Hossein Najmabadi

List of Publications by Year
in descending order

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265 papers	9,847 citations	43973 48 h-index	53109 85 g-index
276 all docs	276 docs citations	276 times ranked	14351 citing authors

#	Article	IF	Citations
1	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.	1.3	19
2	Anticipation Can Be More Common in Hereditary Spastic Paraplegia with SPAST Mutations Than It Appears. Canadian Journal of Neurological Sciences, 2022, 49, 651-661.	0.3	6
3	Genetic etiology of hearing loss in Iran. Human Genetics, 2022, 141, 623-631.	1.8	6
4	ZBTB11 dysfunction: spectrum of brain abnormalities, biochemical signature and cellular consequences. Brain, 2022, 145, 2602-2616.	3.7	5
5	Phenotype and genotype spectrum of variants in guanine nucleotide exchange factor genes in a broad cohort of Iranian patients. Molecular Genetics & Genomic Medicine, 2022, 10, e1894.	0.6	5
6	Intellectual disability associated with craniofacial dysmorphism due to POLR3B mutation and defect in spliceosomal machinery. BMC Medical Genomics, 2022, 15, 89.	0.7	1
7	Understanding the Molecular Basis of Fragile X Syndrome Using Differentiated Mesenchymal Stem Cells.. Iranian Journal of Child Neurology, 2022, 16, 85-95.	0.2	0
8	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.	1.0	2
9	The First Case of BENTA Disease (B Cell Expansion with NF-Î²B and T Cell Anergy) from Iran. Journal of Clinical Immunology, 2021, 41, 811-813.	2.0	2
10	Biallelic mutations in the death domain of PIDD1 impair caspase-2 activation and are associated with intellectual disability. Translational Psychiatry, 2021, 11, 1.	2.4	334
11	Subcellular relocalization and nuclear redistribution of the RNA methyltransferases TRMT1 and TRMT1L upon neuronal activation. RNA Biology, 2021, 18, 1905-1919.	1.5	9
12	POLRMT mutations impair mitochondrial transcription causing neurological disease. Nature Communications, 2021, 12, 1135.	5.8	21
13	Exome sequencing utility in defining the genetic landscape of hearing loss and novelâ€gene discovery in Iran. Clinical Genetics, 2021, 100, 59-78.	1.0	4
14	NGLY1 deficiency: Novel variants and literature review. European Journal of Medical Genetics, 2021, 64, 104146.	0.7	15
15	Clinical and Genetic Characteristics of Splicing Variant in CYP27A1 in an Iranian Family with Cerebrotendinous xanthomatosis. Iranian Biomedical Journal, 2021, 25, 132-139.	0.4	4
16	Identifying the causes of recurrent pregnancy loss in consanguineous couples using whole exome sequencing on the products of miscarriage with no chromosomal abnormalities. Scientific Reports, 2021, 11, 6952.	1.6	15
17	Discovery of a neuromuscular syndrome caused by biallelic variants in ASCC3. Human Genetics and Genomics Advances, 2021, 2, 100024.	1.0	1
18	CEP104 and CEP290; Genes with Ciliary Functions Cause Intellectual Disability in Multiple Families. Archives of Iranian Medicine, 2021, 24, 364-373.	0.2	3

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19	A Novel Î²0-Thalassemia Mutation, HBB: c.356_357delTT [Codon 118 (â€“TT)] in an Iraqi Kurd. Hemoglobin, 2021, 45, 212-214.	0.4	1
20	The recurrent missense mutation p.(Arg367Trp) in YARS1 causes a distinct neurodevelopmental phenotype. Journal of Molecular Medicine, 2021, 99, 1755-1768.	1.7	3
21	The PTRHD1 Mutation in Intellectual Disability. Archives of Iranian Medicine, 2021, 24, 747-751.	0.2	3
22	A novel variant of C12orf4 linked to autosomal recessive intellectual disability type 66 with phenotype expansion. Journal of Gene Medicine, 2021, , e3406.	1.4	0
23	Novel variants in Iranian individuals suspected to have inherited red blood cell disorders, including bone marrow failure syndromes. Haematologica, 2020, 105, e1-e4.	1.7	3
24	The second mutation of SYCE1 gene associated with autosomal recessive nonobstructive azoospermia. Journal of Assisted Reproduction and Genetics, 2020, 37, 451-458.	1.2	34
25	Heterozygosity for the Novel HBA2: c.*91_*92delTA Polyadenylation Site Variant on the Î±2-Globin Gene Expanding the Genetic Spectrum of Î±-Thalassemia in Iran. Hemoglobin, 2020, 44, 423-426.	0.4	2
26	YIF1B mutations cause a post-natal neurodevelopmental syndrome associated with Golgi and primary cilium alterations. Brain, 2020, 143, 2911-2928.	3.7	13
27	Whole genome sequencing identifies a duplicated region encompassing Xq13.2q13.3 in a large Iranian family with intellectual disability. Molecular Genetics & Genomic Medicine, 2020, 8, e1418.	0.6	1
28	FASN-Dependent Lipid Metabolism Links Neurogenic Stem/Progenitor Cell Activity to Learning and Memory Deficits. Cell Stem Cell, 2020, 27, 98-109.e11.	5.2	62
29	Adultâ€onset veryâ€longâ€chain acylâ€CoA dehydrogenase deficiency (VLCADD). European Journal of Neurology, 2020, 27, 2257-2266.	1.7	6
30	When transcripts matter: delineating between non-syndromic hearing loss DFNB32 and hearing impairment infertile male syndrome (HIIMS). Journal of Human Genetics, 2020, 65, 609-617.	1.1	2
31	A novel PTC mutation in the BTB domain of KLHL7 gene in two patients with Bohring-Opitz syndrome-like features. European Journal of Medical Genetics, 2020, 63, 103849.	0.7	3
32	CAPN1 and hereditary spastic paraplegia: a novel variant in an Iranian family and overview of the genotype-phenotype correlation. International Journal of Neuroscience, 2020, 131, 1-13.	0.8	7
33	Limbic System Associated Membrane Protein Mutation in an Iranian Family Diagnosed with MÃ©niÃ¨reâ€™s Disease. Archives of Iranian Medicine, 2020, 23, 319-325.	0.2	6
34	Novel Mutation in LARP7 in Two Iranian Consanguineous Families with Syndromic Intellectual Disability and Facial Dysmorphism. Archives of Iranian Medicine, 2020, 23, 842-847.	0.2	2
35	Molecular Diagnosis of Hereditary Neuropathies by Whole Exome Sequencing and Expanding the Phenotype Spectrum. Archives of Iranian Medicine, 2020, 23, 426-433.	0.2	6
36	GPR126: A novel candidate gene implicated in autosomal recessive intellectual disability. American Journal of Medical Genetics, Part A, 2019, 179, 13-19.	0.7	12

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37	G130V de novo mutation in an Iranian pedigree with nonsyndromic hearing loss without palmoplantar keratoderma. International Journal of Pediatric Otorhinolaryngology, 2019, 126, 109607.	0.4	1
38	Iranome: A catalog of genomic variations in the Iranian population. Human Mutation, 2019, 40, 1968-1984.	1.1	116
39	Bi-allelic Pathogenic Variants in TUBGCP2 Cause Microcephaly and Lissencephaly Spectrum Disorders. American Journal of Human Genetics, 2019, 105, 1005-1015.	2.6	24
40	Distinct genetic variation and heterogeneity of the Iranian population. PLoS Genetics, 2019, 15, e1008385.	1.5	34
41	Chromosomal aberrations in pregnancy and fetal loss: Insight on the effect of consanguinity, review of 1625 cases. Molecular Genetics & Genomic Medicine, 2019, 7, e820.	0.6	10
42	Homozygous variants in the gene SCAPER cause syndromic intellectual disability. American Journal of Medical Genetics, Part A, 2019, 179, 1214-1225.	0.7	10
43	Identification of diseaseâ€causing variants in the EXOSC gene family underlying autosomal recessive intellectual disability in Iranian families. Clinical Genetics, 2019, 95, 718-725.	1.0	5
44	Whole-Transcriptome Analysis Reveals Dysregulation of Actin-Cytoskeleton Pathway in Intellectual Disability Patients. Neuroscience, 2019, 404, 423-444.	1.1	4
45	A splice-altering variant in LARP7 gene leads to exon exclusion. Gene Reports, 2019, 15, 100375.	0.4	1
46	Homozygous Null TBX4 Mutations Lead to Posterior Amelia with Pelvic and Pulmonary Hypoplasia. American Journal of Human Genetics, 2019, 105, 1294-1301.	2.6	17
47	High expression of miRâ€510 was associated with CGG expansion located at upstream of FMR1 into full mutation. Journal of Cellular Biochemistry, 2019, 120, 1916-1923.	1.2	6
48	SZT2 mutation in a boy with intellectual disability, seizures and autistic features. European Journal of Medical Genetics, 2019, 62, 103556.	0.7	12
49	Effect of inbreeding on intellectual disability revisited by trio sequencing. Clinical Genetics, 2019, 95, 151-159.	1.0	49
50	Novel mutations in MYTH4-FERM domains of myosin 15 are associated with autosomal recessive nonsyndromic hearing loss. International Journal of Pediatric Otorhinolaryngology, 2019, 117, 115-126.	0.4	9
51	Genetics of intellectual disability in consanguineous families. Molecular Psychiatry, 2019, 24, 1027-1039.	4.1	131
52	Investigation of Chromosomal Abnormalities and Microdeletion/ Microduplication(s) in Fifty Iranian Patients with Multiple Congenital Anomalies. Cell Journal, 2019, 21, 337-349.	0.2	1
53	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.	0.3	1
54	Novel Mutations in KCNQ4, LHFPL5 and COCH Genes in Iranian Families with Hearing Impairment. Archives of Iranian Medicine, 2019, 22, 189-197.	0.2	9

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55	Contribution of Iran in Elucidating the Genetic Causes of Autosomal Recessive Intellectual Disability. Archives of Iranian Medicine, 2019, 22, 461-471.	0.2	0
56	Novel mutations in mitochondrial carrier family gene SLC25A38, causing congenital sideroblastic anemia in Iranian families, identified by whole exome sequencing. Blood Cells, Molecules, and Diseases, 2018, 71, 39-44.	0.6	7
57	Exonic mutations and exon skipping: Lessons learned from DFNA5. Human Mutation, 2018, 39, 433-440.	1.1	44
58	Genotype and phenotype analysis of 43 Iranian facioscapulohumeral muscular dystrophy patients; Evidence for anticipation. Neuromuscular Disorders, 2018, 28, 303-314.	0.3	6
59	SLC52A2 mutations cause SCABD2 phenotype: A second report. International Journal of Pediatric Otorhinolaryngology, 2018, 104, 195-199.	0.4	9
60	CDC14A phosphatase is essential for hearing and male fertility in mouse and human. Human Molecular Genetics, 2018, 27, 780-798.	1.4	49
61	Old gene, new phenotype: splice-altering variants in CEACAM16 cause recessive non-syndromic hearing impairment. Journal of Medical Genetics, 2018, 55, 555-560.	1.5	48
62	A novel splicing variant in FLNC gene responsible for a highly penetrant familial dilated cardiomyopathy in an extended Iranian family. Gene, 2018, 659, 160-167.	1.0	14
63	De novo and inherited mutations in the X-linked gene CLCN4 are associated with syndromic intellectual disability and behavior and seizure disorders in males and females. Molecular Psychiatry, 2018, 23, 222-230.	4.1	45
64	Variants in CIB2 cause DFNB48 and not USH1J. Clinical Genetics, 2018, 93, 812-821.	1.0	46
65	CNKSR1 gene defect can cause syndromic autosomal recessive intellectual disability. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 691-699.	1.1	4
66	Frequency of Î±-Globin Gene Triplications and Coinheritance with Î²-Globin Gene Mutations in the Iranian Population. Hemoglobin, 2018, 42, 252-256.	0.4	7
67	Grxcr2 is required for stereocilia morphogenesis in the cochlea. PLoS ONE, 2018, 13, e0201713.	1.1	11
68	The power of the Mediator complex-Expanding the genetic architecture and phenotypic spectrum of MED12 -related disorders. Clinical Genetics, 2018, 94, 450-456.	1.0	24
69	Biallelic missense variants in ZBTB11 can cause intellectual disability in humans. Human Molecular Genetics, 2018, 27, 3177-3188.	1.4	19
70	Intellectual Disability and Ataxia: Genetic Collisions. Archives of Iranian Medicine, 2018, 21, 29-40.	0.2	3
71	Calpains: Diverse Functions but Enigmatic. Archives of Iranian Medicine, 2018, 21, 170-179.	0.2	4
72	tRNA Methyltransferase Defects and Intellectual Disability. Archives of Iranian Medicine, 2018, 21, 478-485.	0.2	5

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73	Improved diagnostic yield of neuromuscular disorders applying clinical exome sequencing in patients arising from a consanguineous population. Clinical Genetics, 2017, 91, 386-402.	1.0	41
74	Mutations of the aminoacyl-tRNA-synthetases SARS and WARS2 are implicated in the etiology of autosomal recessive intellectual disability. Human Mutation, 2017, 38, 621-636.	1.1	54
75	Homozygous Truncating Variants in TBC1D23 Cause Pontocerebellar Hypoplasia and Alter Cortical Development. American Journal of Human Genetics, 2017, 101, 428-440.	2.6	39
76	CIB2, defective in isolated deafness, is key for auditory hair cell mechanotransduction and survival. EMBO Molecular Medicine, 2017, 9, 1711-1731.	3.3	66
77	LGMD2E is the most common type of sarcoglycanopathies in the Iranian population. Journal of Neurogenetics, 2017, 31, 161-169.	0.6	18
78	Copy Number Variants in Patients with Autism and Additional Clinical Features: Report of VIPR2 Duplication and a Novel Microduplication Syndrome. Molecular Neurobiology, 2017, 54, 7019-7027.	1.9	20
79	De novo Mutation in CACNA1S Gene in a 20-Year-Old Man Diagnosed with Metabolic Myopathy. Archives of Iranian Medicine, 2017, 20, 617-620.	0.2	1
80	Whole Genome Linkage Analysis Followed by Whole Exome Sequencing Identifies Nicastrin (NCSTN) as a Causative Gene in a Multiplex Family with Î³-Secretase Spectrum of Autoinflammatory Skin Phenotypes. Journal of Investigative Dermatology, 2016, 136, 1283-1286.	0.3	17
81	Prevalence of common MEFV mutations and carrier frequencies in a large cohort of Iranian populations. Journal of Genetics, 2016, 95, 667-674.	0.4	19
82	Investigating the CFH Gene Polymorphisms as a Risk Factor for Age-related Macular Degeneration in an Iranian Population. Ophthalmic Genetics, 2016, 37, 144-149.	0.5	12
83	Characterization of Homozygous Hb Setif (HBA2: c.283G>T) in the Iranian Population. Hemoglobin, 2016, 40, 53-55.	0.4	2
84	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.	0.3	5
85	First Report of a Dominantly Inherited Î²-Thalassemia Caused by a Novel Elongated Î²-Globin Chain. Hemoglobin, 2016, 40, 102-107.	0.4	6
86	Point mutations which should not be overlooked in Hb H disease. Expert Review of Hematology, 2016, 9, 107-113.	1.0	6
87	Identification of Mutations Causing Aberrant Termination and Deficient Splice Donor Site on theHBA1Gene. Hemoglobin, 2016, 40, 38-43.	0.4	2
88	Missense variants in AIMP1 gene are implicated in autosomal recessive intellectual disability without neurodegeneration. European Journal of Human Genetics, 2016, 24, 392-399.	1.4	17
89	BOD1 Is Required for Cognitive Function in Humans and Drosophila. PLoS Genetics, 2016, 12, e1006022.	1.5	18
90	Heterogeneity of Hereditary Hearing Loss in Iran: a Comprehensive Review. Archives of Iranian Medicine, 2016, 19, 720-728.	0.2	18

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91	Genetic Studies in Intellectual Disability and Behavioral Impairment. Archives of Iranian Medicine, 2016, 19, 363-75.	0.2	10
92	Two novel mutations in ILDR1 gene cause autosomal recessive nonsyndromic hearing loss in consanguineous Iranian families. Journal of Genetics, 2015, 94, 483-487.	0.4	14
93	PDZD7 and hearing loss: More than just a modifier. American Journal of Medical Genetics, Part A, 2015, 167, 2957-2965.	0.7	54
94	Diagnostic pitfalls of less well recognized HbH disease. Blood Cells, Molecules, and Diseases, 2015, 55, 387-395.	0.6	13
95	Association Study of the TREM2 Gene and Identification of a Novel Variant in Exon 2 in Iranian Patients with Late-Onset Alzheimer's Disease. Medical Principles and Practice, 2015, 24, 351-354.	1.1	19
96	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.	0.4	4
97	Mutation Screening of theKrÃ¼ppel-Like Factor 1Gene Using Single-Strand Conformational Polymorphism in a Cohort of IranianÎ²-Thalassemia Patients. Hemoglobin, 2015, 39, 24-29.	0.4	9
98	Homozygous SLC6A17 Mutations Cause Autosomal-Recessive Intellectual Disability with Progressive Tremor, Speech Impairment, and Behavioral Problems. American Journal of Human Genetics, 2015, 96, 386-396.	2.6	27
99	Redefining the MED13L syndrome. European Journal of Human Genetics, 2015, 23, 1308-1317.	1.4	53
100	Mutations in the histamineN-methyltransferase gene,HNMT, are associated with nonsyndromic autosomal recessive intellectual disability. Human Molecular Genetics, 2015, 24, 5697-5710.	1.4	27
101	Interaction of anÎ±-Globin Gene Triplication withÎ²-Globin Gene Mutations in Iranian Patients withÎ²-Thalassemia Intermedia. Hemoglobin, 2015, 39, 201-206.	0.4	21
102	Copy number variations of six and seven Î±-globin genes in a family with intermedia and major thalassemia phenotypes. Expert Review of Hematology, 2015, 8, 693-698.	1.0	9
103	Characterising the spectrum of autosomal recessive hereditary hearing loss in Iran. Journal of Medical Genetics, 2015, 52, 823-829.	1.5	87
104	Hb Dartmouth (HBA2: c.200T>C): An Î±2-Globin Gene Associated with Hb H Disease in One Homozygous Patient. Hemoglobin, 2015, 39, 152-155.	0.4	8
105	A 21 Nucleotide Duplication on theÎ±1- andÎ±2-Globin Genes Involves a Variety of Hypochromic Microcytic Anemias, From Mild to Hb H Disease. Hemoglobin, 2015, 39, 196-200.	0.4	2
106	Mutations on theÎ±2-Globin Gene That May TriggerÎ±+-Thalassemia. Hemoglobin, 2015, 39, 398-402.	0.4	3
107	A defect in the CLIP1 gene (CLIP-170) can cause autosomal recessive intellectual disability. European Journal of Human Genetics, 2015, 23, 331-336.	1.4	22
108	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.	1.1	56

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109	Homozygosity for the AATAAAâ€‰›â€‰AATA-â€‰- Polyadenylation Site Mutation on the Î±2-Globin Gene Causing Transfusion-Dependent Hb H Disease in an Iranian Patient: A Case Report. Hemoglobin, 2015, 39, 355-8.	0.4	8
110	Report of a patient with limb-girdle muscular dystrophy, ptosis and ophthalmoparesis caused by plectinopathy. Archives of Iranian Medicine, 2015, 18, 60-4.	0.2	8
111	New evidence for the role of calpain 10 in autosomal recessive intellectual disability: identification of two novel nonsense variants by exome sequencing in Iranian families. Archives of Iranian Medicine, 2015, 18, 179-84.	0.2	6
112	Carrier Testing in Known Autosomal Recessive Intellectual Disability Genes in an Iranian Healthy Individual Using Exome Sequencing. Archives of Iranian Medicine, 2015, 18, 643-69.	0.2	2
113	Exome Sequencing and Linkage Analysis Identified Novel Candidate Genes in Recessive Intellectual Disability Associated with Ataxia. Archives of Iranian Medicine, 2015, 18, 670-82.	0.2	4
114	Genetic Investigation of an Iranian Supercentenarian by Whole Exome Sequencing. Archives of Iranian Medicine, 2015, 18, 688-97.	0.2	6
115	Impact of whole exome sequencing among Iranian patients with autosomal recessive retinitis pigmentosa. Archives of Iranian Medicine, 2015, 18, 776-85.	0.2	15
116	Evaluation of chromosomal aberrations caused by air pollutants in some taxi drivers from two polluted districts of urban Tehran and its comparison with drivers from rural areas of Lahijan: a pilot study. Journal of Environmental Health Science & Engineering, 2014, 12, 144.	1.4	6
117	Retinitis Pigmentosa, Cutis Laxa, and Pseudoxanthoma Elasticumâ€“Like Skin Manifestations Associated with GGCX Mutations. Journal of Investigative Dermatology, 2014, 134, 2331-2338.	0.3	20
118	Integrated Sequence Analysis Pipeline Provides One-Stop Solution for Identifying Disease-Causing Mutations. Human Mutation, 2014, 35, 1427-1435.	1.1	31
119	A novel mutation of the USH2C (GPR98) gene in an Iranian family with Usher syndrome type II. Journal of Genetics, 2014, 93, 837-841.	0.4	14
120	Pattern of immunoglobulin and T-cell receptor-Î´Íˆ gene rearrangements in Iranian children with B-precursor acute lymphoblastic leukemia. Hematology, 2014, 19, 259-266.	0.7	7
121	Investigation of ATP6V1B1 and ATP6V0A4 genes causing hereditary hearing loss associated with distal renal tubular acidosis in Iranian families. Journal of Laryngology and Otology, 2014, 128, 1056-1059.	0.4	5
122	Genetics of non-syndromic hearing loss in the Middle East. International Journal of Pediatric Otorhinolaryngology, 2014, 78, 2026-2036.	0.4	45
123	NDST1 missense mutations in autosomal recessive intellectual disability. American Journal of Medical Genetics, Part A, 2014, 164, 2753-2763.	0.7	34
124	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.	1.1	33
125	Characterizing a Cohort of Î±-Thalassemia Couples Collected During Screening for Hemoglobinopathies: 14 Years of an Iranian Experience. Hemoglobin, 2014, 38, 153-157.	0.4	9
126	Utilizing Ethnic-Specific Differences in Minor Allele Frequency to Recategorize Reported Pathogenic Deafness Variants. American Journal of Human Genetics, 2014, 95, 445-453.	2.6	137

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127	Identification of a founder mutation for Pendred syndrome in families from northwest Iran. International Journal of Pediatric Otorhinolaryngology, 2014, 78, 1828-1832.	0.4	9
128	Optimizing A Lipocomplex-Based Gene Transfer Method into HeLa Cell Line. Cell Journal, 2014, 15, 372-7.	0.2	5
129	Association of polymorphisms at LDLR locus with coronary artery disease independently from lipid profile. Acta Medica Iranica, 2014, 52, 352-9.	0.8	11
130	Utility Of The Multivariate Approach In Predicting Î²-Thalassemia Intermedia Or Î²-Thalassemia Major Types In Iranian Patients. Hemoglobin, 2013, 37, 413-422.	0.4	12
131	The influence of the BCL11A polymorphism on the phenotype of patients with beta thalassemia could be affected by the beta globin locus control region and/or the Xmn1-HBG2 genotypic background. Blood Cells, Molecules, and Diseases, 2013, 51, 80-84.	0.6	11
132	The Spectrum ofÎ±-Thalassemia Mutations in the Kurdish Population of Northeastern Iraq. Hemoglobin, 2013, 37, 56-64.	0.4	11
133	Hydroxyurea responsiveness in Â-thalassemic patients is determined by the stress response adaptation of erythroid progenitors and their differentiation propensity. Haematologica, 2013, 98, 696-704.	1.7	49
134	Investigation of primary microcephaly in Bushehr province of Iran: novel <scp>STIL</scp> andÂ<scp>ASPM</scp> mutations. Clinical Genetics, 2013, 83, 488-490.	1.0	16
135	A novel ALDH5A1 mutation is associated with succinic semialdehyde dehydrogenase deficiency and severe intellectual disability in an Iranian family. American Journal of Medical Genetics, Part A, 2013, 161, 1915-1922.	0.7	16
136	Genomic characterization of some Iranian children with idiopathic mental retardation using array comparative genomic hybridization. Indian Journal of Human Genetics, 2013, 19, 443.	0.7	1
137	A Mutation Causes MuSK Reduced Sensitivity to Agrin and Congenital Myasthenia. PLoS ONE, 2013, 8, e53826.	1.1	36
138	Identification of Chromosome Abnormalities in Subtelomeric Regions Using Multiplex Ligation Dependent Probe Amplification (MLPA) Technique in 100 Iranian Patients With Idiopathic Mental Retardation. Iranian Red Crescent Medical Journal, 2013, 15, e8221.	0.5	3
139	A Novel Deletion Mutation in ASPM Gene in an Iranian Family with Autosomal Recessive Primary Microcephaly. Iranian Journal of Child Neurology, 2013, 7, 23-30.	0.2	2
140	Novel mutations responsible for Î±-thalassemia in Iranian families. Hemoglobin, 2013, 37, 148-59.	0.4	12
141	NPHP4 Variants Are Associated With Pleiotropic Heart Malformations. Circulation Research, 2012, 110, 1564-1574.	2.0	46
142	Diagnostic values of GHSR DNA methylation pattern in breast cancer. Breast Cancer Research and Treatment, 2012, 135, 705-713.	1.1	16
143	The modifying effect of Xmn1-HBG2 on thalassemic phenotype is associated with its linked elements in the beta globin locus control region, including the palindromic site at 5â€²HS4. Blood Cells, Molecules, and Diseases, 2012, 48, 1-5.	0.6	10
144	The spectrum of GJB2 mutations in the Iranian population with non-syndromic hearing lossâ€”A twelve year study. International Journal of Pediatric Otorhinolaryngology, 2012, 76, 1164-1174.	0.4	38

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145	Spectrum of GJB2 (Cx26) gene mutations in Iranian Azeri patients with nonsyndromic autosomal recessive hearing loss. International Journal of Pediatric Otorhinolaryngology, 2012, 76, 268-271.	0.4	24
146	Two novel SLC26A4 mutations in Iranian families with autosomal recessive hearing loss. International Journal of Pediatric Otorhinolaryngology, 2012, 76, 845-850.	0.4	13
147	Core promoter STRs: Novel mechanism for inter-individual variation in gene expression in humans. Gene, 2012, 492, 195-198.	1.0	25
148	The XmnI andBCL11ASingle Nucleotide Polymorphisms May Help Predict Hydroxyurea Response in Iranian Î²-Thalassemia Patients. Hemoglobin, 2012, 36, 371-380.	0.4	35
149	A comprehensive study to determine heterogeneity of autosomal recessive nonsyndromic hearing loss in Iran. American Journal of Medical Genetics, Part A, 2012, 158A, 2485-2492.	0.7	55
150	Update of the pompe disease mutation database with 60 novel GAA sequence variants and additional studies on the functional effect of 34 previously reported variants. Human Mutation, 2012, 33, 1161-1165.	1.1	67
151	Screening for MYO15A gene mutations in autosomal recessive nonsyndromic, GJB2 negative Iranian deaf population. American Journal of Medical Genetics, Part A, 2012, 158A, 1857-1864.	0.7	54
152	Congenital myasthenic syndrome with tubular aggregates caused by GFPT1 mutations. Journal of Neurology, 2012, 259, 838-850.	1.8	72
153	Mutations in NSUN2 Cause Autosomal- Recessive Intellectual Disability. American Journal of Human Genetics, 2012, 90, 847-855.	2.6	243
154	c<scp>AMP</scp> response elementâ€binding protein 1 is required for hydroxyureaâ€mediated induction of Î³â€globin expression in <scp>K</scp>562 cells. Clinical and Experimental Pharmacology and Physiology, 2012, 39, 510-517.	0.9	8
155	Hemoglobin Q-Iran detected in family members from Northern Iran: a case report. Journal of Medical Case Reports, 2012, 6, 47.	0.4	3
156	Investigation of genetic causes of intellectual disability in kerman province, South East of iran. Iranian Red Crescent Medical Journal, 2012, 14, 79-85.	0.5	1
157	Genotype-Phenotype Correlation In Iranian Patients With Hb H Disease. Hemoglobin, 2011, 35, 40-46.	0.4	20
158	Mutation of the conserved polyadenosine RNA binding protein, ZC3H14/dNab2, impairs neural function in Drosophila and humans. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12390-12395.	3.3	77
159	Exceptional human core promoter nucleotide compositions. Gene, 2011, 475, 79-86.	1.0	19
160	Analyzing 5â€²HS3 and 5â€²HS4 LCR core regions and NF-E2 in Iranian thalassemia intermedia patients with normal or carrier status for beta-globin mutations. Blood Cells, Molecules, and Diseases, 2011, 46, 201-205.	0.6	3
161	Deep sequencing reveals 50 novel genes for recessive cognitive disorders. Nature, 2011, 478, 57-63.	13.7	805
162	Reversion of the human calreticulin gene promoter to the ancestral type as a result of a novel psychosis-associated mutation. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2011, 35, 541-544.	2.5	18

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163	Support for down-tuning of the calreticulin gene in the process of human evolution. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2011, 35, 1770-1773.	2.5	15
164	Reducing the exome search space for Mendelian diseases using genetic linkage analysis of exome genotypes. Genome Biology, 2011, 12, R85.	13.9	72
165	Chromosome abnormality rate among Iranian patients with idiopathic mental retardation from consanguineous marriages. Archives of Medical Science, 2011, 2, 321-325.	0.4	11
166	Next generation sequencing in a family with autosomal recessive Kahrizi syndrome (OMIM 612713) reveals a homozygous frameshift mutation in SRD5A3. European Journal of Human Genetics, 2011, 19, 115-117.	1.4	52
167	The human caveolin 1 gene upstream purine complex and neurodegenerationâ€"A common signature. Journal of Neuroimmunology, 2011, 236, 106-110.	1.1	7
168	Loss-of-Function Mutations of ILDR1 Cause Autosomal-Recessive Hearing Impairment DFNB42. American Journal of Human Genetics, 2011, 88, 127-137.	2.6	108
169	Hexosamine Biosynthetic Pathway Mutations Cause Neuromuscular Transmission Defect. American Journal of Human Genetics, 2011, 88, 162-172.	2.6	153
170	Mutations in the Alpha 1,2-Mannosidase Gene, MAN1B1, Cause Autosomal-Recessive Intellectual Disability. American Journal of Human Genetics, 2011, 89, 176-182.	2.6	73
171	Mutations in the Alpha 1,2-Mannosidase Gene, MAN1B1, Cause Autosomal-Recessive Intellectual Disability. American Journal of Human Genetics, 2011, 89, 348.	2.6	0
172	ST3GAL3 Mutations Impair the Development of Higher Cognitive Functions. American Journal of Human Genetics, 2011, 89, 407-414.	2.6	89
173	Autosomal recessive mental retardation: homozygosity mapping identifies 27 single linkage intervals, at least 14 novel loci and several mutation hotspots. Human Genetics, 2011, 129, 141-148.	1.8	45
174	Homozygous THAP1 mutations as cause of earlyâ€onset generalized dystonia. Movement Disorders, 2011, 26, 858-861.	2.2	39
175	Two Iranian families with a novel mutation in GJB2 causing autosomal dominant nonsyndromic hearing loss. American Journal of Medical Genetics, Part A, 2011, 155, 1202-1211.	0.7	9
176	A novel nonsense mutation in TUSC3 is responsible for nonâ€syndromic autosomal recessive mental retardation in a consanguineous Iranian family. American Journal of Medical Genetics, Part A, 2011, 155, 1976-1980.	0.7	43
177	Did the GJB2 35delG mutation originate in Iran?. American Journal of Medical Genetics, Part A, 2011, 155, 2453-2458.	0.7	15
178	Prevalence of GJB2-associated deafness and outcomes of cochlear implantation in Iran. Journal of Laryngology and Otology, 2011, 125, 455-459.	0.4	18
179	Novel extreme homozygote haplotypes at the human caveolin 1 gene upstream purine complex in sporadic Alzheimer's disease. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 347-349.	1.1	11
180	Common MEFV mutation analysis in 36 Iranian patients with familial Mediterranean fever: clinical and demographic significance. Modern Rheumatology, 2010, 20, 566-572.	0.9	9

#	Article	IF	Citations
181	Mutations in Grxcr1 Are The Basis for Inner Ear Dysfunction in the Pirouette Mouse. American Journal of Human Genetics, 2010, 86, 148-160.	2.6	49
182	Brachyphalangy, polydactyly and tibial aplasia/hypoplasia syndrome (OMIM 609945): case report and review of the literature. European Journal of Pediatrics, 2010, 169, 1535-1539.	1.3	3
183	miRNA mutations are not a common cause of deafness. American Journal of Medical Genetics, Part A, 2010, 152A, 646-652.	0.7	14
184	Novel mutations in the calreticulin gene core promoter and coding sequence in schizoaffective disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 706-709.	1.1	10
185	Carrier frequency of SMA by quantitative analysis of the SMN1 deletion in the Iranian population. European Journal of Neurology, 2010, 17, 160-162.	1.7	18
186	Genetic male infertility and mutation of CATSPER ion channels. European Journal of Human Genetics, 2010, 18, 1178-1184.	1.4	139
187	A novel mutation adjacent to the Bth mouse mutation in the TMC1 gene makes this mouse an excellent model of human deafness at the DFNA36 locus. Clinical Genetics, 2010, 77, 395-398.	1.0	29
188	Variable hearing impairment in a DFNB2 family with a novel MYO7A missense mutation. Clinical Genetics, 2010, 77, 563-571.	1.0	37
189	Epigenetically Deregulated microRNA-375 Is Involved in a Positive Feedback Loop with Estrogen Receptor Î± in Breast Cancer Cells. Cancer Research, 2010, 70, 9175-9184.	0.4	260
190	Mutations in TMC1 are a Common Cause of DFNB7/11 Hearing Loss in the Iranian Population. Annals of Otology, Rhinology and Laryngology, 2010, 119, 830-835.	0.6	29
191	A REPORT OF 8 CASES WITH HEMOGLOBIN H DISEASE IN AN IRANIAN FAMILY. Pediatric Hematology and Oncology, 2010, 27, 405-412.	0.3	4
192	Î²-Thalassemia Mutations in the Kurdish Population of Northeastern Iraq. Hemoglobin, 2010, 34, 469-476.	0.4	27
193	Frequency of Positive XmnI Gî³ polymorphism and coinheritance of common alpha thalassemia mutations do not show statistically significant difference between thalassemia major and intermedia cases with homozygous IVSII-1 mutation. Blood Cells, Molecules, and Diseases, 2010, 44, 95-99.	0.6	20
194	A clinical and molecular genetic study of 112 Iranian families with primary microcephaly. Journal of Medical Genetics, 2010, 47, 823-828.	1.5	87
195	Common MEFV mutation analysis in 36 Iranian patients with familial Mediterranean fever: clinical and demographic significance. Modern Rheumatology, 2010, 20, 566-572.	0.9	11
196	Î±-Thalassemia Mutation Analyses in Mazandaran Province, North Iran. Hemoglobin, 2009, 33, 115-123.	0.4	43
197	Î±-Thalassemia Mutations in Gilan Province, North Iran. Hemoglobin, 2009, 33, 235-241.	0.4	30
198	A large deletion in GPR98 causes type IIC Usher syndrome in male and female members of an Iranian family. Journal of Medical Genetics, 2009, 46, 272-276.	1.5	36

#	Article	IF	Citations
199	Skew in the human caveolin 1 gene upstream purine complex homozygote haplotype compartment in multiple sclerosis. Journal of Neuroimmunology, 2009, 216, 103-107.	1.1	10
200	A novel splice site mutation in the RDX gene causes DFNB24 hearing loss in an Iranian family. American Journal of Medical Genetics, Part A, 2009, 149A, 555-558.	0.7	18
201	A novel polymorphic purine complex at the 1.5 kb upstream region of the human caveolinâ€1 gene and risk of Alzheimer's disease; Extraâ€short alleles and accumulated allele homozygosity. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 248-253.	1.1	16
202	Expanded mutational spectrum in Cohen syndrome, tissue expression, and transcript variants ofCOH1. Human Mutation, 2009, 30, E404-E420.	1.1	44
203	Mutations in the first MyTH4 domain of MYO15A are a common cause of DFNB3 hearing loss. Laryngoscope, 2009, 119, 727-733.	1.1	48
204	Identification of SLC26A4 gene mutations in Iranian families with hereditary hearing impairment. European Journal of Pediatrics, 2009, 168, 651-653.	1.3	30
205	An autosomal recessive syndrome of severe mental retardation, cataract, coloboma and kyphosis maps to the pericentromeric region of chromosome 4. European Journal of Human Genetics, 2009, 17, 125-128.	1.4	17
206	Human Male Infertility Caused by Mutations in the CATSPER1 Channel Protein. American Journal of Human Genetics, 2009, 84, 505-510.	2.6	206
207	Mutations in LOXHD1, an Evolutionarily Conserved Stereociliary Protein, Disrupt Hair Cell Function in Mice and Cause Progressive Hearing Loss in Humans. American Journal of Human Genetics, 2009, 85, 328-337.	2.6	129
208	Identification of Mutations in TRAPPC9, which Encodes the NIK- and IKK-Î²-Binding Protein, in Nonsyndromic Autosomal-Recessive Mental Retardation. American Journal of Human Genetics, 2009, 85, 909-915.	2.6	120
209	Fragile X syndrome screening of families with consanguineous and non-consanguineous parents in the Iranian population. European Journal of Medical Genetics, 2009, 52, 170-173.	0.7	13
210	Molecular Characterization of Î±-Thalassemia in the Dohuk Region of Iraq. Hemoglobin, 2009, 33, 37-44.	0.4	41
211	Sensorineural deafness and male infertility: a contiguous gene deletion syndrome. BMJ Case Reports, 2009, 2009, bcr0820080645-bcr0820080645.	0.2	12
212	GJB2 mutations in Baluchi population. Journal of Genetics, 2008, 87, 195-197.	0.4	17
213	Interstitial deletion of the short arm of chromosome 10 del(10)(p11.2p12.32) in a patient with congenital heart disease, minor dysmorphism, and mental retardation. American Journal of Medical Genetics, Part A, 2008, 146A, 3223-3226.	0.7	1
214	No association between the DAT1 10-repeat allele and ADHD in the Iranian population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 110-111.	1.1	11
215	Identification of a nonsense mutation in the very low-density lipoprotein receptor gene (VLDLR) in an Iranian family with dysequilibrium syndrome. European Journal of Human Genetics, 2008, 16, 270-273.	1.4	50
216	Alopeciamental retardation syndrome: clinical and molecular characterization of four patients. British Journal of Dermatology, 2008, 159, ???-???.	1.4	4

#	Article	IF	Citations
217	A mutation in the calreticulin gene promoter in a family case of schizoaffective disorder leads to its aberrant transcriptional activation. Brain Research, 2008, 1239, 36-41.	1.1	9
218	Prenatal Diagnosis for Î²-Thalassemia Major in the Iranian Province of Hormozgan. Hemoglobin, 2008, 32, 539-545.	0.4	14
219	A Defect in the TUSC3 Gene Is Associated with Autosomal Recessive Mental Retardation. American Journal of Human Genetics, 2008, 82, 1158-1164.	2.6	127
220	M-banding characterization of a 16p11.2p13.1 tandem duplication in a child with autism, neurodevelopmental delay and dysmorphism. European Journal of Medical Genetics, 2008, 51, 608-614.	0.7	14
221	Molecular Mechanisms Underlying Thalassemia Intermedia in Iran. Genetic Testing and Molecular Biomarkers, 2008, 12, 549-556.	1.7	21
222	Î±-Thalassemia Mutations in Khuzestan Province, Southwest Iran. Hemoglobin, 2008, 32, 546-552.	0.4	31
223	A catechol- O -methyltransferase that is essential for auditory function in mice and humans. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 14609-14614.	3.3	62
224	Screening of Iranian Thalassemic Families for the Most Common Deletions of the Î²-Globin Gene Cluster. Hemoglobin, 2007, 31, 463-469.	0.4	4
225	Mutation analysis of the DBC2 gene in sporadic and familial breast cancer. Acta OncolÃ³gica, 2007, 46, 770-772.	0.8	13
226	A Forward Genetics Screen in Mice Identifies Recessive Deafness Traits and Reveals That Pejvakin Is Essential for Outer Hair Cell Function. Journal of Neuroscience, 2007, 27, 2163-2175.	1.7	159
227	Genetic Screening of Leber Congenital Amaurosis in a Large Consanguineous Iranian Family. Ophthalmic Genetics, 2007, 28, 224-228.	0.5	7
228	Sensorineural deafness and male infertility: a contiguous gene deletion syndrome. Journal of Medical Genetics, 2007, 44, 233-240.	1.5	98
229	Validation of a reverse-hybridization StripAssay for the simultaneous analysis of common Î±-thalassemia point mutations and deletions. Clinical Chemistry and Laboratory Medicine, 2007, 45, 605-10.	1.4	33
230	Elucidating the spectrum of Â-thalassemia mutations in Iran. Haematologica, 2007, 92, 992-993.	1.7	52
231	Thalassemia in Iran. Journal of Pediatric Hematology/Oncology, 2007, 29, 233-238.	0.3	181
232	Ethnic-specific distribution of mutations in 716 patients with congenital adrenal hyperplasia owing to 21-hydroxylase deficiency. Molecular Genetics and Metabolism, 2007, 90, 414-421.	0.5	116
233	A Defect in the Ionotropic Glutamate Receptor 6 Gene (GRIK2) Is Associated with Autosomal Recessive Mental Retardation. American Journal of Human Genetics, 2007, 81, 792-798.	2.6	137
234	LRP6 Mutation in a Family with Early Coronary Disease and Metabolic Risk Factors. Science, 2007, 315, 1278-1282.	6.0	567

#	Article	IF	Citations
235	Globin Chain Synthesis is a Useful Complementary Tool in the Differential Diagnosis of Thalassemias. Hemoglobin, 2007, 31, 333-341.	0.4	6
236	Identification of three novelTECTA mutations in Iranian families with autosomal recessive nonsyndromic hearing impairment at the DFNB21 locus. American Journal of Medical Genetics, Part A, 2007, 143A, 1623-1629.	0.7	43
237	Gender dimorphism in the DAT1 â^'67 T-allele homozygosity and predisposition to bipolar disorder. Brain Research, 2007, 1144, 142-145.	1.1	17
238	Homozygosity mapping in consanguineous families reveals extreme heterogeneity of non-syndromic autosomal recessive mental retardation and identifies 8 novel gene loci. Human Genetics, 2007, 121, 43-48.	1.8	92
239	A novel DFNA5 mutation does not cause hearing loss in an Iranian family. Journal of Human Genetics, 2007, 52, 549-552.	1.1	26
240	Fourteen-Year Experience of Prenatal Diagnosis of Thalassemia in Iran. Public Health Genomics, 2006, 9, 93-97.	0.6	55
241	The Coxsackievirus and Adenovirus Receptor: A new adhesion protein in cochlear development. Hearing Research, 2006, 215, 1-9.	0.9	28
242	C677T Methylentetrahydrofulate Reductase and Angiotensin Converting Enzyme Gene Polymorphisms in Patients with Alzheimerâ€™s Disease in Iranian Population. Neurochemical Research, 2006, 31, 1079-1083.	1.6	22
243	SNP array-based homozygosity mapping reveals MCPH1 deletion in family with autosomal recessive mental retardation and mild microcephaly. Human Genetics, 2006, 118, 708-715.	1.8	67
244	Attention-deficit/hyperactivity disorder (ADHD) association with the DAT1 core promoter â^'67 T allele. Brain Research, 2006, 1101, 1-4.	1.1	20
245	The Cypriot and Iranian National Mutation Frequency Databases. Human Mutation, 2006, 27, 598-599.	1.1	32
246	Association of AKT1 haplotype with the risk of schizophrenia in Iranian population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 383-386.	1.1	74
247	A point mutation at the calreticulin gene core promoter conserved sequence in a case of schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 294-295.	1.1	18
248	Determining the cause of patchwork HBA1 and HBA2 genes: recurrent gene conversion or crossing over fixation events. Haematologica, 2006, 91, 297-302.	1.7	22
249	GJB2 mutations: Passage through Iran. American Journal of Medical Genetics, Part A, 2005, 133A, 132-137.	0.7	77
250	Association between the DRD2 A1 allele and opium addiction in the Iranian population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 134B, 39-41.	1.1	32
251	Association analysis of the dopamine transporter (DAT1)-67A/T polymorphism in bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 135B, 47-49.	1.1	31
252	Mutation of COL11A2 causes autosomal recessive non-syndromic hearing loss at the DFNB53 locus. Journal of Medical Genetics, 2005, 42, e61-e61.	1.5	75

#	Article	IF	Citations
253	The frequency of GJB2 mutations and the Î” (GJB6-D13S1830) deletion as a cause of autosomal recessive non-syndromic deafness in the Kurdish population. Clinical Genetics, 2004, 65, 506-508.	1.0	34
254	Association of the dopamine transporter gene (DAT1) core promoter polymorphism ?67T variant with schizophrenia. American Journal of Medical Genetics Part A, 2004, 129B, 10-12.	2.4	32
255	Hb Dhonburi (Neapolis) [Î²126(H4)Valâ†'Gly] Identified in a Family from Northern Iran. Hemoglobin, 2004, 28, 353-356.	0.4	7
256	The Iranian Human Mutation Gene Bank: A data and sample resource for worldwide collaborative genetics research. Human Mutation, 2003, 21, 146-150.	1.1	25
257	A novel homozygous missense mutation in the GNE gene of a patient with quadriceps-sparing hereditary inclusion body myopathy associated with muscle inflammation. Neuromuscular Disorders, 2003, 13, 830-834.	0.3	65
258	High Prevalence of the â^'Î±3.7Deletion Among Thalassemia Patients in Iran. Hemoglobin, 2003, 27, 53-55.	0.4	21
259	alpha-globin gene deletion and point mutation analysis among in Iranian patients with microcytic hypochromic anemia. Haematologica, 2003, 88, 1196-7.	1.7	19
260	GJB2 mutations in Iranians with autosomal recessive non-syndromic sensorineural hearing loss. Human Mutation, 2002, 19, 572-572.	1.1	40
261	Rare and unexpected mutations among Iranian beta-thalassemia patients and prenatal samples discovered by reverse-hybridization and DNA sequencing. Haematologica, 2002, 87, 1113-4.	1.7	26
262	THE Î²-THALASSEMIA MUTATION SPECTRUM IN THE IRANIAN POPULATION. Hemoglobin, 2001, 25, 285-296.	0.4	134
263	Structure and Organization of theRBMYGenes on the Human Y Chromosome: Transposition and Amplification of an Ancestral AutosomalhnRNPGGene. Genomics, 1998, 49, 283-289.	1.3	57
264	Substantial prevalence of microdeletions of the Y-chromosome in infertile men with idiopathic azoospermia and oligozoospermia detected using a sequence-tagged site-based mapping strategy. Journal of Clinical Endocrinology and Metabolism, 1996, 81, 1347-1352.	1.8	133
265	Genomic structure of a Y-specific ribonucleic acid binding motif- containing gene: a putative candidate for a subset of male infertility. Journal of Clinical Endocrinology and Metabolism, 1996, 81, 2159-2164.	1.8	14