

# Souvik Ghosh

## List of Publications by Year in descending order

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Version: 2024-02-01

14  
papers

562  
citations

1039406

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1199166

12  
g-index

20  
all docs

20  
docs citations

20  
times ranked

1115  
citing authors

#	ARTICLE	IF	CITATIONS
1	CFIm-mediated alternative polyadenylation remodels cellular signaling and miRNA biogenesis. <i>Nucleic Acids Research</i> , 2022, 50, 3096-3114.	6.5	13
2	Transcriptional Enhancer Factor Domain Family member 4 Exerts an Oncogenic Role in Hepatocellular Carcinoma by Hippo-Independent Regulation of Heat Shock Protein 70 Family Members. <i>Hepatology Communications</i> , 2021, 5, 661-674.	2.0	13
3	The transcriptional landscape of a hepatoma cell line grown on scaffolds of extracellular matrix proteins. <i>BMC Genomics</i> , 2021, 22, 238.	1.2	0
4	GW182 Proteins Restrict Extracellular Vesicle-Mediated Export of MicroRNAs in Mammalian Cancer Cells. <i>Molecular and Cellular Biology</i> , 2021, 41, .	1.1	10
5	MiR-CLIP reveals <i>iso</i> -miR selective regulation in the miR-124 targetome. <i>Nucleic Acids Research</i> , 2021, 49, 25-37.	6.5	10
6	Mitochondria Control mTORC1 Activity Linked Compartmentalization of eIF4E to Regulate Extracellular Export of microRNAs. <i>Journal of Cell Science</i> , 2020, 133, .	1.2	1
7	Prevention of dsRNA-induced interferon signaling by AGO1x is linked to breast cancer cell proliferation. <i>EMBO Journal</i> , 2020, 39, e103922.	3.5	22
8	Single-cell mRNA profiling reveals the hierarchical response of miRNA targets to miRNA induction. <i>Molecular Systems Biology</i> , 2018, 14, e8266.	3.2	24
9	Discovery of physiological and cancer-related regulators of 3' UTR processing with KAPAC. <i>Genome Biology</i> , 2018, 19, 44.	3.8	54
10	3' End Sequencing Library Preparation with A-seq2. <i>Journal of Visualized Experiments</i> , 2017, , .	0.2	7
11	TFAP2A is a component of the ZEB1/2 network that regulates TGF $\beta$ 1-induced epithelial to mesenchymal transition. <i>Biology Direct</i> , 2017, 12, 8.	1.9	28
12	A comprehensive analysis of 3' end sequencing data sets reveals novel polyadenylation signals and the repressive role of heterogeneous ribonucleoprotein C on cleavage and polyadenylation. <i>Genome Research</i> , 2016, 26, 1145-1159.	2.4	196
13	Reversible HuR-miRNA binding controls extracellular export of miR-122 and augments stress response. <i>EMBO Reports</i> , 2016, 17, 1184-1203.	2.0	139
14	Polysome arrest restricts miRNA turnover by preventing exosomal export of miRNA in growth-retarded mammalian cells. <i>Molecular Biology of the Cell</i> , 2015, 26, 1072-1083.	0.9	41