Kolsoum Inanloorahatloo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4941183/publications.pdf

Version: 2024-02-01

1307594 1199594 16 324 12 7 citations g-index h-index papers 16 16 16 844 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	iPSC-derived cardiomyocytes reveal abnormal TGF- \hat{l}^2 signalling in left ventricular non-compaction cardiomyopathy. Nature Cell Biology, 2016, 18, 1031-1042.	10.3	148
2	Sex-based differences in myocardial gene expression in recently deceased organ donors with no prior cardiovascular disease. PLoS ONE, 2017, 12, e0183874.	2.5	43
3	A Rapid, High-Quality, Cost-Effective, Comprehensive and Expandable Targeted Next-Generation Sequencing Assay for Inherited Heart Diseases. Circulation Research, 2015, 117, 603-611.	4.5	34
4	Study on SARS-CoV-2 strains in Iran reveals potential contribution of co-infection with and recombination between different strains to the emergence of new strains. Virology, 2021, 562, 63-73.	2.4	26
5	Mutation in ST6GALNAC5 identified in family with coronary artery disease. Scientific Reports, 2014, 4, 3595.	3.3	23
6	Mutation in CYP27A1 identified in family with coronary artery disease. European Journal of Medical Genetics, 2013, 56, 655-660.	1.3	15
7	Evolution of SARSâ€CoVâ€2 genome from December 2019 to late March 2020: EmergedÂhaplotypes and informative TagÂnucleotide variations. Journal of Medical Virology, 2021, 93, 2010-2020.	5.0	10
8	Polymorphisms of cystathionine beta-synthase gene are associated with susceptibility to sepsis. European Journal of Human Genetics, 2016, 24, 1041-1048.	2.8	8
9	Expression of Dmrt Family Genes During Gonadal Differentiation in Two Species of Artemia (Branchiopoda, Anostraca) from Urmia Lake (Iran). Crustaceana, 2010, 83, 1153-1165.	0.3	6
10	Whole-Transcriptome Analysis Reveals Dysregulation of Actin-Cytoskeleton Pathway in Intellectual Disability Patients. Neuroscience, 2019, 404, 423-444.	2.3	4
11	Worldâ€wide tracking of major SARSâ€CoVâ€2 genome haplotypes in sequences of June 1 to November 15, 2020 and discovery of rapid expansion of a new haplotype. Journal of Medical Virology, 2021, 93, 3251-3256.	5.0	4
12	Deep geno- and phenotyping in two consanguineous families with CMT2 reveals HADHA as an unusual disease-causing gene and an intronic variant in GDAP1 as an unusual mutation. Journal of Neurology, 2021, 268, 640-650.	3.6	2
13	Intellectual disability associated with craniofacial dysmorphism due to POLR3B mutation and defect in spliceosomal machinery. BMC Medical Genomics, 2022, 15, 89.	1.5	1
14	Comparison of the protein profile of the reproductive system between bisexual and parthenogenetic species of Artemia (Branchiopoda, Anostraca). Crustaceana, 2009, 82, 1237-1248.	0.3	0
15	Abstract 248: Aberrant TGF \hat{I}^2 Signaling as an Etiology of Left Ventricular Non-compaction Cardiomyopathy. Circulation Research, 2015, 117, .	4.5	О
16	Intermittent white noise exposure is associated with rat cochleae damage and changes in the gene expression. Egyptian Journal of Medical Human Genetics, 2022, 23, .	1.0	0