in Mice. American Journal of Respiratory Cell and Molecular Biology, 2021, 64, 368-378.	1.4	7
31	Differential Activity and Utilization of Glucocorticoid Receptor Binding Sites Yields Transcriptional Heterogeneity. Journal of the Endocrine Society, 2021, 5, A499-A500.	0.1	1
32	Abstract 3560: The chromatin architecture at gucocorticoid receptor binding sites , 2013, , .		0