

Viviana I Risca

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8459771/publications.pdf>

Version: 2024-02-01

10
papers

2,102
citations

1307594

7
h-index

1372567

10
g-index

18
all docs

18
docs citations

18
times ranked

5668
citing authors

#	ARTICLE	IF	CITATIONS
1	An improved ATAC-seq protocol reduces background and enables interrogation of frozen tissues. <i>Nature Methods</i> , 2017, 14, 959-962.	19.0	1,653
2	Variable chromatin structure revealed by in situ spatially correlated DNA cleavage mapping. <i>Nature</i> , 2017, 541, 237-241.	27.8	132
3	Chromatin-associated RNA sequencing (ChAR-seq) maps genome-wide RNA-to-DNA contacts. <i>ELife</i> , 2018, 7, .	6.0	121
4	Unraveling the 3D genome: genomics tools for multiscale exploration. <i>Trends in Genetics</i> , 2015, 31, 357-372.	6.7	62
5	BLIMP1 Induces Transient Metastatic Heterogeneity in Pancreatic Cancer. <i>Cancer Discovery</i> , 2017, 7, 1184-1199.	9.4	53
6	The dynamic, combinatorial cis-regulatory lexicon of epidermal differentiation. <i>Nature Genetics</i> , 2021, 53, 1564-1576.	21.4	45
7	Nucleosome Orientation Map Finds Two New Chromatin Folding Motifs. <i>Cell</i> , 2019, 176, 412-413.	28.9	16
8	A Conditional System to Specifically Link Disruption of Protein-Coding Function with Reporter Expression in Mice. <i>Cell Reports</i> , 2014, 7, 2078-2086.	6.4	9
9	Chromatin-associated RNA Sequencing (ChAR-seq). <i>Current Protocols in Molecular Biology</i> , 2019, 126, e87.	2.9	7
10	Beyond the Linear Genome: Paired-End Sequencing as a Biophysical Tool. <i>Trends in Cell Biology</i> , 2015, 25, 716-719.	7.9	1