

# Shaoli Das

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4037118/publications.pdf>

Version: 2024-02-01

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	A long noncoding RNAâ€™microRNA expression signature predicts metastatic signature in pheochromocytomas and paragangliomas. <i>Endocrine</i> , 2021, , 1.	1.1	1
2	Long intergenic noncoding RNA profiles of pheochromocytoma and paraganglioma: A novel prognostic biomarker. <i>International Journal of Cancer</i> , 2020, 146, 2326-2335.	2.3	14
3	Cancer-Specific Immune Prognostic Signature in Solid Tumors and Its Relation to Immune Checkpoint Therapies. <i>Cancers</i> , 2020, 12, 2476.	1.7	39
4	Synthetic Lethal Drug Combinations Targeting Proteasome and Histone Deacetylase Inhibitors in TP53-Mutated Cancers. , 2020, 1, 42-47.		0
5	DiscoverSL: an R package for multi-omic data driven prediction of synthetic lethality in cancers. <i>Bioinformatics</i> , 2019, 35, 701-702.	1.8	41
6	SL-BioDP: Multi-Cancer Interactive Tool for Prediction of Synthetic Lethality and Response to Cancer Treatment. <i>Cancers</i> , 2019, 11, 1682.	1.7	20
7	Pan-Cancer Analysis of Potential Synthetic Lethal Drug Targets Specific to Alterations in DNA Damage Response. <i>Frontiers in Oncology</i> , 2019, 9, 1136.	1.3	14
8	Radiation-Induced Long Noncoding RNAs in a Mouse Model after Whole-Body Irradiation. <i>Radiation Research</i> , 2018, 189, 251.	0.7	44
9	Alteration of MicroRNA Biogenesis Pathways in Cancers. , 2018, , 47-58.		2
10	Correlation of altered expression of a long non-coding RNA, NEAT1, in peripheral blood mononuclear cells with dengue disease progression. <i>Journal of Infection</i> , 2017, 75, 541-554.	1.7	30
11	miRepress: modelling gene expression regulation by microRNA with non-conventional binding sites. <i>Scientific Reports</i> , 2016, 6, 22334.	1.6	7
12	Dynamic changes in global microRNAome and transcriptome reveal complex miRNA-mRNA regulated host response to Japanese Encephalitis Virus in microglial cells. <i>Scientific Reports</i> , 2016, 6, 20263.	1.6	54
13	InCeDB: Database of Human Long Noncoding RNA Acting as Competing Endogenous RNA. <i>PLoS ONE</i> , 2014, 9, e98965.	1.1	150
14	Competing Endogenous RNA: The Key to Posttranscriptional Regulation. <i>Scientific World Journal</i> , The, 2014, 2014, 1-6.	0.8	242
15	HumanViCe: host ceRNA network in virus infected cells in human. <i>Frontiers in Genetics</i> , 2014, 5, 249.	1.1	41
16	An siRNA designing tool with a unique functional off-target filtering approach. <i>Journal of Biomolecular Structure and Dynamics</i> , 2013, 31, 1343-1357.	2.0	4
17	Long Noncoding RNAs: New Players in the Molecular Mechanism for Maintenance and Differentiation of Pluripotent Stem Cells. <i>Stem Cells and Development</i> , 2013, 22, 2240-2253.	1.1	93
18	SeedSeq: Off-Target Transcriptome Database. <i>BioMed Research International</i> , 2013, 2013, 1-9.	0.9	8

#	ARTICLE	IF	CITATIONS
19	Circ2Traits: a comprehensive database for circular RNA potentially associated with disease and traits. <i>Frontiers in Genetics</i> , 2013, 4, 283.	1.1	417
20	Computational Approaches for Designing Efficient and Specific siRNAs. , 2012, , .		1
21	HNOCDDB: A comprehensive database of genes and miRNAs relevant to head and neck and oral cancer. <i>Oral Oncology</i> , 2012, 48, 117-119.	0.8	22
22	Automated Classification of Disease Patterns from Echo-cardiography Images Based on Shape Features of the Left Ventricle. , 2010, , .		0