

Zsolt Bã;nfai

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6455813/publications.pdf>

Version: 2024-02-01

11
papers

145
citations

1478505

6
h-index

1199594

12
g-index

15
all docs

15
docs citations

15
times ranked

326
citing authors

#	ARTICLE	IF	CITATIONS
1	Interleukins and interleukin receptors in rheumatoid arthritis: Research, diagnostics and clinical implications. <i>World Journal of Orthopedics</i> , 2014, 5, 516.	1.8	60
2	Refining the South Asian Origin of the Romani people. <i>BMC Genetics</i> , 2017, 18, 82.	2.7	26
3	Marked Differences of Haplotype Tagging SNP Distribution, Linkage, and Haplotype Profile of APOA5 Gene in Roma Population Samples. <i>Pathology and Oncology Research</i> , 2017, 23, 853-861.	1.9	10
4	Revealing the Genetic Impact of the Ottoman Occupation on Ethnic Groups of East-Central Europe and on the Roma Population of the Area. <i>Frontiers in Genetics</i> , 2019, 10, 558.	2.3	9
5	Genome-wide autosomal, mtDNA, and Y chromosome analysis of King Bela III of the Hungarian Arpad dynasty. <i>Scientific Reports</i> , 2021, 11, 19210.	3.3	9
6	Novel phenotypic variant in the MYH7 spectrum due to a stop-loss mutation in the C-terminal region: a case report. <i>BMC Medical Genetics</i> , 2017, 18, 105.	2.1	8
7	Genetic polymorphisms in promoter and intronic regions of CYP1A2 gene in Roma and Hungarian population samples. <i>Environmental Toxicology and Pharmacology</i> , 2014, 38, 814-820.	4.0	7
8	Small supernumerary marker chromosome 15 and a ring chromosome 15 associated with a 15q26.3 deletion excluding the IGF1R gene. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 443-449.	1.2	7
9	Revealing the impact of the Caucasus region on the genetic legacy of Romani people from genome-wide data. <i>PLoS ONE</i> , 2018, 13, e0202890.	2.5	5
10	Investigating the genetic characteristics of the Csangos, a traditionally Hungarian speaking ethnic group residing in Romania. <i>Journal of Human Genetics</i> , 2020, 65, 1093-1103.	2.3	2
11	Genome-Wide Marker Data-Based Comparative Population Analysis of Szeklers From Korond, Transylvania, and From Transylvania Living Non-Szekler Hungarians. <i>Frontiers in Genetics</i> , 2022, 13, 841769.	2.3	1