## Zohreh Fattahi

List of Publications by Year in descending order

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687363 501196 30 984 13 28 citations h-index g-index papers 32 32 32 2443 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Biallelic mutations in the death domain of PIDD1 impair caspase-2 activation and are associated with intellectual disability. Translational Psychiatry, 2021, 11, 1.	4.8	334
2	Genetics of intellectual disability in consanguineous families. Molecular Psychiatry, 2019, 24, 1027-1039.	7.9	131
3	Iranome: A catalog of genomic variations in the Iranian population. Human Mutation, 2019, 40, 1968-1984.	2.5	116
4	The Role of a Novel TRMT1 Gene Mutation and Rare GRM1 Gene Defect in Intellectual Disability in Two Azeri Families. PLoS ONE, 2015, 10, e0129631.	2.5	56
5	Screening for <i>MYO15A</i> gene mutations in autosomal recessive nonsyndromic, <i>GJB2</i> negative Iranian deaf population. American Journal of Medical Genetics, Part A, 2012, 158A, 1857-1864.	1.2	54
6	Effect of inbreeding on intellectual disability revisited by trio sequencing. Clinical Genetics, 2019, 95, 151-159.	2.0	49
7	Distinct genetic variation and heterogeneity of the Iranian population. PLoS Genetics, 2019, 15, e1008385.	3.5	34
8	Mutation profile of BBS genes in Iranian patients with Bardetâ€"Biedl syndrome: genetic characterization and report of nine novel mutations in five BBS genes. Journal of Human Genetics, 2014, 59, 368-375.	2.3	33
9	A defect in the CLIP1 gene (CLIP-170) can cause autosomal recessive intellectual disability. European Journal of Human Genetics, 2015, 23, 331-336.	2.8	22
10	Biallelic missense variants in ZBTB11 can cause intellectual disability in humans. Human Molecular Genetics, 2018, 27, 3177-3188.	2.9	19
11	SARSâ€CoVâ€2 outbreak in Iran: The dynamics of the epidemic and evidence on two independent introductions. Transboundary and Emerging Diseases, 2022, 69, 1375-1386.	3.0	19
12	Whole Genome Linkage Analysis Followed by Whole Exome Sequencing Identifies Nicastrin (NCSTN) as a Causative Gene in a Multiplex Family with <sup>13</sup> -Secretase Spectrum of Autoinflammatory Skin Phenotypes. Journal of Investigative Dermatology, 2016, 136, 1283-1286.	0.7	17
13	Missense variants in AIMP1 gene are implicated in autosomal recessive intellectual disability without neurodegeneration. European Journal of Human Genetics, 2016, 24, 392-399.	2.8	17
14	<i>GPR126</i> : A novel candidate gene implicated in autosomal recessive intellectual disability. American Journal of Medical Genetics, Part A, 2019, 179, 13-19.	1.2	12
15	SZT2 mutation in a boy with intellectual disability, seizures and autistic features. European Journal of Medical Genetics, 2019, 62, 103556.	1.3	12
16	Report of a patient with limb-girdle muscular dystrophy, ptosis and ophthalmoparesis caused by plectinopathy. Archives of Iranian Medicine, 2015, 18, 60-4.	0.6	8
17	Molecular Diagnosis of Hereditary Neuropathies by Whole Exome Sequencing and Expanding the Phenotype Spectrum. Archives of Iranian Medicine, 2020, 23, 426-433.	0.6	6
18	Genetic Investigation of an Iranian Supercentenarian by Whole Exome Sequencing. Archives of Iranian Medicine, 2015, 18, 688-97.	0.6	6

#	Article	IF	CITATIONS
19	Report of limb girdle muscular dystrophy type 2a in 6 Iranian patients, one with a novel deletion in CAPN3 gene. Neuromuscular Disorders, 2016, 26, 277-282.	0.6	5
20	Identification of diseaseâ€causing variants in the <i>EXOSC</i> gene family underlying autosomal recessive intellectual disability in Iranian families. Clinical Genetics, 2019, 95, 718-725.	2.0	5
21	ZBTB11 dysfunction: spectrum of brain abnormalities, biochemical signature and cellular consequences. Brain, 2022, 145, 2602-2616.	7.6	5
22	Finding mutation within non-coding region of GJB2 reveals its importance in genetic testing of Hearing Loss in Iranian population. International Journal of Pediatric Otorhinolaryngology, 2015, 79, 136-138.	1.0	4
23	<i>CNKSR1</i> gene defect can cause syndromic autosomal recessive intellectual disability. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 691-699.	1.7	4
24	Exome Sequencing and Linkage Analysis Identified Novel Candidate Genes in Recessive Intellectual Disability Associated with Ataxia. Archives of Iranian Medicine, 2015, 18, 670-82.	0.6	4
25	Novel variants in Iranian individuals suspected to have inherited red blood cell disorders, including bone marrow failure syndromes. Haematologica, 2020, 105, e1-e4.	3.5	3
26	<i>CEP104</i> and <i>CEP290</i> ; Genes with Ciliary Functions Cause Intellectual Disability in Multiple Families. Archives of Iranian Medicine, 2021, 24, 364-373.	0.6	3
27	Comprehensive <scp>genotypeâ€phenotype</scp> correlation in <scp>AP</scp> â€4 deficiency syndrome; Adding data from a large cohort of Iranian patients. Clinical Genetics, 2021, 99, 187-192.	2.0	2
28	Whole genome sequencing identifies a duplicated region encompassing Xq13.2q13.3 in a large Iranian family with intellectual disability. Molecular Genetics & Enomic Medicine, 2020, 8, e1418.	1.2	1
29	Brief Report of Variants Detected in Hereditary Hearing Loss Cases in Iran over a 3-Year Period. Iranian Journal of Public Health, 2019, 48, 1910-1915.	0.5	1
30	De novo Mutation in CACNA1S Gene in a 20-Year-Old Man Diagnosed with Metabolic Myopathy. Archives of Iranian Medicine, 2017, 20, 617-620.	0.6	1