

Suman Ghosal

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4201964/publications.pdf>

Version: 2024-02-01

15
papers

1,276
citations

840585

11
h-index

996849

15
g-index

15
all docs

15
docs citations

15
times ranked

2368
citing authors

#	ARTICLE	IF	CITATIONS
1	Circ2Traits: a comprehensive database for circular RNA potentially associated with disease and traits. <i>Frontiers in Genetics</i> , 2013, 4, 283.	1.1	417
2	Competing Endogenous RNA: The Key to Posttranscriptional Regulation. <i>Scientific World Journal</i> , The, 2014, 2014, 1-6.	0.8	242
3	InCeDB: Database of Human Long Noncoding RNA Acting as Competing Endogenous RNA. <i>PLoS ONE</i> , 2014, 9, e98965.	1.1	150
4	The Human CCHC-type Zinc Finger Nucleic Acid-Binding Protein Binds G-Rich Elements in Target mRNA Coding Sequences and Promotes Translation. <i>Cell Reports</i> , 2017, 18, 2979-2990.	2.9	106
5	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
6	Genotype-phenotype correlations in pheochromocytoma and paraganglioma: a systematic review and individual patient meta-analysis. <i>Endocrine-Related Cancer</i> , 2019, 26, 539-550.	1.6	87
7	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
8	HumanViCe: host ceRNA network in virus infected cells in human. <i>Frontiers in Genetics</i> , 2014, 5, 249.	1.1	41
9	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
10	HNOCDB: 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
11	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
12	SeedSeq: Off-Target Transcriptome Database. <i>BioMed Research International</i> , 2013, 2013, 1-9.	0.9	8
13	miRepress: modelling gene expression regulation by microRNA with non-conventional binding sites. <i>Scientific Reports</i> , 2016, 6, 22334.	1.6	7
14	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
15	A long noncoding RNA-microRNA expression signature predicts metastatic signature in pheochromocytomas and paragangliomas. <i>Endocrine</i> , 2021, , 1.	1.1	1