

Bã©la Melegh

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3529397/publications.pdf>

Version: 2024-02-01

64
papers

3,476
citations

430874

18
h-index

175258

52
g-index

72
all docs

72
docs citations

72
times ranked

6335
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical and genetic spectrum of a large cohort of patients with $\hat{\Gamma}$ -sarcoglycan muscular dystrophy. <i>Brain</i> , 2022, 145, 596-606.	7.6	11
2	The need for recognition of core professional groups in genetics healthcare services in Europe. <i>European Journal of Human Genetics</i> , 2022, 30, 639-640.	2.8	2
3	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
4	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.	27.8	929
5	TUBB4B gene mutation in Leber phenotype of congenital amaurosis syndrome associated with early-onset deafness. <i>European Journal of Medical Genetics</i> , 2022, 65, 104471.	1.3	3
6	Äœber die Notwendigkeit der Anerkennung von sog. Kernberufsgruppen innerhalb der genetischen Gesundheitsversorgung in Europa. <i>Medizinische Genetik</i> , 2022, 34, 81-83.	0.2	1
7	Erratum zu: Äœber die Notwendigkeit der Anerkennung von sog. Kernberufsgruppen innerhalb der genetischen Gesundheitsversorgung in Europa. <i>Medizinische Genetik</i> , 2022, 34, 189-191.	0.2	0
8	Xp11.2 Duplication in Females: Unique Features of a Rare Copy Number Variation. <i>Frontiers in Genetics</i> , 2021, 12, 635458.	2.3	2
9	Genotype-Phenotype Associations in Patients With Type-1, Type-2, and Atypical NF1 Microdeletions. <i>Frontiers in Genetics</i> , 2021, 12, 673025.	2.3	11
10	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
11	Whole Exome Sequencing in a Series of Patients with a Clinical Diagnosis of Tuberous Sclerosis Not Confirmed by Targeted TSC1/TSC2 Sequencing. <i>Genes</i> , 2021, 12, 1401.	2.4	4
12	Intrafamilial variability of limb-girdle muscular dystrophy, LGMD1D type. <i>European Journal of Medical Genetics</i> , 2020, 63, 103655.	1.3	10
13	Mutation spectrum of the SCN1A gene in a Hungarian population with epilepsy. <i>Seizure: the Journal of the British Epilepsy Association</i> , 2020, 74, 8-13.	2.0	8
14	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
15	New genotype-phenotype correlations in a large European cohort of patients with sarcoglycanopathy. <i>Brain</i> , 2020, 143, 2696-2708.	7.6	45
16	Conversion of individuals at risk for spinocerebellar ataxia types 1, 2, 3, and 6 to manifest ataxia (RISCA): a longitudinal cohort study. <i>Lancet Neurology</i> , The, 2020, 19, 738-747.	10.2	41
17	Mass Spectrometric Analysis of L-carnitine and its Esters: Potential Biomarkers of Disturbances in Carnitine Homeostasis. <i>Current Molecular Medicine</i> , 2020, 20, 336-354.	1.3	15
18	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

#	ARTICLE	IF	CITATIONS
19	Possible Phenotypic Consequences of Structural Differences in Idic(15) in a Small Cohort of Patients. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4935.	4.1	1
20	Prediction of Survival With Long-Term Disease Progression in Most Common Spinocerebellar Ataxia. <i>Movement Disorders</i> , 2019, 34, 1220-1227.	3.9	14
21	A novel pathogenic variant in TNPO3 in a Hungarian family with limb-girdle muscular dystrophy 1F. <i>European Journal of Medical Genetics</i> , 2019, 62, 103662.	1.3	15
22	Hearing impairment locus heterogeneity and identification of PLS1 as a new autosomal dominant gene in Hungarian Roma. <i>European Journal of Human Genetics</i> , 2019, 27, 869-878.	2.8	10
23	DNA profiling of Hungarian King BáĀla III and other skeletal remains originating from the Royal Basilica of SzĀkesfehĀrvĀr. <i>Archaeological and Anthropological Sciences</i> , 2019, 11, 1345-1357.	1.8	24
24	Age-Related Hearing Impairment Associated NAT2, GRM7, GRHL2 Susceptibility Gene Polymorphisms and Haplotypes in Roma and Hungarian Populations. <i>Pathology and Oncology Research</i> , 2019, 25, 1349-1355.	1.9	5
25	Role of carnitine and its derivatives in the development and management of type 2 diabetes. <i>Nutrition and Diabetes</i> , 2018, 8, 8.	3.2	121
26	Survival in patients with spinocerebellar ataxia types 1, 2, 3, and 6 (EUROSCA): a longitudinal cohort study. <i>Lancet Neurology</i> , The, 2018, 17, 327-334.	10.2	69
27	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
28	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
29	Long-term evolution of patient-reported outcome measures in spinocerebellar ataxias. <i>Journal of Neurology</i> , 2018, 265, 2040-2051.	3.6	34
30	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
31	CYP2C9 and VKORC1 in therapeutic dosing and safety of acenocoumarol treatment: implication for clinical practice in Hungary. <i>Environmental Toxicology and Pharmacology</i> , 2017, 56, 282-289.	4.0	4
32	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
33	On the distribution of intranuclear and cytoplasmic aggregates in the brainstem of patients with spinocerebellar ataxia type 2 and 3. <i>Brain Pathology</i> , 2017, 27, 345-355.	4.1	36
34	Refining the South Asian Origin of the Romani people. <i>BMC Genetics</i> , 2017, 18, 82.	2.7	26
35	Initiating an undiagnosed diseases program in the Western Australian public health system. <i>Orphanet Journal of Rare Diseases</i> , 2017, 12, 83.	2.7	24
36	Kleefstra syndrome in Hungarian patients: additional symptoms besides the classic phenotype. <i>Molecular Cytogenetics</i> , 2016, 9, 22.	0.9	14

#	ARTICLE	IF	CITATIONS
37	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. <i>JAMA Psychiatry</i> , 2016, 73, 497.	11.0	51
38	Cytochrome P450 Drug Metabolizing Enzymes in Roma Population Samples: Systematic Review of the Literature. <i>Current Medicinal Chemistry</i> , 2016, 23, 3632-3652.	2.4	7
39	Partial tetrasomy of the proximal long arm of chromosome 15 in two patients: the significance of the gene dosage in terms of phenotype. <i>Molecular Cytogenetics</i> , 2015, 8, 41.	0.9	3
40	Marked differences in frequencies of statin therapy relevant SLCO1B1 variants and haplotypes between Roma and Hungarian populations. <i>BMC Genetics</i> , 2015, 16, 108.	2.7	10
41	Genome Sequences of Three Turkey Orthoreovirus Strains Isolated in Hungary. <i>Genome Announcements</i> , 2015, 3, .	0.8	3
42	MARVELD2 (DFNB49) Mutations in the Hearing Impaired Central European Roma Population - Prevalence, Clinical Impact and the Common Origin. <i>PLoS ONE</i> , 2015, 10, e0124232.	2.5	17
43	Identification of <i>SLC22A5</i> Gene Mutation in a Family with Carnitine Uptake Defect. <i>Case Reports in Genetics</i> , 2015, 2015, 1-5.	0.2	14
44	Towards a European consensus for reporting incidental findings during clinical NGS testing. <i>European Journal of Human Genetics</i> , 2015, 23, 1601-1606.	2.8	85
45	Extreme differences in SLCO1B3 functional polymorphisms in Roma and Hungarian populations. <i>Environmental Toxicology and Pharmacology</i> , 2015, 39, 1246-1251.	4.0	4
46	Undiagnosed Diseases Network International (UDNI): White paper for global actions to meet patient needs. <i>Molecular Genetics and Metabolism</i> , 2015, 116, 223-225.	1.1	69
47	Interethnic variability of CYP4F2 (V433M) in admixed population of Roma and Hungarians. <i>Environmental Toxicology and Pharmacology</i> , 2015, 40, 280-283.	4.0	2
48	Interethnic differences of cytochrome P450 gene polymorphisms may influence outcome of taxane therapy in Roma and Hungarian populations. <i>Drug Metabolism and Pharmacokinetics</i> , 2015, 30, 453-456.	2.2	3
49	Increased prevalence of functional minor allele variants of drug metabolizing CYP2B6 and CYP2D6 genes in Roma population samples. <i>Pharmacological Reports</i> , 2015, 67, 460-464.	3.3	14
50	Phenotypic variability in a Hungarian patient with the 4q21 microdeletion syndrome. <i>Molecular Cytogenetics</i> , 2015, 8, 16.	0.9	11
51	Long-term disease progression in spinocerebellar ataxia types 1, 2, 3, and 6: a longitudinal cohort study. <i>Lancet Neurology</i> , The, 2015, 14, 1101-1108.	10.2	213
52	Interleukins and interleukin receptors in rheumatoid arthritis: Research, diagnostics and clinical implications. <i>World Journal of Orthopedics</i> , 2014, 5, 516.	1.8	60
53	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
54	Deletion of 4q28.3-31.23 in the background of multiple malformations with pulmonary hypertension. <i>Molecular Cytogenetics</i> , 2014, 7, 36.	0.9	15

#	ARTICLE	IF	CITATIONS
55	Ancient human genomes suggest three ancestral populations for present-day Europeans. <i>Nature</i> , 2014, 513, 409-413.	27.8	1,179
56	Marked differences of haplotype tagging SNP distribution, linkage, and haplotype profile of IL23 receptor gene in Roma and Hungarian population samples. <i>Cytokine</i> , 2014, 65, 148-152.	3.2	4
57	Exome sequencing identifies Laing distal myopathy MYH7 mutation in a Roma family previously diagnosed with distal neuronopathy. <i>Neuromuscular Disorders</i> , 2014, 24, 156-161.	0.6	17
58	Interleukin and interleukin receptor gene polymorphisms in inflammatory bowel diseases susceptibility. <i>World Journal of Gastroenterology</i> , 2014, 20, 3208.	3.3	23
59	Reconstructing Roma History from Genome-Wide Data. <i>PLoS ONE</i> , 2013, 8, e58633.	2.5	61
60	Non-syndromic Hearing Impairment in a Hungarian Family with the m.7510T>C Mutation of Mitochondrial tRNASer(UCN) and Review of Published Cases. <i>JIMD Reports</i> , 2012, 9, 105-111.	1.5	3
61	Analysis of mtDNA A3243G mutation frequency in Hungary. <i>Open Medicine (Poland)</i> , 2010, 5, 322-328.	1.3	3
62	Tel Hashomer camptodactyly syndrome: 12-year follow-up of a Hungarian patient and review. <i>American Journal of Medical Genetics, Part A</i> , 2005, 135A, 320-323.	1.2	4
63	Molecular epidemiology of human P[8],G9 rotaviruses in Hungary between 1998 and 2001. <i>Journal of Medical Microbiology</i> , 2004, 53, 791-801.	1.8	24
64	Phenotypic manifestations of the OCTN2 V295X mutation: Sudden infant death and carnitine-responsive cardiomyopathy in Roma families. <i>American Journal of Medical Genetics Part A</i> , 2004, 131A, 121-126.	2.4	34