Basharat A Bhat

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/4381968/publications.pdf

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15	161	1651377 6	1336881 12
papers	citations	h-index	g-index

16	16	16	232
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Data Analysis of DNA Methylation Epigenome-Wide Association Studies (EWAS): A Guide to the Principles of Best Practice. Methods in Molecular Biology, 2022, 2458, 23-45.	0.4	5
2	Genome wide expression analysis of circular RNAs in mammary epithelial cells of cattle revealed difference in milk synthesis. PeerJ, 2022, 10, e13029.	0.9	7
3	Mammary epithelial cell transcriptome reveals potential roles of lncRNAs in regulating milk synthesis pathways in Jersey and Kashmiri cattle. BMC Genomics, 2022, 23, 176.	1.2	4
4	Tuberous sclerosis complex: a complex case.. Cold Spring Harbor Molecular Case Studies, 2022, 8, .	0.7	1
5	Identification of potential key genes and pathways associated with the Pashmina fiber initiation using RNA-Seq and integrated bioinformatics analysis. Scientific Reports, 2021, 11, 1766.	1.6	14
6	Changthangi Pashmina Goat Genome: Sequencing, Assembly, and Annotation. Frontiers in Genetics, 2021, 12, 695178.	1.1	0
7	SNPs in Mammary Gland Epithelial Cells Unraveling Potential Difference in Milk Production Between Jersey and Kashmiri Cattle Using RNA Sequencing. Frontiers in Genetics, 2021, 12, 666015.	1.1	5
8	Abiotic stress induced miRNA-TF-gene regulatory network: A structural perspective. Genomics, 2020, 112, 412-422.	1.3	13
9	DNA methylation profiling identifies a high effect genetic variant for lipoprotein(a) levels. Epigenetics, 2020, 15, 949-958.	1.3	14
10	Comparative transcriptome analysis reveals the genetic basis of coat color variation in Pashmina goat. Scientific Reports, 2019, 9, 6361.	1.6	22
11	Comparative transcriptome analysis of mammary epithelial cells at different stages of lactation reveals wide differences in gene expression and pathways regulating milk synthesis between Jersey and Kashmiri cattle. PLoS ONE, 2019, 14, e0211773.	1.1	39
12	Draft genome sequence of Dichelobacter nodosus JKS-07 serogroup E from India. Journal of Global Antimicrobial Resistance, 2019, 16, 199-201.	0.9	0
13	Structural, functional and molecular dynamics analysis of cathepsin B gene SNPs associated with tropical calcific pancreatitis, a rare disease of tropics. PeerJ, 2019, 7, e7425.	0.9	7
14	mutTCPdb: a comprehensive database for genomic variants of a tropical country neglected diseaseâ€"tropical calcific pancreatitis. Database: the Journal of Biological Databases and Curation, 2018, 2018, .	1.4	6
15	TM-Aligner: Multiple sequence alignment tool for transmembrane proteins with reduced time and improved accuracy. Scientific Reports, 2017, 7, 12543.	1.6	23