

# Danny Leung

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3579654/publications.pdf>

Version: 2024-02-01

10  
papers

1,582  
citations

1163117

8  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

2996  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proviral silencing in embryonic stem cells requires the histone methyltransferase ESET. <i>Nature</i> , 2010, 464, 927-931.	27.8	681
2	DNA Methylation and SETDB1/H3K9me3 Regulate Predominantly Distinct Sets of Genes, Retroelements, and Chimeric Transcripts in mESCs. <i>Cell Stem Cell</i> , 2011, 8, 676-687.	11.1	427
3	Integrative analysis of haplotype-resolved epigenomes across human tissues. <i>Nature</i> , 2015, 518, 350-354.	27.8	201
4	Replication timing maintains the global epigenetic state in human cells. <i>Science</i> , 2021, 372, 371-378.	12.6	103
5	Regulation of DNA methylation turnover at LTR retrotransposons and imprinted loci by the histone methyltransferase Setdb1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 6690-6695.	7.1	89
6	Impact of flanking chromosomal sequences on localization and silencing by the human non-coding RNA XIST. <i>Genome Biology</i> , 2015, 16, 208.	8.8	36
7	Humanizing the yeast origin recognition complex. <i>Nature Communications</i> , 2021, 12, 33.	12.8	24
8	DNA Repair Inhibition Leads to Active Export of Repetitive Sequences to the Cytoplasm Triggering an Inflammatory Response. <i>Journal of Neuroscience</i> , 2021, 41, 9286-9307.	3.6	13
9	Mouse strain-specific polymorphic provirus functions as cis-regulatory element leading to epigenomic and transcriptomic variations. <i>Nature Communications</i> , 2021, 12, 6462.	12.8	4
10	Epigenomic and transcriptomic analysis of chronic inflammatory diseases. <i>Genes and Genomics</i> , 2021, 43, 227-236.	1.4	3