

Bader Almuzzaini

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

414
citations

933447

10
h-index

794594

19
g-index

22
all docs

22
docs citations

22
times ranked

436
citing authors

#	ARTICLE	IF	CITATIONS
1	Blockade of p38 MAPK overcomes AML stem cell line KG1a resistance to 5-Fluorouridine and the impact on miRNA profiling. <i>PLoS ONE</i> , 2022, 17, e0267855.	2.5	7
2	Identification of the TTC26 Splice Variant in a Novel Complex Ciliopathy Syndrome with Biliary, Renal, Neurological, and Skeletal Manifestations. <i>Molecular Syndromology</i> , 2021, 12, 133-140.	0.8	5
3	Pancytopenia, Recurrent Infection, Poor Wound Healing, Heterotopia of the Brain Probably Associated with A Candidate Novel de Novo CDC42 Gene Defect: Expanding the Molecular and Phenotypic Spectrum. <i>Genes</i> , 2021, 12, 294.	2.4	9
4	Identification of Novel Mutations in Colorectal Cancer Patients Using AmpliSeq Comprehensive Cancer Panel. <i>Journal of Personalized Medicine</i> , 2021, 11, 535.	2.5	3
5	Biallelic variant in DACH1, encoding Dachshund Homolog 1, defines a novel candidate locus for recessive postaxial polydactyly type A. <i>Genomics</i> , 2021, 113, 2495-2502.	2.9	16
6	Herbal melanin induces interleukin-1 β secretion and production by human THP-1 monocytes via Toll-like receptor-2 and p38 MAPK activation. <i>Experimental and Therapeutic Medicine</i> , 2021, 22, 1081.	1.8	6
7	Interferon-induced transmembrane protein-3 genetic variant rs12252 is associated with COVID-19 mortality. <i>Genomics</i> , 2021, 113, 1733-1741.	2.9	39
8	Identification of CSF3R Mutations in B-Lineage Acute Lymphoblastic Leukemia Using Comprehensive Cancer Panel and Next-Generation Sequencing. <i>Genes</i> , 2021, 12, 1326.	2.4	3
9	A classification system for split-hand/ foot malformation (SHFM): A proposal based on 3 pedigrees with WNT10B mutations. <i>European Journal of Medical Genetics</i> , 2020, 63, 103738.	1.3	6
10	Blood pressure-lowering activity of statins: a systematic literature review and meta-analysis of placebo-randomized controlled trials. <i>European Journal of Clinical Pharmacology</i> , 2020, 76, 1745-1754.	1.9	3
11	The effect of the VKORC1 promoter variant on warfarin responsiveness in the Saudi Warfarin Pharmacogenetic (SWAP) cohort. <i>Scientific Reports</i> , 2020, 10, 11613.	3.3	9
12	Biallelic variants in four genes underlying recessive osteogenesis imperfecta. <i>European Journal of Medical Genetics</i> , 2020, 63, 103954.	1.3	26
13	Mutated <i>RAP1GDS1</i> causes a new syndrome of dysmorphic feature, intellectual disability & speech delay. <i>Annals of Clinical and Translational Neurology</i> , 2020, 7, 956-964.	3.7	21
14	Population pharmacokinetics of busulfan in Saudi pediatric patients undergoing hematopoietic stem cell transplantation. <i>International Journal of Clinical Pharmacy</i> , 2020, 42, 703-712.	2.1	3
15	A novel interstitial deletion of chromosome 2q21.1-q23.3: Case report and literature review. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1135.	1.2	6
16	Epidemiology of cancer in Saudi Arabia thru 2010-2019: a systematic review with constrained meta-analysis; Running title: epidemiology of cancer in Saudi Arabia thru 2010-2019. <i>AIMS Public Health</i> , 2020, 7, 679-696.	2.6	65
17	Targeted <i>SLC19A3</i> gene sequencing of 3000 Saudi newborn: a pilot study toward newborn screening. <i>Annals of Clinical and Translational Neurology</i> , 2019, 6, 2097-2103.	3.7	38
18	Actin-dependent global chromatin organization and gene expression programs control cellular identity. <i>FASEB Journal</i> , 2018, 32, 1296-1314.	0.5	50

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19	Nuclear Wiskottâ€Aldrich syndrome protein co-regulates T cell factor 1-mediated transcription in T cells. <i>Genome Medicine</i> , 2017, 9, 91.	8.2	16
20	In Î²â€actin knockouts, epigenetic reprogramming and rDNA transcription inactivation lead to growth and proliferation defects. <i>FASEB Journal</i> , 2016, 30, 2860-2873.	0.5	34
21	Nuclear myosin 1 contributes to a chromatin landscape compatible with RNA polymerase II transcription activation. <i>BMC Biology</i> , 2015, 13, 35.	3.8	48