

Shipra Bhatia

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3397176/publications.pdf>

Version: 2024-02-01

11
papers

421
citations

1163117

8
h-index

1281871

11
g-index

13
all docs

13
docs citations

13
times ranked

875
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification and functional modelling of plausibly causative cis-regulatory variants in a highly-selected cohort with X-linked intellectual disability. PLoS ONE, 2021, 16, e0256181.	2.5	3
2	Quantitative spatial and temporal assessment of regulatory element activity in zebrafish. ELife, 2021, 10, .	6.0	14
3	Zebrafish: A Powerful Model for Understanding the Functional Relevance of Noncoding Region Mutations in Human Genetic Diseases. Biomedicines, 2019, 7, 71.	3.2	6
4	Lampreys, the jawless vertebrates, contain three Pax6 genes with distinct expression in eye, brain and pancreas. Scientific Reports, 2019, 9, 19559.	3.3	23
5	Psp1/p52 regulates posterior Hoxa genes through activation of lncRNA Hottip. PLoS Genetics, 2017, 13, e1006677.	3.5	30
6	SBE6: a novel long-range enhancer involved in driving sonic hedgehog expression in neural progenitor cells. Open Biology, 2016, 6, 160197.	3.6	17
7	Functional Assessment of Disease-Associated Regulatory Variants In Vivo Using a Versatile Dual Colour Transgenesis Strategy in Zebrafish. PLoS Genetics, 2015, 11, e1005193.	3.5	31
8	Disruption of SATB2 or its long-range cis-regulation by SOX9 causes a syndromic form of Pierre Robin sequence. Human Molecular Genetics, 2014, 23, 2569-2579.	2.9	51
9	Disruption of long-range gene regulation in human genetic disease: a kaleidoscope of general principles, diverse mechanisms and unique phenotypic consequences. Human Genetics, 2014, 133, 815-845.	3.8	32
10	Disruption of Autoregulatory Feedback by a Mutation in a Remote, Ultraconserved PAX6 Enhancer Causes Aniridia. American Journal of Human Genetics, 2013, 93, 1126-1134.	6.2	172
11	Sequencing of Pax6 Loci from the Elephant Shark Reveals a Family of Pax6 Genes in Vertebrate Genomes, Forged by Ancient Duplications and Divergences. PLoS Genetics, 2013, 9, e1003177.	3.5	40