

# Kaia Mattioli

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4567655/publications.pdf>

Version: 2024-02-01

6  
papers

405  
citations

1683354

5  
h-index

1872312

6  
g-index

10  
all docs

10  
docs citations

10  
times ranked

754  
citing authors

#	ARTICLE	IF	CITATIONS
1	Chromatin environment, transcriptional regulation, and splicing distinguish lincRNAs and mRNAs. <i>Genome Research</i> , 2017, 27, 27-37.	2.4	207
2	High-throughput functional analysis of lincRNA core promoters elucidates rules governing tissue specificity. <i>Genome Research</i> , 2019, 29, 344-355.	2.4	100
3	Cis and trans effects differentially contribute to the evolution of promoters and enhancers. <i>Genome Biology</i> , 2020, 21, 210.	3.8	35
4	Resolving mechanisms of immune-mediated disease in primary <scp>CD</scp> 4 T cells. <i>EMBO Molecular Medicine</i> , 2020, 12, e12112.	3.3	30
5	Whole genome sequencing delineates regulatory, copy number, and cryptic splice variants in early onset cardiomyopathy. <i>Npj Genomic Medicine</i> , 2022, 7, 18.	1.7	14
6	Genome-wide CRISPR interference screen identifies long non-coding RNA loci required for differentiation and pluripotency. <i>PLoS ONE</i> , 2021, 16, e0252848.	1.1	12