Ikhlak Ahmed

List of Publications by Year in descending order

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Version: 2024-02-01

26 papers	3,504 citations	430874 18 h-index	501196 28 g-index
37 all docs	37 docs citations	37 times ranked	8089 citing authors

#	Article	IF	Citations
1	Melanocortinâ€4 receptor complexity in energy homeostasis,obesity and drug development strategies. Diabetes, Obesity and Metabolism, 2022, 24, 583-598.	4.4	8
2	Qatar genome: Insights on genomics from the Middle East. Human Mutation, 2022, 43, 499-510.	2.5	29
3	A population study of clinically actionable genetic variation affecting drug response from the Middle East. Npj Genomic Medicine, 2022, 7, 10.	3.8	20
4	Insights Into the Role of CircRNAs: Biogenesis, Characterization, Functional, and Clinical Impact in Human Malignancies. Frontiers in Cell and Developmental Biology, 2021, 9, 617281.	3.7	53
5	Regulation of Circular RNA CircNFATC3 in Cancer Cells Alters Proliferation, Migration, and Oxidative Phosphorylation. Frontiers in Cell and Developmental Biology, 2021, 9, 595156.	3.7	19
6	Role of non-coding RNA networks in leukemia progression, metastasis and drug resistance. Molecular Cancer, 2020, 19, 57.	19.2	68
7	Cell Type-Specific TGF-Î² Mediated EMT in 3D and 2D Models and Its Reversal by TGF-Î² Receptor Kinase Inhibitor in Ovarian Cancer Cell Lines. International Journal of Molecular Sciences, 2019, 20, 3568.	4.1	19
8	Identification of human genetic variants controlling circular RNA expression. Rna, 2019, 25, 1765-1778.	3.5	23
9	High-resolution proteinâ€“protein interaction mapping using all-versus-all sequencing (AVA-Seq). Journal of Biological Chemistry, 2019, 294, 11549-11558.	3.4	10
10	Silencing of ANKRD12 circRNA induces molecular and functional changes associated with invasive phenotypes. BMC Cancer, 2019, 19, 565.	2.6	33
11	Genus-wide sequencing supports a two-locus model for sex-determination in Phoenix. Nature Communications, 2018, 9, 3969.	12.8	86
12	DET1-mediated degradation of a SAGA-like deubiquitination module controls H2Bub homeostasis. ELife, 2018, 7, .	6.0	63
13	Altered expression pattern of circular RNAs in primary and metastatic sites of epithelial ovarian carcinoma. Oncotarget, 2016, 7, 36366-36381.	1.8	148
14	Ensembl regulation resources. Database: the Journal of Biological Databases and Curation, 2016, 2016, bav119.	3.0	45
15	DNA DAMAGE BINDING PROTEIN2 Shapes the DNA Methylation Landscape. Plant Cell, 2016, 28, 2043-2059.	6.6	16
16	K-core decomposition of a protein domain co-occurrence network reveals lower cancer mutation rates for interior cores. Journal of Clinical Bioinformatics, 2015, 5, 1.	1.2	16
17	Characterization Of Circular Rnas In Ovarian Cancer. , 2014, , .		1
18	Insights into the role of DNA methylation in diatoms by genome-wide profiling in Phaeodactylum tricornutum. Nature Communications, 2013, 4, 2091.	12.8	113

#	Article	IF	Citations
19	Histone H2B Monoubiquitination Facilitates the Rapid Modulation of Gene Expression during Arabidopsis Photomorphogenesis. PLoS Genetics, 2012, 8, e1002825.	3.5	115
20	Ensembl 2013. Nucleic Acids Research, 2012, 41, D48-D55.	14.5	856
21	Integrative epigenomic mapping defines four main chromatin states in Arabidopsis. EMBO Journal, 2011, 30, 1928-1938.	7.8	600
22	Genome-wide evidence for local DNA methylation spreading from small RNA-targeted sequences in Arabidopsis. Nucleic Acids Research, 2011, 39, 6919-6931.	14.5	142
23	Integrative Transcript and Metabolite Analysis of Nutritionally Enhanced DE-ETIOLATED1 Downregulated Tomato Fruit. Plant Cell, 2010, 22, 1190-1215.	6.6	160
24	Utilizing linkage disequilibrium information from Indian Genome Variation Database for mapping mutations: SCA12 case study. Journal of Genetics, 2009, 88, 55-60.	0.7	4
25	Mapping Human Genetic Diversity in Asia. Science, 2009, 326, 1541-1545.	12.6	557
26	Genetic landscape of the people of India: a canvas for disease gene exploration. Journal of Genetics, 2008, 87, 3-20.	0.7	282