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 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
2	Molecular classification of blood and bleeding disorder genes. Npj Genomic Medicine, 2021, 6, 62.	3.8	2
3	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
4	Integrated Analysis of Whole Exome Sequencing and Copy Number Evaluation in Parkinsonâ€™s Disease. Scientific Reports, 2019, 9, 3344.	3.3	31
5	Molecular yield of targeted sequencing for Glanzmann thrombasthenia patients. Npj Genomic Medicine, 2019, 4, 4.	3.8	9
6	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
7	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
8	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
9	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
10	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
11	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
12	Data on common variants associated with coronary artery disease/myocardial infarction in ethnic Arabs. Data in Brief, 2016, 7, 172-176.	1.0	0
13	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
14	The Affymetrix DMET Plus Platform Reveals Unique Distribution of ADME-Related Variants in Ethnic Arabs. Disease Markers, 2015, 2015, 1-8.	1.3	8
15	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
16	A study of the role of GATA2 gene polymorphism in coronary artery disease risk traits. Gene, 2014, 544, 152-158.	2.2	33
17	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
18	Autozygome maps dispensable DNA and reveals potential selective bias against nullizygosity. Genetics in Medicine, 2012, 14, 515-519.	2.4	10

#	Article	IF	Citations
19	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