

AyÅe NazlÄ± BaÅak

List of Publications by Year in descending order

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46
papers

1,938
citations

471509

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289244

40
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48
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docs citations

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times ranked

3704
citing authors

#	ARTICLE	IF	CITATIONS
1	De Novo and Dominantly Inherited <i>SPTAN1</i> Mutations Cause Spastic Paraplegia and Cerebellar Ataxia. <i>Movement Disorders</i> , 2022, 37, 1175-1186.	3.9	9
2	Impact of the Amyotrophic Lateral Sclerosis Disease on the Biomechanical Properties and Oxidative Stress Metabolism of the Lung Tissue Correlated With the Human Mutant SOD1G93A Protein Accumulation. <i>Frontiers in Bioengineering and Biotechnology</i> , 2022, 10, 810243.	4.1	3
3	Comprehensive Research on Past and Future Therapeutic Strategies Devoted to Treatment of Amyotrophic Lateral Sclerosis. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2400.	4.1	32
4	Genome-wide study of DNA methylation shows alterations in metabolic, inflammatory, and cholesterol pathways in ALS. <i>Science Translational Medicine</i> , 2022, 14, eabj0264.	12.4	38
5	Vestibulo-ocular reflex impairment in SPG7 hereditary spastic paraplegia. <i>Clinical Neurophysiology</i> , 2021, 132, 77-79.	1.5	1
6	A novel PNPLA6 mutation in a Turkish family with intractable Holmes tremor and spastic ataxia. <i>Neurological Sciences</i> , 2021, 42, 1535-1539.	1.9	13
7	Two cases of early-onset autosomal recessive spastic ataxia of Charlevoix-Saguenay diagnosed in adulthood. <i>Clinical Neurology and Neurosurgery</i> , 2021, 201, 106423.	1.4	1
8	A rare case of juvenile amyotrophic lateral sclerosis. <i>Turkish Journal of Pediatrics</i> , 2021, 63, 495.	0.6	2
9	The Effect of <i>SMN</i> Gene Dosage on ALS Risk and Disease Severity. <i>Annals of Neurology</i> , 2021, 89, 686-697.	5.3	10
10	A combined clinical and computational approach to understand the SOD1A4T-mediated pathogenesis of rapidly progressive familial amyotrophic lateral sclerosis. <i>Acta Neurologica Belgica</i> , 2021, , 1.	1.1	1
11	The Complex Genetic Landscape of Hereditary Ataxias in Turkey and Implications in Clinical Practice. <i>Movement Disorders</i> , 2021, 36, 1676-1688.	3.9	9
12	Bi-allelic variants in HOPS complex subunit VPS41 cause cerebellar ataxia and abnormal membrane trafficking. <i>Brain</i> , 2021, 144, 769-780.	7.6	33
13	A novel pathogenic variant in the 3' end of the <i>AGTPBP1</i> gene gives rise to neurodegeneration without cerebellar atrophy: an expansion of the disease phenotype?. <i>Neurogenetics</i> , 2021, 22, 127-132.	1.4	7
14	Varied phenotypic spectrum presenting of paroxysmal exercise-induced dyskinesia: a Turkish family with <i>SLC2A1</i> mutation. <i>Neurological Sciences</i> , 2021, 42, 4751-4754.	1.9	1
15	The genetic structure of the Turkish population reveals high levels of variation and admixture. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	42
16	Association of Variants in the <i>SPTLC1</i> Gene With Juvenile Amyotrophic Lateral Sclerosis. <i>JAMA Neurology</i> , 2021, 78, 1236.	9.0	46
17	Early-Onset Parkinson's Disease: A Novel Deletion Comprising the <i>DJ</i> and <i>TNFRSF9</i> Genes. <i>Movement Disorders</i> , 2021, 36, 2973-2976.	3.9	5
18	The first biallelic missense mutation in the <i>FXN</i> gene in a consanguineous Turkish family with Charcot-Marie-Tooth-like phenotype. <i>Neurogenetics</i> , 2020, 21, 73-78.	1.4	11

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19	Cerebellar cognitive-affective syndrome preceding ataxia associated with complex extrapyramidal features in a Turkish SCA48 family. <i>Neurogenetics</i> , 2020, 21, 51-58.	1.4	22
20	Revisiting the complex architecture of ALS in Turkey: Expanding genotypes, shared phenotypes, molecular networks, and a public variant database. <i>Human Mutation</i> , 2020, 41, e7-e45.	2.5	10
21	Reply to letter to the editor by De Michele et al. <i>Neurogenetics</i> , 2020, 21, 147-147.	1.4	0
22	Ataxia telangiectasia like disorder: Another dopa-responsive disorder look-alike?. <i>Parkinsonism and Related Disorders</i> , 2020, 74, 22-24.	2.2	1
23	A novel homozygous FBXO38 variant causes an early-onset distal hereditary motor neuronopathy type IID. <i>Journal of Human Genetics</i> , 2019, 64, 1141-1144.	2.3	9
24	Mutations in the Glycosyltransferase Domain of GLT8D1 Are Associated with Familial Amyotrophic Lateral Sclerosis. <i>Cell Reports</i> , 2019, 26, 2298-2306.e5.	6.4	57
25	ERLIN1 mutations cause teenage-onset slowly progressive ALS in a large Turkish pedigree. <i>European Journal of Human Genetics</i> , 2018, 26, 745-748.	2.8	25
26	Homozygous <i>CAPN1</i> mutations causing a spastic-ataxia phenotype in 2 families. <i>Neurology: Genetics</i> , 2018, 4, e218.	1.9	19
27	Elevated Global DNA Methylation Is Not Exclusive to Amyotrophic Lateral Sclerosis and Is Also Observed in Spinocerebellar Ataxia Types 1 and 2. <i>Neurodegenerative Diseases</i> , 2018, 18, 38-48.	1.4	27
28	Genome-wide Analyses Identify KIF5A as a Novel ALS Gene. <i>Neuron</i> , 2018, 97, 1268-1283.e6.	8.1	517
29	Phenotypic and Genotypic Analysis of Hereditary Ataxia Patients in Sakarya City, Turkey. <i>Noropsikiyatri Arsivi</i> , 2017, 56, 106-109.	0.3	0
30	Impact of Genetic Defects on Coronary Atherosclerosis among Turkish Cypriots. <i>Heart Surgery Forum</i> , 2017, 20, 223.	0.5	4
31	Atypical Features in a Large Turkish Family Affected with Friedreich Ataxia. <i>Case Reports in Neurological Medicine</i> , 2016, 2016, 1-7.	0.4	4
32	The genetic basis of asymptomatic codon 8 frame-shift (<i>HBB</i> :c25_26delAA) β^0 thalassaemia homozygotes. <i>British Journal of Haematology</i> , 2016, 172, 958-965.	2.5	4
33	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016, 48, 1043-1048.	21.4	494
34	NEK1 variants confer susceptibility to amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016, 48, 1037-1042.	21.4	218
35	Successful treatment of Juvenile parkinsonism with bilateral subthalamic deep brain stimulation in a 14-year-old patient with parkin gene mutation. <i>Parkinsonism and Related Disorders</i> , 2016, 24, 137-138.	2.2	8
36	Braitâ€Fahnâ€Schwarz disease: Parkinsonâ€™s disease and amyotrophic lateral sclerosis complex. <i>Acta Neurologica Belgica</i> , 2016, 116, 401-403.	1.1	4

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37	A Turkish Family with a Familial ALS-positive UBQLN2-S340I Mutation. <i>Noropsikiyatri Arsivi</i> , 2016, 53, 283-285.	0.7	5
38	Both Ubiquitin Ligases FBXW8 and PARK2 Are Sequestered into Insolubility by ATXN2 PolyQ Expansions, but Only FBXW8 Expression Is Dysregulated. <i>PLoS ONE</i> , 2015, 10, e0121089.	2.5	18
39	Turkish families with juvenile motor neuron disease broaden the phenotypic spectrum of <i>SPG11</i> . <i>Neurology: Genetics</i> , 2015, 1, e25.	1.9	10
40	The distinct genetic pattern of ALS in Turkey and novel mutations. <i>Neurobiology of Aging</i> , 2015, 36, 1764.e9-1764.e18.	3.1	78
41	Coexistence of autosomal recessive spastic ataxia of Charlevoix Saguenay and spondyloepiphyseal dysplasia in a Turkish patient. <i>Journal of the Neurological Sciences</i> , 2015, 357, 290-291.	0.6	4
42	FBXO7â€R498X mutation: Phenotypic variability from chorea to early onset parkinsonism within a family. <i>Parkinsonism and Related Disorders</i> , 2014, 20, 1253-1256.	2.2	29
43	ATXN2 and Its Neighbouring Gene SH2B3 Are Associated with Increased ALS Risk in the Turkish Population. <i>PLoS ONE</i> , 2012, 7, e42956.	2.5	43
44	Î²-THALASSEMIA IN TURKEY: A REVIEW OF THE CLINICAL, EPIDEMIOLOGICAL, MOLECULAR, AND EVOLUTIONARY ASPECTS. <i>Hemoglobin</i> , 2001, 25, 227-239.	0.8	36
45	Rare Î²-thalassemia mutation IVS-II-848 (C-A) first reported in a Turkish cypriot family. , 1997, 54, 338-339.		3
46	PRENATAL DIAGNOSIS OF Î²-THALASSAEMIA AND SICKLE CELL ANAEMIA IN TURKEY. , 1996, 16, 252-258.		21