

# Yaniv Lubling

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8838136/publications.pdf>

Version: 2024-02-01

18  
papers

6,435  
citations

471509

17  
h-index

839539

18  
g-index

23  
all docs

23  
docs citations

23  
times ranked

10697  
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-cell Hi-C reveals cell-to-cell variability in chromosome structure. <i>Nature</i> , 2013, 502, 59-64.	27.8	1,347
2	Multiscale 3D Genome Rewiring during Mouse Neural Development. <i>Cell</i> , 2017, 171, 557-572.e24.	28.9	1,060
3	Integrative Analysis of the <i>Caenorhabditis elegans</i> Genome by the modENCODE Project. <i>Science</i> , 2010, 330, 1775-1787.	12.6	912
4	Dysfunctional CD8 T Cells Form a Proliferative, Dynamically Regulated Compartment within Human Melanoma. <i>Cell</i> , 2019, 176, 775-789.e18.	28.9	760
5	Cell-cycle dynamics of chromosomal organization at single-cell resolution. <i>Nature</i> , 2017, 547, 61-67.	27.8	636
6	Distinct Modes of Regulation by Chromatin Encoded through Nucleosome Positioning Signals. <i>PLoS Computational Biology</i> , 2008, 4, e1000216.	3.2	393
7	Single-cell characterization of haematopoietic progenitors and their trajectories in homeostasis and perturbed haematopoiesis. <i>Nature Cell Biology</i> , 2018, 20, 836-846.	10.3	267
8	MetaCell: analysis of single-cell RNA-seq data using K-nn graph partitions. <i>Genome Biology</i> , 2019, 20, 206.	8.8	218
9	Single-cell Hi-C for genome-wide detection of chromatin interactions that occur simultaneously in a single cell. <i>Nature Protocols</i> , 2015, 10, 1986-2003.	12.0	135
10	Gene expression divergence in yeast is coupled to evolution of DNA-encoded nucleosome organization. <i>Nature Genetics</i> , 2009, 41, 438-445.	21.4	132
11	Capturing pairwise and multi-way chromosomal conformations using chromosomal walks. <i>Nature</i> , 2016, 540, 296-300.	27.8	128
12	An integrated open framework for thermodynamics of reactions that combines accuracy and coverage. <i>Bioinformatics</i> , 2012, 28, 2037-2044.	4.1	108
13	p53 binds preferentially to genomic regions with high DNA-encoded nucleosome occupancy. <i>Genome Research</i> , 2010, 20, 1361-1368.	5.5	86
14	UMI-4C for quantitative and targeted chromosomal contact profiling. <i>Nature Methods</i> , 2016, 13, 685-691.	19.0	78
15	Compensation for differences in gene copy number among yeast ribosomal proteins is encoded within their promoters. <i>Genome Research</i> , 2011, 21, 2114-2128.	5.5	51
16	Nucleosome maps of the human cytomegalovirus genome reveal a temporal switch in chromatin organization linked to a major IE protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 13126-13131.	7.1	43
17	High nucleosome occupancy is encoded at X-linked gene promoters in <i>C. elegans</i> . <i>Genome Research</i> , 2011, 21, 237-244.	5.5	35
18	Single-cell analysis of regions of interest (SCARI) using a photosensitive tag. <i>Nature Chemical Biology</i> , 2021, 17, 1139-1147.	8.0	13