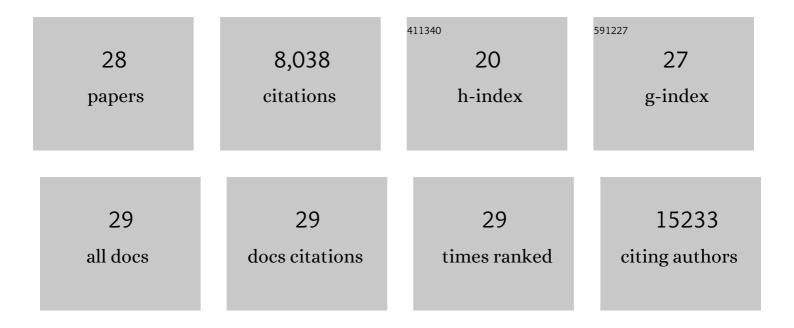
## Tae Hoon Kim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9072522/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Oxidative Stress and the Intersection of Oncogenic Signaling and Metabolism in Squamous Cell Carcinomas. Cells, 2021, 10, 606.	1.8	3
2	YAP and TAZ maintain PROX1 expression in the developing lymphatic and lymphovenous valves in response to VEGF-C signaling. Development (Cambridge), 2020, 147, .	1.2	28
3	The S-phase-induced lncRNA SUNO1 promotes cell proliferation by controlling YAP1/Hippo signaling pathway. ELife, 2020, 9, .	2.8	21
4	p63 and SOX2 Dictate Glucose Reliance and Metabolic Vulnerabilities in Squamous Cell Carcinomas. Cell Reports, 2019, 28, 1860-1878.e9.	2.9	68
5	Electrophysiological and transcriptomic correlates of neuropathic pain in human dorsal root ganglion neurons. Brain, 2019, 142, 1215-1226.	3.7	198
6	Global transcriptional activity dynamics reveal functional enhancer RNAs. Genome Research, 2018, 28, 1799-1811.	2.4	34
7	Complementary Wnt Sources Regulate Lymphatic Vascular Development via PROX1-Dependent Wnt/β-Catenin Signaling. Cell Reports, 2018, 25, 571-584.e5.	2.9	55
8	Biotin tagging of MeCP2 in mice reveals contextual insights into the Rett syndrome transcriptome. Nature Medicine, 2017, 23, 1203-1214.	15.2	102
9	Chromosome Conformation Capture for Research on Innate Antiviral Immunity. Methods in Molecular Biology, 2017, 1656, 195-208.	0.4	0
10	Mechanotransduction activates canonical Wnt/β-catenin signaling to promote lymphatic vascular patterning and the development of lymphatic and lymphovenous valves. Genes and Development, 2016, 30, 1454-1469.	2.7	121
11	Histone Deacetylases Positively Regulate Transcription through the Elongation Machinery. Cell Reports, 2015, 13, 1444-1455.	2.9	138
12	Boundary Associated Long Noncoding RNA Mediates Long-Range Chromosomal Interactions. PLoS ONE, 2015, 10, e0136104.	1.1	7
13	A novel virus-inducible enhancer of the interferon-β gene with tightly linked promoter and enhancer activities. Nucleic Acids Research, 2014, 42, 12537-12554.	6.5	30
14	Oct-1 Regulates IL-17 Expression by Directing Interchromosomal Associations in Conjunction with CTCF in T Cells. Molecular Cell, 2014, 54, 56-66.	4.5	44
15	Conserved, developmentally regulated mechanism couples chromosomal looping and heterochromatin barrier activity at the homeobox gene A locus. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7391-7396.	3.3	101
16	Genomic imprinting at a boundary element flanking the SDHD locus. Human Molecular Genetics, 2011, 20, 4452-4461.	1.4	21
17	From sextant to GPS: Twentyâ€five years of mapping the genome with ChIP. Journal of Cellular Biochemistry, 2009, 107, 6-10.	1.2	7
18	Global position analysis of the <i>Pseudomonas aeruginosa</i> quorumâ€sensing transcription factor LasR. Molecular Microbiology, 2009, 73, 1072-1085.	1.2	207

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19	Towards a genome-wide reconstruction of cis-regulatory networks in the human genome. Seminars in Cell and Developmental Biology, 2009, 20, 842-848.	2.3	11
20	Analysis of the Vertebrate Insulator Protein CTCF-Binding Sites in the Human Genome. Cell, 2007, 128, 1231-1245.	13.5	910
21	ChlP hip for Genomeâ€Wide Analysis of Protein Binding in Mammalian Cells. Current Protocols in Molecular Biology, 2007, 79, Unit 21.13.	2.9	22
22	Identification and analysis of functional elements in 1% of the human genome by the ENCODE pilot project. Nature, 2007, 447, 799-816.	13.7	4,709
23	An all-round view of eukaryotic transcription. Genome Biology, 2006, 7, 323.	13.9	2
24	Genome-Wide Analysis of Protein-DNA Interactions. Annual Review of Genomics and Human Genetics, 2006, 7, 81-102.	2.5	148
25	Positive Inter-Regulation between Î <sup>2</sup> -Catenin/T Cell Factor-4 Signaling and Endothelin-1 Signaling Potentiates Proliferation and Survival of Prostate Cancer Cells. Molecular Pharmacology, 2006, 69, 520-531.	1.0	44
26	β-Catenin activates the growth factor endothelin-1 in colon cancer cells. Oncogene, 2005, 24, 597-604.	2.6	90
27	A high-resolution map of active promoters in the human genome. Nature, 2005, 436, 876-880.	13.7	841
28	Direct isolation and identification of promoters in the human genome. Genome Research, 2005, 15, 830-839.	2.4	76