## Ofir Hakim

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7211240/publications.pdf

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414303 361296 2,785 32 20 32 h-index citations g-index papers 35 35 35 5259 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Ep300 sequestration to functionally distinct glucocorticoid receptor binding loci underlie rapid gene activation and repression. Nucleic Acids Research, 2022, 50, 6702-6714.	6.5	7
2	Genomic retargeting of p53 and CTCF is associated with transcriptional changes during oncogenic HRas-induced transformation. Communications Biology, 2020, 3, 696.	2.0	4
3	The Position and Complex Genomic Architecture of Plant T-DNA Insertions Revealed by 4SEE. International Journal of Molecular Sciences, 2020, 21, 2373.	1.8	7
4	Chromatin dynamics enable transcriptional rhythms in the cnidarian Nematostella vectensis. PLoS Genetics, 2019, 15, e1008397.	1.5	9
5	Regulatory chromatin landscape in Arabidopsis thaliana roots uncovered by coupling INTACT and ATAC-seq. Plant Methods, 2018, 14, 113.	1.9	45
6	Genomic effects of glucocorticoids. Protoplasma, 2017, 254, 1175-1185.	1.0	18
7	OCT4 impedes cell fate redirection by the melanocyte lineage master regulator MITF in mouse ESCs. Nature Communications, 2017, 8, 1022.	5.8	6
8	Hierarchical role for transcription factors and chromatin structure in genome organization along adipogenesis. FEBS Journal, 2017, 284, 3230-3244.	2.2	10
9	Mapping Genome-wide Accessible Chromatin in Primary Human T Lymphocytes by ATAC-Seq. Journal of Visualized Experiments, 2017, , .	0.2	8
10	Neuronal CTCF Is Necessary for Basal and Experience-Dependent Gene Regulation, Memory Formation, and Genomic Structure of BDNF and Arc. Cell Reports, 2016, 17, 2418-2430.	2.9	78
11	Cajal bodies are linked to genome conformation. Nature Communications, 2016, 7, 10966.	5.8	127
12	FIE, a nuclear PRC2 protein, forms cytoplasmic complexes in <i>Arabidopsis thaliana </i> Experimental Botany, 2016, 67, 6111-6123.	2.4	16
13	Comparative analysis of T4 DNA ligases and DNA polymerases used in chromosome conformation capture assays. BioTechniques, 2015, 58, 195-199.	0.8	8
14	Dynamics of chromatin accessibility and long-range interactions in response to glucocorticoid pulsing. Genome Research, 2015, 25, 845-857.	2.4	86
15	Epigenetic targeting of activation-induced cytidine deaminase. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 18667-18672.	3.3	48
16	B Cell Super-Enhancers and Regulatory Clusters Recruit AID Tumorigenic Activity. Cell, 2014, 159, 1524-1537.	13.5	234
17	3D view of chromosomes, DNA damage, and translocations. Current Opinion in Genetics and Development, 2014, 25, 118-125.	1.5	11
18	A Macrohistone Variant Links Dynamic Chromatin Compaction to BRCA1-Dependent Genome Maintenance. Cell Reports, 2014, 8, 1049-1062.	2.9	179

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19	A Genome-wide Map of CTCF Multivalency Redefines the CTCF Code. Cell Reports, 2013, 3, 1678-1689.	2.9	270
20	Interactome Maps of Mouse Gene Regulatory Domains Reveal Basic Principles of Transcriptional Regulation. Cell, 2013, 155, 1507-1520.	13.5	299
21	Cycles in spatial and temporal chromosomal organization driven by the circadian clock. Nature Structural and Molecular Biology, 2013, 20, 1206-1213.	3.6	110
22	Spatial congregation of STAT binding directs selective nuclear architecture during T-cell functional differentiation. Genome Research, 2013, 23, 462-472.	2.4	44
23	The Origin of B Cell Recurrent Chromosomal Translocations: Proximity versus DNA Damage. Molecular Cell, 2013, 51, 275-276.	4.5	5
24	Mouse model of endemic Burkitt translocations reveals the long-range boundaries of <i>lg</i> -mediated oncogene deregulation. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 10972-10977.	3.3	25
25	DNA damage defines sites of recurrent chromosomal translocations in B lymphocytes. Nature, 2012, 484, 69-74.	13.7	186
26	SnapShot: Chromosome Conformation Capture. Cell, 2012, 148, 1068-1068.e2.	13.5	53
27	Diverse gene reprogramming events occur in the same spatial clusters of distal regulatory elements. Genome Research, 2011, 21, 697-706.	2.4	132
28	Early Th1 Cell Differentiation Is Marked by a Tfh Cell-like Transition. Immunity, 2011, 35, 919-931.	6.6	364
29	3D shortcuts to gene regulation. Current Opinion in Cell Biology, 2010, 22, 305-313.	2.6	33
30	Glucocorticoid Receptor Activation of the Ciz1-Lcn2 Locus by Long Range Interactions. Journal of Biological Chemistry, 2009, 284, 6048-6052.	1.6	61
31	FIE and CURLY LEAF polycomb proteins interact in the regulation of homeobox gene expression during sporophyte development. Plant Journal, 2004, 37, 707-719.	2.8	229
32	From flour to flower: how Polycomb group proteins influence multiple aspects of plant development. Trends in Plant Science, 2003, 8, 439-445.	4.3	68