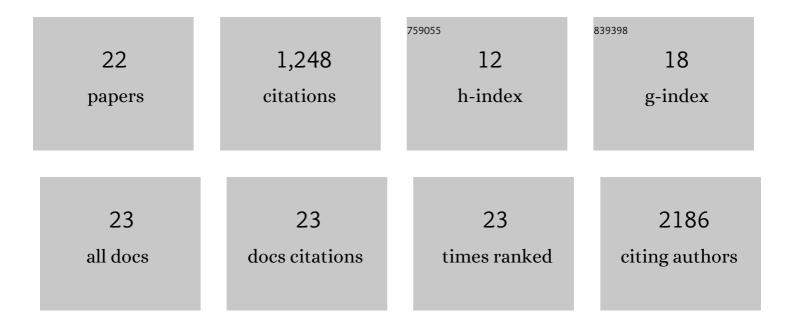
## Shaoli Das

List of Publications by Year in descending order

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**SHAOLI DAS** 

| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | A long noncoding RNA–microRNA expression signature predicts metastatic signature in pheochromocytomas and paragangliomas. Endocrine, 2021, , 1.   | 1.1 | 1         |
| 2  | Long intergenic noncoding RNA profiles of pheochromocytoma and paraganglioma: A novel prognostic biomarker. International Journal of Cancer, 2020, 146, 2326-2335.                                | 2.3 | 14        |
| 3  | Cancer-Specific Immune Prognostic Signature in Solid Tumors and Its Relation to Immune Checkpoint<br>Therapies. Cancers, 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.<br>Bioinformatics, 2019, 35, 701-702.   | 1.8 | 41        |
| 6  | SL-BioDP: Multi-Cancer Interactive Tool for Prediction of Synthetic Lethality and Response to Cancer<br>Treatment. Cancers, 2019, 11, 1682.   | 1.7 | 20        |
| 7  | Pan-Cancer Analysis of Potential Synthetic Lethal Drug Targets Specific to Alterations in DNA Damage<br>Response. Frontiers in Oncology, 2019, 9, 1136.   | 1.3 | 14        |
| 8  | Radiation-Induced Long Noncoding RNAs in a Mouse Model after Whole-Body Irradiation. Radiation<br>Research, 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. Journal of Infection, 2017, 75, 541-554.                | 1.7 | 30        |
| 11 | miRepress: modelling gene expression regulation by microRNA with non-conventional binding sites.<br>Scientific Reports, 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. Scientific Reports, 2016, 6, 20263. | 1.6 | 54        |
| 13 | InCeDB: Database of Human Long Noncoding RNA Acting as Competing Endogenous RNA. PLoS ONE,<br>2014, 9, e98965.  | 1.1 | 150       |
| 14 | Competing Endogenous RNA: The Key to Posttranscriptional Regulation. Scientific World Journal, The, 2014, 2014, 1-6.  | 0.8 | 242       |
| 15 | HumanViCe: host ceRNA network in virus infected cells in human. Frontiers in Genetics, 2014, 5, 249.  | 1.1 | 41        |
| 16 | An siRNA designing tool with a unique functional off-target filtering approach. Journal of<br>Biomolecular Structure and Dynamics, 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. Stem Cells and Development, 2013, 22, 2240-2253.                       | 1.1 | 93        |
| 18 | SeedSeq: Off-Target Transcriptome Database. BioMed Research International, 2013, 2013, 1-9.   | 0.9 | 8         |

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| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 19 | Circ2Traits: a comprehensive database for circular RNA potentially associated with disease and traits.<br>Frontiers in Genetics, 2013, 4, 283. | 1.1 | 417       |
| 20 | Computational Approaches for Designing Efficient and Specific siRNAs. , 2012, , .  |     | 1         |
| 21 | HNOCDB: A comprehensive database of genes and miRNAs relevant to head and neck and oral cancer.<br>Oral Oncology, 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         |