

Scott T Younger

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5380960/publications.pdf>

Version: 2024-02-01

14
papers

793
citations

933447

10
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

1767
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiplexable, locus-specific targeting of long RNAs with CRISPR-Display. <i>Nature Methods</i> , 2015, 12, 664-670.	19.0	268
2	Deubiquitinases Maintain Protein Homeostasis and Survival of Cancer Cells upon Glutathione Depletion. <i>Cell Metabolism</i> , 2019, 29, 1166-1181.e6.	16.2	121
3	Cells Lacking the <i>RB1</i> Tumor Suppressor Gene Are Hyperdependent on Aurora B Kinase for Survival. <i>Cancer Discovery</i> , 2019, 9, 230-247.	9.4	119
4	Integrative genomic analysis reveals widespread enhancer regulation by p53 in response to DNA damage. <i>Nucleic Acids Research</i> , 2015, 43, 4447-4462.	14.5	84
5	p53 regulates enhancer accessibility and activity in response to DNA damage. <i>Nucleic Acids Research</i> , 2017, 45, 9889-9900.	14.5	61
6	Genomic answers for children: Dynamic analyses of >1000 pediatric rare disease genomes. <i>Genetics in Medicine</i> , 2022, 24, 1336-1348.	2.4	37
7	'Lnc'-ing enhancers to MYC regulation. <i>Cell Research</i> , 2014, 24, 643-644.	12.0	23
8	SMAD4 represses FOSL1 expression and pancreatic cancer metastatic colonization. <i>Cell Reports</i> , 2021, 36, 109443.	6.4	23
9	Pooled Lentiviral Delivery Genetic Screens. <i>Current Protocols in Molecular Biology</i> , 2018, 121, 32.1.1-32.1.21.	2.9	20
10	Identification of functional regulatory elements in the human genome using pooled CRISPR screens. <i>BMC Genomics</i> , 2020, 21, 107.	2.8	12
11	Expediting rare disease diagnosis: a call to bridge the gap between clinical and functional genomics. <i>Molecular Medicine</i> , 2020, 26, 117.	4.4	10
12	Silent pericentromeric repeats speak out. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 15008-15009.	7.1	8
13	FOXC2 Autoregulates Its Expression in the Pulmonary Endothelium After Endotoxin Stimulation in a Histone Acetylation-Dependent Manner. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 657662.	3.7	6
14	Massively parallel identification of functionally consequential noncoding genetic variants in undiagnosed rare disease patients. <i>Scientific Reports</i> , 2022, 12, 7576.	3.3	1