Betoul Baz

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/7245652/publications.pdf

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19	276	933447 10	1058476 14
papers	citations	h-index	g-index

20	20	20	644
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	A genome-wide association study reveals susceptibility loci for myocardial infarction/coronary artery disease in Saudi Arabs. Atherosclerosis, 2016, 245, 62-70.	0.8	61
2	A study of the role of GATA2 gene polymorphism in coronary artery disease risk traits. Gene, 2014, 544, 152-158.	2.2	33
3	Integrated Analysis of Whole Exome Sequencing and Copy Number Evaluation in Parkinsonâ€™s Disease. Scientific Reports, 2019, 9, 3344.	3.3	31
4	Accelerated Endothelial to Mesenchymal Transition Increased Fibrosis via Deleting Notch Signaling in Wound Vasculature. Journal of Investigative Dermatology, 2018, 138, 1166-1175.	0.7	29
5	Novel homozygous mutation in DSP causing skin fragility-woolly hair syndrome: report of a large family and review of the desmoplakin-related phenotypes. Clinical Genetics, 2011, 80, 50-58.	2.0	27
6	A novel splice site mutation in ERLIN2 causes hereditary spastic paraplegia in a Saudi family. European Journal of Medical Genetics, 2013, 56, 43-45.	1.3	21
7	A New Susceptibility Locus for Myocardial Infarction, Hypertension, Type 2 Diabetes Mellitus, and Dyslipidemia on Chromosome 12q24. Disease Markers, 2014, 2014, 1-10.	1.3	17
8	Whole exome sequencing in ADHD trios from single and multi-incident families implicates new candidate genes and highlights polygenic transmission. European Journal of Human Genetics, 2020, 28, 1098-1110.	2.8	13
9	Validation of Ion TorrentTM Inherited Disease Panel with the PGMTM Sequencing Platform for Rapid and Comprehensive Mutation Detection. Genes, 2018, 9, 267.	2.4	12
10	Autozygome maps dispensable DNA and reveals potential selective bias against nullizygosity. Genetics in Medicine, 2012, 14, 515-519.	2.4	10
11	Molecular yield of targeted sequencing for Glanzmann thrombasthenia patients. Npj Genomic Medicine, 2019, 4, 4.	3.8	9
12	The Affymetrix DMET Plus Platform Reveals Unique Distribution of ADME-Related Variants in Ethnic Arabs. Disease Markers, 2015, 2015, 1-8.	1.3	8
13	Genetic variation in the mitogen-activated protein kinase/extracellular signal-regulated kinase pathway affects contact hypersensitivity responses. Journal of Allergy and Clinical Immunology, 2018, 142, 981-984.e7.	2.9	2
14	Molecular classification of blood and bleeding disorder genes. Npj Genomic Medicine, 2021, 6, 62.	3.8	2
15	574 Genetic associations of skin wound healing in the collaborative cross identifies aldose reductase family members as key regulators of healing speed. Journal of Investigative Dermatology, 2016, 136, S258.	0.7	1
16	Data on common variants associated with coronary artery disease/myocardial infarction in ethnic Arabs. Data in Brief, 2016, 7, 172-176.	1.0	0
17	183 Genome wide association identifies MAPKinase pathway regulators as key genetic determinants of allergic contact dermatitis. Journal of Investigative Dermatology, 2017, 137, S224.	0.7	0
18	662 Deletion of Notch signalling in the vasculature accelerates Endothelial to Mesenchymal Transition in skin wound healing. Journal of Investigative Dermatology, 2017, 137, S306.	0.7	0

#	Article	IF	Citations
19	A Murine Kitl Allele Regulates Skin Mast Cell Density across 58 Collaborative Mouse Cross Strains. Journal of Investigative Dermatology, 2022, 142, 2275-2280.e4.	0.7	0