Nuha Alrayes

List of Publications by Year in descending order

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Version: 2024-02-01

10	154	1478505	1372567
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#	Article	IF	CITATIONS
1	Exome analysis identified a novel missense mutation in the CLPP gene in a consanguineous Saudi family expanding the clinical spectrum of Perrault Syndrome type-3. Journal of the Neurological Sciences, 2015, 353, 149-154.	0.6	37
2	A missense mutation in TRAPPC6A leads to build-up of the protein, in patients with a neurodevelopmental syndrome and dysmorphic features. Scientific Reports, 2018, 8, 2053.	3.3	30
3	The alkylglycerol monooxygenase (AGMO) gene previously involved in autism also causes a novel syndromic form of primary microcephaly in a consanguineous Saudi family. Journal of the Neurological Sciences, 2016, 363, 240-244.	0.6	18
4	Truncating mutation in intracellular phospholipase A1 gene (DDHD2) in hereditary spastic paraplegia with intellectual disability (SPG54). BMC Research Notes, 2015, 8, 271.	1.4	17
5	Myocardial infarction biomarker discovery with integrated gene expression, pathways and biological networks analysis. Genomics, 2020, 112, 5072-5085.	2.9	17
6	Molecular insights into the coding region mutations of lowâ€density lipoprotein receptor adaptor protein 1 (LDLRAP1) linked to familial hypercholesterolemia. Journal of Gene Medicine, 2020, 22, e3176.	2.8	12
7	Whole exome sequencing of a Saudi family and systems biology analysis identifies CPED1 as a putative causative gene to Celiac Disease. Saudi Journal of Biological Sciences, 2020, 27, 1494-1502.	3.8	8
8	Exome Analysis Identifies a Novel Compound Heterozygous Alteration in TGM1 Gene Leading to Lamellar Ichthyosis in a Child From Saudi Arabia: Case Presentation. Frontiers in Pediatrics, 2019, 7, 44.	1.9	7
9	Novel missense alteration in <i>LRP4</i> gene underlies Cenani–Lenz syndactyly syndrome in a consanguineous family. Journal of Gene Medicine, 2020, 22, e3143.	2.8	5
10	Exome Sequencing Identifies the Extremely Rare ITGAV and FN1 Variants in Early Onset Inflammatory Bowel Disease Patients. Frontiers in Pediatrics, 2022, 10, .	1.9	3