

Jamal Nasir

List of Publications by Year in descending order

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45
papers

4,014
citations

257450

24
h-index

265206

42
g-index

46
all docs

46
docs citations

46
times ranked

4907
citing authors

#	ARTICLE	IF	CITATIONS
1	A YAC Mouse Model for Huntington's Disease with Full-Length Mutant Huntingtin, Cytoplasmic Toxicity, and Selective Striatal Neurodegeneration. <i>Neuron</i> , 1999, 23, 181-192.	8.1	789
2	Targeted disruption of the Huntington's disease gene results in embryonic lethality and behavioral and morphological changes in heterozygotes. <i>Cell</i> , 1995, 81, 811-823.	28.9	758
3	A gene encoding a putative GTPase regulator is mutated in familial amyotrophic lateral sclerosis 2. <i>Nature Genetics</i> , 2001, 29, 166-173.	21.4	635
4	Cell death attenuation by 'Usurpin', a mammalian DED-caspase homologue that precludes caspase-8 recruitment and activation by the CD-95 (Fas, APO-1) receptor complex. <i>Cell Death and Differentiation</i> , 1998, 5, 271-288.	11.2	293
5	Interaction between chromatin proteins MECP2 and ATRX is disrupted by mutations that cause inherited mental retardation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 2709-2714.	7.1	231
6	CMIP and ATP2C2 Modulate Phonological Short-Term Memory in Language Impairment. <i>American Journal of Human Genetics</i> , 2009, 85, 264-272.	6.2	173
7	Differential 3' polyadenylation of the Huntington disease gene results in two mRNA species with variable tissue expression. <i>Human Molecular Genetics</i> , 1993, 2, 1541-1545.	2.9	94
8	GA4GH: International policies and standards for data sharing across genomic research and healthcare. <i>Cell Genomics</i> , 2021, 1, 100029.	6.5	94
9	Neuronal degeneration in the basal ganglia and loss of pallido-subthalamic synapses in mice with targeted disruption of the Huntington's disease gene. <i>Brain Research</i> , 1999, 818, 468-479.	2.2	77
10	Targeted disruption of Huntingtin-associated protein-1 (Hap1) results in postnatal death due to depressed feeding behavior. <i>Human Molecular Genetics</i> , 2002, 11, 945-959.	2.9	73
11	Huntington disease: new insights into the relationship between CAG expansion and disease. <i>Human Molecular Genetics</i> , 1996, 5, 1431-1435.	2.9	60
12	Identification of an Alu retrotransposition event in close proximity to a strong candidate gene for Huntington's disease. <i>Nature</i> , 1993, 362, 370-373.	27.8	50
13	Cloning and Characterization of Three Novel Genes, ALS2CR1, ALS2CR2, and ALS2CR3, in the Juvenile Amyotrophic Lateral Sclerosis (ALS2) Critical Region at Chromosome 2q33-q34: Candidate Genes for ALS2. <i>Genomics</i> , 2001, 71, 200-213.	2.9	46
14	Abnormal vibration-induced illusion of movement in idiopathic focal dystonia: An endophenotypic marker?. <i>Movement Disorders</i> , 2008, 23, 373-377.	3.9	45
15	Severe receptive language disorder in childhood—familial aspects and long-term outcomes: results from a Scottish study. <i>Archives of Disease in Childhood</i> , 2007, 92, 614-619.	1.9	43
16	Structural analysis of the 5' region of mouse and human huntington disease genes reveals conservation of putative promoter region and di- and trinucleotide polymorphisms. <i>Genomics</i> , 1995, 25, 707-715.	2.9	41
17	Genomic organization of the human β -adducin gene and its alternately spliced isoforms. <i>Genomics</i> , 1995, 25, 93-99.	2.9	37
18	Life Without Huntingtin. Normal Differentiation into Functional Neurons. <i>Journal of Neurochemistry</i> , 1999, 72, 1009-1018.	3.9	37

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19	Exome analysis identified a novel missense mutation in the CLPP gene in a consanguineous Saudi family expanding the clinical spectrum of Perrault Syndrome type-3. <i>Journal of the Neurological Sciences</i> , 2015, 353, 149-154.	0.6	37
20	De novo single-nucleotide and copy number variation in discordant monozygotic twins reveals disease-related genes. <i>European Journal of Human Genetics</i> , 2019, 27, 1121-1133.	2.8	37
21	The Data Use Ontology to streamline responsible access to human biomedical datasets. <i>Cell Genomics</i> , 2021, 1, 100028.	6.5	31
22	Murine α -L-Iduronidase: cDNA Isolation and Expression. <i>Genomics</i> , 1994, 24, 311-316.	2.9	30
23	A missense mutation in TRAPPC6A leads to build-up of the protein, in patients with a neurodevelopmental syndrome and dysmorphic features. <i>Scientific Reports</i> , 2018, 8, 2053.	3.3	30
24	Case-based interprofessional learning for undergraduate healthcare professionals in the clinical setting. <i>Journal of Interprofessional Care</i> , 2017, 31, 125-128.	1.7	28
25	A mutation in the major autophagy gene, WIPI2, associated with global developmental abnormalities. <i>Brain</i> , 2019, 142, 1242-1254.	7.6	28
26	Unbalanced whole arm translocation resulting in loss of 18p in dystonia. <i>Movement Disorders</i> , 2006, 21, 859-863.	3.9	22
27	The Murine Homologues of the Huntington Disease Gene (Hdh) and the α -Adducin Gene (Add1) Map to Mouse Chromosome 5 within a Region of Conserved Synteny with Human Chromosome 4p16.3. <i>Genomics</i> , 1994, 22, 198-201.	2.9	21
28	Co-amplification of L1 line elements with localised low copy repeats in Giemsa dark bands: implications for genome organisation. <i>Nucleic Acids Research</i> , 1991, 19, 3255-3260.	14.5	19
29	ALG-2 interacting protein AIP1: a novel link between D1 and D3 signalling. <i>European Journal of Neuroscience</i> , 2008, 27, 1626-1633.	2.6	19
30	A Yeast Artificial Chromosome-Based Physical Map of the Juvenile Amyotrophic Lateral Sclerosis (ALS2) Critical Region on Human Chromosome 2q33-q34. <i>Genomics</i> , 1999, 55, 106-112.	2.9	18
31	The alkylglycerol monooxygenase (AGMO) gene previously involved in autism also causes a novel syndromic form of primary microcephaly in a consanguineous Saudi family. <i>Journal of the Neurological Sciences</i> , 2016, 363, 240-244.	0.6	18
32	Truncating mutation in intracellular phospholipase A1 gene (DDHD2) in hereditary spastic paraplegia with intellectual disability (SPG54). <i>BMC Research Notes</i> , 2015, 8, 271.	1.4	17
33	ZizB, a novel RacGEF regulates development, cell motility and cytokinesis in <i>Dictyostelium</i> .. <i>Journal of Cell Science</i> , 2012, 125, 2457-65.	2.0	15
34	Clinically Significant Missense Variants in Human <i>GALNT3</i> , <i>GALNT8</i> , <i>GALNT12</i> , and <i>GALNT13</i> Genes: Intriguing In Silico Findings. <i>Journal of Cellular Biochemistry</i> , 2014, 115, 313-327.	2.6	15
35	Genomic organization of the human caspase-9 gene on Chromosome 1p36.1-p36.3. <i>Mammalian Genome</i> , 1999, 10, 757-760.	2.2	14
36	SARS-CoV-2 Susceptibility and ACE2 Gene Variations Within Diverse Ethnic Backgrounds. <i>Frontiers in Genetics</i> , 2022, 13, 888025.	2.3	14

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37	Interleukin-1 β -converting enzyme (ICE) and related cell death genes ICERel-II and ICERel-III map to the same PAC clone at band 11q22.2-22.3. <i>Mammalian Genome</i> , 1997, 8, 611-613.	2.2	11
38	Distinct proteomic profiles in monozygotic twins discordant for ischaemic stroke. <i>Molecular and Cellular Biochemistry</i> , 2019, 456, 157-165.	3.1	7
39	Mutations in <i>MINAR2</i> encoding membrane integral NOTCH2-associated receptor 2 cause deafness in humans and mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	3
40	Identical non-identical twins and non-identical identical twins. <i>BMJ</i> , The, 2015, , h6589.	6.0	1
41	Elevated \hat{I}^3 -Glutamyltransferase and Erythrocyte Sedimentation Rate in Ischemic Stroke in Discordant Monozygotic Twin Study. <i>International Journal of Stroke</i> , 2015, 10, E32-E33.	5.9	1
42	Overexpression of the dopamine receptor α -interacting protein Alix/AIP1 modulatesNMDAreceptor α -triggered cell death. <i>FEBS Letters</i> , 2019, 593, 1381-1391.	2.8	1
43	Genetics of speech sounds great. <i>Clinical Genetics</i> , 2008, 54, 117-118.	2.0	0
44	Whole exome sequencing reveals a homozygous SGCB variant in a Pakhtun family with limb girdle muscular dystrophy (LGMDR4) phenotype. <i>Gene Reports</i> , 2021, 22, 101014.	0.8	0
45	The Rac GEF ZizB regulates development, cell motility and cytokinesis in <i>Dictyostelium</i> . <i>Development (Cambridge)</i> , 2012, 139, e1-e1.	2.5	0