Shaoli Das

List of Publications by Year
in descending order

22 papers	1,248 citations	759055 12 h-index	839398 18 g-index
23 all docs	23 docs citations	23 times ranked	2186 citing authors

#	Article	IF	Citations
1	Circ2Traits: a comprehensive database for circular RNA potentially associated with disease and traits. Frontiers in Genetics, 2013, 4, 283.	1.1	417
2	Competing Endogenous RNA: The Key to Posttranscriptional Regulation. Scientific World Journal, The, 2014, 2014, 1-6.	0.8	242
3	lnCeDB: Database of Human Long Noncoding RNA Acting as Competing Endogenous RNA. PLoS ONE, 2014, 9, e98965.	1.1	150
4	Long Noncoding RNAs: New Players in the Molecular Mechanism for Maintenance and Differentiation of Pluripotent Stem Cells. Stem Cells and Development, 2013, 22, 2240-2253.	1.1	93
5	Dynamic changes in global microRNAome and transcriptome reveal complex miRNA-mRNA regulated host response to Japanese Encephalitis Virus in microglial cells. Scientific Reports, 2016, 6, 20263.	1.6	54
6	Radiation-Induced Long Noncoding RNAs in a Mouse Model after Whole-Body Irradiation. Radiation Research, 2018, 189, 251.	0.7	44
7	HumanViCe: host ceRNA network in virus infected cells in human. Frontiers in Genetics, 2014, 5, 249.	1.1	41
8	DiscoverSL: an R package for multi-omic data driven prediction of synthetic lethality in cancers. Bioinformatics, 2019, 35, 701-702.	1.8	41
9	Cancer-Specific Immune Prognostic Signature in Solid Tumors and Its Relation to Immune Checkpoint Therapies. Cancers, 2020, 12, 2476.	1.7	39
10	Correlation of altered expression of a long non-coding RNA, NEAT1, in peripheral blood mononuclear cells with dengue disease progression. Journal of Infection, 2017, 75, 541-554.	1.7	30
11	HNOCDB: A comprehensive database of genes and miRNAs relevant to head and neck and oral cancer. Oral Oncology, 2012, 48, 117-119.	0.8	22
12	SL-BioDP: Multi-Cancer Interactive Tool for Prediction of Synthetic Lethality and Response to Cancer Treatment. Cancers, 2019, 11, 1682.	1.7	20
13	Pan-Cancer Analysis of Potential Synthetic Lethal Drug Targets Specific to Alterations in DNA Damage Response. Frontiers in Oncology, 2019, 9, 1136.	1.3	14
14	Long intergenic noncoding RNA profiles of pheochromocytoma and paraganglioma: A novel prognostic biomarker. International Journal of Cancer, 2020, 146, 2326-2335.	2.3	14
15	SeedSeq: Off-Target Transcriptome Database. BioMed Research International, 2013, 2013, 1-9.	0.9	8
16	miRepress: modelling gene expression regulation by microRNA with non-conventional binding sites. Scientific Reports, 2016, 6, 22334.	1.6	7
17	An siRNA designing tool with a unique functional off-target filtering approach. Journal of Biomolecular Structure and Dynamics, 2013, 31, 1343-1357.	2.0	4
18	Alteration of MicroRNA Biogenesis Pathways in Cancers. , 2018, , 47-58.		2

#	Article	IF	Citations
19	Computational Approaches for Designing Efficient and Specific siRNAs. , 2012, , .		1
20	A long noncoding RNAâ€"microRNA expression signature predicts metastatic signature in pheochromocytomas and paragangliomas. Endocrine, 2021, , 1.	1.1	1
21	Automated Classification of Disease Patterns from Echo-cardiography Images Based on Shape Features of the Left Ventricle. , 2010, , .		0
22	Synthetic Lethal Drug Combinations Targeting Proteasome and Histone Deacetylase Inhibitors in TP53-Mutated Cancers. , 2020, 1, 42-47.		0