

Zeeshan Fazal

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7234330/publications.pdf>

Version: 2024-02-01

14
papers

343
citations

1040056

9
h-index

1281871

11
g-index

15
all docs

15
docs citations

15
times ranked

574
citing authors

#	ARTICLE	IF	CITATIONS
1	Towards a Unified Understanding of Lithium Action in Basic Biology and its Significance for Applied Biology. <i>Journal of Membrane Biology</i> , 2017, 250, 587-604.	2.1	103
2	POLYAR, a new computer program for prediction of poly(A) sites in human sequences. <i>BMC Genomics</i> , 2010, 11, 646.	2.8	51
3	Mechanisms of cisplatin sensitivity and resistance in testicular germ cell tumors. , 2019, 2, 580-594.		35
4	Epigenetic Remodeling through Downregulation of Polycomb Repressive Complex 2 Mediates Chemotherapy Resistance in Testicular Germ Cell Tumors. <i>Cancers</i> , 2019, 11, 796.	3.7	25
5	Toward a Mechanistic Understanding of Poly- and Perfluoroalkylated Substances and Cancer. <i>Cancers</i> , 2022, 14, 2919.	3.7	25
6	A phase 1 study of combined guadecitabine and cisplatin in platinum refractory germ cell cancer. <i>Cancer Medicine</i> , 2021, 10, 156-163.	2.8	23
7	Hypermethylation and global remodelling of DNA methylation is associated with acquired cisplatin resistance in testicular germ cell tumours. <i>Epigenetics</i> , 2021, 16, 1071-1084.	2.7	21
8	Between a Rock and a Hard Place: An Epigenetic-Centric View of Testicular Germ Cell Tumors. <i>Cancers</i> , 2021, 13, 1506.	3.7	18
9	Three-Dimensional Encapsulation of <i>Saccharomyces cerevisiae</i> in Silicate Matrices Creates Distinct Metabolic States as Revealed by Gene Chip Analysis. <i>ACS Nano</i> , 2017, 11, 3560-3575.	14.6	17
10	Reciprocal epigenetic remodeling controls testicular cancer hypersensitivity to hypomethylating agents and chemotherapy. <i>Molecular Oncology</i> , 2022, 16, 683-698.	4.6	10
11	Using Optimal F-Measure and Random Resampling in Gene Ontology Enrichment Calculations. <i>Frontiers in Applied Mathematics and Statistics</i> , 2019, 5, .	1.3	3
12	Model of ion intensity from tandem mass spectra for improved peptide identification and simulation. , 2011, , .		1
13	Multifactorial Understanding of Ion Abundance in Tandem Mass Spectrometry Experiments. <i>Journal of Proteomics and Bioinformatics</i> , 2013, 06, 23-29.	0.4	1
14	Hierarchical modeling of alternative exon usage associations with survival. , 2011, , .		0