

Stanley F Nelson

List of Publications by Year in descending order

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Version: 2024-02-01

298
papers

43,282
citations

1883

102
h-index

2375

198
g-index

316
all docs

316
docs citations

316
times ranked

57838
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994. | 9.4 | 2,067 |
| 2 | Shotgun bisulphite sequencing of the Arabidopsis genome reveals DNA methylation patterning. <i>Nature</i> , 2008, 452, 215-219. | 13.7 | 2,039 |
| 3 | Melanomas acquire resistance to B-RAF(V600E) inhibition by RTK or N-RAS upregulation. <i>Nature</i> , 2010, 468, 973-977. | 13.7 | 1,944 |
| 4 | Functional impact of global rare copy number variation in autism spectrum disorders. <i>Nature</i> , 2010, 466, 368-372. | 13.7 | 1,803 |
| 5 | Rare Structural Variants Disrupt Multiple Genes in Neurodevelopmental Pathways in Schizophrenia. <i>Science</i> , 2008, 320, 539-543. | 6.0 | 1,654 |
| 6 | Mapping autism risk loci using genetic linkage and chromosomal rearrangements. <i>Nature Genetics</i> , 2007, 39, 319-328. | 9.4 | 1,272 |
| 7 | Assessing the significance of chromosomal aberrations in cancer: Methodology and application to glioma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 20007-20012. | 3.3 | 927 |
| 8 | High-throughput oncogene mutation profiling in human cancer. <i>Nature Genetics</i> , 2007, 39, 347-351. | 9.4 | 927 |
| 9 | Resolving Individuals Contributing Trace Amounts of DNA to Highly Complex Mixtures Using High-Density SNP Genotyping Microarrays. <i>PLoS Genetics</i> , 2008, 4, e1000167. | 1.5 | 892 |
| 10 | Clinical Exome Sequencing for Genetic Identification of Rare Mendelian Disorders. <i>JAMA - Journal of the American Medical Association</i> , 2014, 312, 1880. | 3.8 | 842 |
| 11 | DNA sequencing of maternal plasma to detect Down syndrome: An international clinical validation study. <i>Genetics in Medicine</i> , 2011, 13, 913-920. | 1.1 | 809 |
| 12 | Linkage, Association, and Gene-Expression Analyses Identify CNTNAP2 as an Autism-Susceptibility Gene. <i>American Journal of Human Genetics</i> , 2008, 82, 150-159. | 2.6 | 738 |
| 13 | Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. <i>Nature Neuroscience</i> , 2015, 18, 199-209. | 7.1 | 701 |
| 14 | Gene Expression Profiling of Gliomas Strongly Predicts Survival. <i>Cancer Research</i> , 2004, 64, 6503-6510. | 0.4 | 659 |
| 15 | Analysis of oncogenic signaling networks in glioblastoma identifies ASPM as a molecular target. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 17402-17407. | 3.3 | 606 |
| 16 | Melanoma whole-exome sequencing identifies V600EB-RAF amplification-mediated acquired B-RAF inhibitor resistance. <i>Nature Communications</i> , 2012, 3, 724. | 5.8 | 567 |
| 17 | A genome-wide scan for common alleles affecting risk for autism. <i>Human Molecular Genetics</i> , 2010, 19, 4072-4082. | 1.4 | 538 |
| 18 | Genome Sequencing Highlights the Dynamic Early History of Dogs. <i>PLoS Genetics</i> , 2014, 10, e1004016. | 1.5 | 481 |

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|----|--|------|-----------|
| 19 | DNA sequencing of maternal plasma reliably identifies trisomy 18 and trisomy 13 as well as Down syndrome: an international collaborative study. <i>Genetics in Medicine</i> , 2012, 14, 296-305. | 1.1 | 471 |
| 20 | Targeted Therapy Resistance Mediated by Dynamic Regulation of Extrachromosomal Mutant EGFR DNA. <i>Science</i> , 2014, 343, 72-76. | 6.0 | 460 |
| 21 | BFAST: An Alignment Tool for Large Scale Genome Resequencing. <i>PLoS ONE</i> , 2009, 4, e7767. | 1.1 | 444 |
| 22 | Cold Urticaria, Immunodeficiency, and Autoimmunity Related to <i>PLCG2</i> Deletions. <i>New England Journal of Medicine</i> , 2012, 366, 330-338. | 13.9 | 391 |
| 23 | Gene Expression Profile Correlates with T-Cell Infiltration and Relative Survival in Glioblastoma Patients Vaccinated with Dendritic Cell Immunotherapy. <i>Clinical Cancer Research</i> , 2011, 17, 1603-1615. | 3.2 | 378 |
| 24 | C-terminal truncations in human DNA exonuclease TREX1 cause autosomal dominant retinal vasculopathy with cerebral leukodystrophy. <i>Nature Genetics</i> , 2007, 39, 1068-1070. | 9.4 | 366 |
| 25 | Mutations in a Human ROBO Gene Disrupt Hindbrain Axon Pathway Crossing and Morphogenesis. <i>Science</i> , 2004, 304, 1509-1513. | 6.0 | 361 |
| 26 | Exome sequencing-based copy-number variation and loss of heterozygosity detection: ExomeCNV. <i>Bioinformatics</i> , 2011, 27, 2648-2654. | 1.8 | 357 |
| 27 | Individual common variants exert weak effects on the risk for autism spectrum disorders. <i>Human Molecular Genetics</i> , 2012, 21, 4781-4792. | 1.4 | 334 |
| 28 | Genome-wide copy number variation study associates metabotropic glutamate receptor gene networks with attention deficit hyperactivity disorder. <i>Nature Genetics</i> , 2012, 44, 78-84. | 9.4 | 334 |
| 29 | Gene connectivity, function, and sequence conservation: predictions from modular yeast co-expression networks. <i>BMC Genomics</i> , 2006, 7, 40. | 1.2 | 327 |
| 30 | Gain-of-function mutations in TRPV4 cause autosomal dominant brachyolmia. <i>Nature Genetics</i> , 2008, 40, 999-1003. | 9.4 | 320 |
| 31 | A Single CRISPR-Cas9 Deletion Strategy that Targets the Majority of DMD Patients Restores Dystrophin Function in hiPSC-Derived Muscle Cells. <i>Cell Stem Cell</i> , 2016, 18, 533-540. | 5.2 | 307 |
| 32 | A Genomewide Scan for Loci Involved in Attention-Deficit/Hyperactivity Disorder. <i>American Journal of Human Genetics</i> , 2002, 70, 1183-1196. | 2.6 | 304 |
| 33 | Identification of inflammatory gene modules based on variations of human endothelial cell responses to oxidized lipids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 12741-12746. | 3.3 | 303 |
| 34 | Epidermal Growth Factor Receptor Activation in Glioblastoma through Novel Missense Mutations in the Extracellular Domain. <i>PLoS Medicine</i> , 2006, 3, e485. | 3.9 | 298 |
| 35 | Gene expression profiling identifies molecular subtypes of gliomas. <i>Oncogene</i> , 2003, 22, 4918-4923. | 2.6 | 264 |
| 36 | Effect of Genetic Diagnosis on Patients with Previously Undiagnosed Disease. <i>New England Journal of Medicine</i> , 2018, 379, 2131-2139. | 13.9 | 261 |

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|----|---|-----|-----------|
| 37 | Whole genome amplification using a degenerate oligonucleotide primer allows hundreds of genotypes to be performed on less than one nanogram of genomic DNA. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 14676-14679. | 3.3 | 256 |
| 38 | Oxysterols Are Novel Activators of the Hedgehog Signaling Pathway in Pluripotent Mesenchymal Cells. Journal of Biological Chemistry, 2007, 282, 8959-8968. | 1.6 | 254 |
| 39 | Identification of molecular subtypes of glioblastoma by gene expression profiling. Oncogene, 2003, 22, 2361-2373. | 2.6 | 247 |
| 40 | A Genetic Analysis of Neural Progenitor Differentiation. Neuron, 2001, 29, 325-339. | 3.8 | 243 |
| 41 | Genome-Wide Analysis of Copy Number Variants in Attention Deficit Hyperactivity Disorder: The Role of Rare Variants and Duplications at 15q13.3. American Journal of Psychiatry, 2012, 169, 195-204. | 4.0 | 242 |
| 42 | U87MG Decoded: The Genomic Sequence of a Cytogenetically Aberrant Human Cancer Cell Line. PLoS Genetics, 2010, 6, e1000832. | 1.5 | 229 |
| 43 | Mutations in the RNA exosome component gene EXOSC3 cause pontocerebellar hypoplasia and spinal motor neuron degeneration. Nature Genetics, 2012, 44, 704-708. | 9.4 | 216 |
| 44 | Primary Glioblastomas Express Mesenchymal Stem-Like Properties. Molecular Cancer Research, 2006, 4, 607-619. | 1.5 | 215 |
| 45 | ISPD loss-of-function mutations disrupt dystroglycan O-mannosylation and cause Walker-Warburg syndrome. Nature Genetics, 2012, 44, 575-580. | 9.4 | 212 |
| 46 | Mutations in PYCR1 cause cutis laxa with progeroid features. Nature Genetics, 2009, 41, 1016-1021. | 9.4 | 211 |
| 47 | Exome Sequencing in the Clinical Diagnosis of Sporadic or Familial Cerebellar Ataxia. JAMA Neurology, 2014, 71, 1237. | 4.5 | 211 |
| 48 | Genetic Linkage of Attention-Deficit/Hyperactivity Disorder on Chromosome 16p13, in a Region Implicated in Autism. American Journal of Human Genetics, 2002, 71, 959-963. | 2.6 | 210 |
| 49 | A Genomewide Scan for Attention-Deficit/Hyperactivity Disorder in an Extended Sample: Suggestive Linkage on 17p11. American Journal of Human Genetics, 2003, 72, 1268-1279. | 2.6 | 206 |
| 50 | Evidence that the dopamine D4 receptor is a susceptibility gene in attention deficit hyperactivity disorder. Molecular Psychiatry, 1998, 3, 427-430. | 4.1 | 198 |
| 51 | SGK196 Is a Glycosylation-Specific <i>O</i> -Mannose Kinase Required for Dystroglycan Function. Science, 2013, 341, 896-899. | 6.0 | 197 |
| 52 | Progressive Ataxia Due to a Missense Mutation in a Calcium-Channel Gene. American Journal of Human Genetics, 1997, 61, 1078-1087. | 2.6 | 191 |
| 53 | Joint mouse-human phenome-wide association to test gene function and disease risk. Nature Communications, 2016, 7, 10464. | 5.8 | 190 |
| 54 | DNA-microarray analysis of brain cancer: molecular classification for therapy. Nature Reviews Neuroscience, 2004, 5, 782-792. | 4.9 | 189 |

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|----|--|-----|-----------|
| 55 | Identification of EpCAM as the Gene for Congenital Tufting Enteropathy. <i>Gastroenterology</i> , 2008, 135, 429-437. | 0.6 | 185 |
| 56 | Distinct Transcription Profiles of Primary and Secondary Glioblastoma Subgroups. <i>Cancer Research</i> , 2006, 66, 159-167. | 0.4 | 182 |
| 57 | Gene expression analysis of glioblastomas identifies the major molecular basis for the prognostic benefit of younger age. <i>BMC Medical Genomics</i> , 2008, 1, 52. | 0.7 | 181 |
| 58 | A novel approach of homozygous haplotype sharing identifies candidate genes in autism spectrum disorder. <i>Human Genetics</i> , 2012, 131, 565-579. | 1.8 | 180 |
| 59 | Insulin growth factor-binding protein 2 is a candidate biomarker for PTEN status and PI3K/Akt pathway activation in glioblastoma and prostate cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 5563-5568. | 3.3 | 173 |
| 60 | Exome Sequencing for the Diagnosis of 46,XY Disorders of Sex Development. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, E333-E344. | 1.8 | 172 |
| 61 | Detection of differentially expressed genes in primary tumor tissues using representational differences analysis coupled to microarray hybridization. <i>Nucleic Acids Research</i> , 1998, 26, 3059-3065. | 6.5 | 169 |
| 62 | Evidence for linkage of a tandem duplication polymorphism upstream of the dopamine D4 receptor gene (DRD4) with attention deficit hyperactivity disorder (ADHD). <i>Molecular Psychiatry</i> , 2000, 5, 531-536. | 4.1 | 169 |
| 63 | Mutations in the PCNA-binding domain of CDKN1C cause IMAGe syndrome. <i>Nature Genetics</i> , 2012, 44, 788-792. | 9.4 | 169 |
| 64 | Joint Analysis of the DRD5 Marker Concludes Association with Attention-Deficit/Hyperactivity Disorder Confined to the Predominantly Inattentive and Combined Subtypes. <i>American Journal of Human Genetics</i> , 2004, 74, 348-356. | 2.6 | 168 |
| 65 | Evidence for Sex-Specific Risk Alleles in Autism Spectrum Disorder. <i>American Journal of Human Genetics</i> , 2004, 75, 1117-1123. | 2.6 | 165 |
| 66 | Meta-analysis of genome-wide linkage scans of attention deficit hyperactivity disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1392-1398. | 1.1 | 160 |
| 67 | CEP41 is mutated in Joubert syndrome and is required for tubulin glutamylation at the cilium. <i>Nature Genetics</i> , 2012, 44, 193-199. | 9.4 | 157 |
| 68 | FXR regulates organic solute transporters $\hat{1}$ and $\hat{2}$ in the adrenal gland, kidney, and intestine. <i>Journal of Lipid Research</i> , 2006, 47, 201-214. | 2.0 | 153 |
| 69 | ERBB3 and NGFR mark a distinct skeletal muscle progenitor cell in human development and hPSCs. <i>Nature Cell Biology</i> , 2018, 20, 46-57. | 4.6 | 151 |
| 70 | Ciliary Abnormalities Due to Defects in the Retrograde Transport Protein DYNC2H1 in Short-Rib Polydactyly Syndrome. <i>American Journal of Human Genetics</i> , 2009, 84, 542-549. | 2.6 | 149 |
| 71 | Genome-wide analysis of single-nucleotide polymorphisms in human expressed sequences. <i>Nature Genetics</i> , 2000, 26, 233-236. | 9.4 | 147 |
| 72 | Multiple self-healing squamous epithelioma is caused by a disease-specific spectrum of mutations in TGFBR1. <i>Nature Genetics</i> , 2011, 43, 365-369. | 9.4 | 147 |

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|----|--|------|-----------|
| 73 | High-throughput profiling of influenza A virus hemagglutinin gene at single-nucleotide resolution. <i>Scientific Reports</i> , 2014, 4, 4942. | 1.6 | 147 |
| 74 | Relationship between Gene Expression and Enhancement in Glioblastoma Multiforme: Exploratory DNA Microarray Analysis. <i>Radiology</i> , 2008, 249, 268-277. | 3.6 | 146 |
| 75 | Maternal embryonic leucine zipper kinase is a key regulator of the proliferation of malignant brain tumors, including brain tumor stem cells. <i>Journal of Neuroscience Research</i> , 2008, 86, 48-60. | 1.3 | 144 |
| 76 | Familial episodic ataxia: Clinical heterogeneity in four families linked to chromosome 19p. <i>Annals of Neurology</i> , 1997, 41, 8-16. | 2.8 | 143 |
| 77 | Replication of Autism Linkage: Fine-Mapping Peak at 17q21. <i>American Journal of Human Genetics</i> , 2005, 76, 1050-1056. | 2.6 | 142 |
| 78 | The Undiagnosed Diseases Network: Accelerating Discovery about Health and Disease. <i>American Journal of Human Genetics</i> , 2017, 100, 185-192. | 2.6 | 142 |
| 79 | Prevalence and Psychiatric Comorbidity of Attention-Deficit/Hyperactivity Disorder in an Adolescent Finnish Population. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2007, 46, 1575-1583. | 0.3 | 141 |
| 80 | The apolipoprotein E 24 allele is not a significant risk factor for frontotemporal dementia. <i>Annals of Neurology</i> , 1998, 44, 134-138. | 2.8 | 136 |
| 81 | Maternal embryonic leucine zipper kinase (MELK) regulates multipotent neural progenitor proliferation. <i>Journal of Cell Biology</i> , 2005, 170, 413-427. | 2.3 | 136 |
| 82 | Fyn and Src Are Effectors of Oncogenic Epidermal Growth Factor Receptor Signaling in Glioblastoma Patients. <i>Cancer Research</i> , 2009, 69, 6889-6898. | 0.4 | 136 |
| 83 | Biomarkers to Predict Antidepressant Response. <i>Current Psychiatry Reports</i> , 2010, 12, 553-562. | 2.1 | 136 |
| 84 | Diagnostic utility of transcriptome sequencing for rare Mendelian diseases. <i>Genetics in Medicine</i> , 2020, 22, 490-499. | 1.1 | 136 |
| 85 | Genomic mismatch scanning: a new approach to genetic linkage mapping. <i>Nature Genetics</i> , 1993, 4, 11-18. | 9.4 | 134 |
| 86 | EGFR Mutation-Induced Alternative Splicing of Max Contributes to Growth of Glycolytic Tumors in Brain Cancer. <i>Cell Metabolism</i> , 2013, 17, 1000-1008. | 7.2 | 130 |
| 87 | Exome Sequencing Identifies PDE4D Mutations in Acrodysostosis. <i>American Journal of Human Genetics</i> , 2012, 90, 746-751. | 2.6 | 128 |
| 88 | Genomic Landscape of Meningiomas. <i>Brain Pathology</i> , 2010, 20, 751-762. | 2.1 | 124 |
| 89 | Lethal Skeletal Dysplasia in Mice and Humans Lacking the Golgin GMAP-210. <i>New England Journal of Medicine</i> , 2010, 362, 206-216. | 13.9 | 122 |
| 90 | Family-Based Genome-Wide Association Scan of Attention-Deficit/Hyperactivity Disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2010, 49, 898-905.e3. | 0.3 | 122 |

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|-----|---|-----|-----------|
| 91 | Attention Deficit Hyperactivity Disorder: Fine Mapping Supports Linkage to 5p13, 6q12, 16p13, and 17p11. <i>American Journal of Human Genetics</i> , 2004, 75, 661-668. | 2.6 | 121 |
| 92 | Assessing the necessity of confirmatory testing for exome-sequencing results in a clinical molecular diagnostic laboratory. <i>Genetics in Medicine</i> , 2014, 16, 510-515. | 1.1 | 121 |
| 93 | A Recessive Skeletal Dysplasia, SEMD Aggrecan Type, Results from a Missense Mutation Affecting the C-Type Lectin Domain of Aggrecan. <i>American Journal of Human Genetics</i> , 2009, 84, 72-79. | 2.6 | 120 |
| 94 | Accuracy of phenotyping of autistic children based on internet implemented parent report. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 1119-1126. | 1.1 | 120 |
| 95 | Annexin A6 modifies muscular dystrophy by mediating sarcolemmal repair. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 6004-6009. | 3.3 | 117 |
| 96 | Disruption of POF1B Binding to Nonmuscle Actin Filaments Is Associated with Premature Ovarian Failure. <i>American Journal of Human Genetics</i> , 2006, 79, 113-119. | 2.6 | 116 |
| 97 | De Novo Nonsense Mutations in KAT6A, a Lysine Acetyl-Transferase Gene, Cause a Syndrome Including Microcephaly and Global Developmental Delay. <i>American Journal of Human Genetics</i> , 2015, 96, 498-506. | 2.6 | 115 |
| 98 | Differential Induction of Glioblastoma Migration and Growth by Two Forms of Pleiotrophin. <i>Journal of Biological Chemistry</i> , 2005, 280, 26953-26964. | 1.6 | 112 |
| 99 | Molecular properties of CD133+ glioblastoma stem cells derived from treatment-refractory recurrent brain tumors. <i>Journal of Neuro-Oncology</i> , 2009, 94, 1-19. | 1.4 | 111 |
| 100 | Relationship between Survival and Edema in Malignant Gliomas: Role of Vascular Endothelial Growth Factor and Neuronal Pentraxin 2. <i>Clinical Cancer Research</i> , 2007, 13, 2592-2598. | 3.2 | 108 |
| 101 | Biased paternal transmission of SNAP-25 risk alleles in attention-deficit hyperactivity disorder. <i>Molecular Psychiatry</i> , 2003, 8, 309-315. | 4.1 | 107 |
| 102 | Transmission disequilibrium testing of dopamine-related candidate gene polymorphisms in ADHD: confirmation of association of ADHD with DRD4 and DRD5. <i>Molecular Psychiatry</i> , 2004, 9, 711-717. | 4.1 | 105 |
| 103 | DYRK1A haploinsufficiency causes a new recognizable syndrome with microcephaly, intellectual disability, speech impairment, and distinct facies. <i>European Journal of Human Genetics</i> , 2015, 23, 1473-1481. | 1.4 | 101 |
| 104 | Association of the cannabinoid receptor gene (CNR1) with ADHD and post-traumatic stress disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1488-1494. | 1.1 | 99 |
| 105 | The functional O-mannose glycan on Î±-dystroglycan contains a phospho-ribitol primed for matriglycan addition. <i>ELife</i> , 2016, 5, . | 2.8 | 98 |
| 106 | Loss of Nardilysin, a Mitochondrial Co-chaperone for Î±-Ketoglutarate Dehydrogenase, Promotes mTORC1 Activation and Neurodegeneration. <i>Neuron</i> , 2017, 93, 115-131. | 3.8 | 95 |
| 107 | Temperament and Character Profiles and the Dopamine D4 Receptor Gene in ADHD. <i>American Journal of Psychiatry</i> , 2005, 162, 906-913. | 4.0 | 94 |
| 108 | PDE5 inhibition alleviates functional muscle ischemia in boys with Duchenne muscular dystrophy. <i>Neurology</i> , 2014, 82, 2085-2091. | 1.5 | 94 |

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|-----|---|-----|-----------|
| 109 | Exome sequencing identifies de novo gain of function missense mutation in KCND2 in identical twins with autism and seizures that slows potassium channel inactivation. <i>Human Molecular Genetics</i> , 2014, 23, 3481-3489. | 1.4 | 90 |
| 110 | Detecting tissue-specific regulation of alternative splicing as a qualitative change in microarray data. <i>Nucleic Acids Research</i> , 2004, 32, e180-e180. | 6.5 | 87 |
| 111 | Stem cell associated gene expression in glioblastoma multiforme: relationship to survival and the subventricular zone. <i>Journal of Neuro-Oncology</i> , 2010, 96, 359-367. | 1.4 | 86 |
| 112 | Next-generation mapping: a novel approach for detection of pathogenic structural variants with a potential utility in clinical diagnosis. <i>Genome Medicine</i> , 2017, 9, 90. | 3.6 | 86 |
| 113 | Pontocerebellar hypoplasia type 1. <i>Neurology</i> , 2013, 80, 438-446. | 1.5 | 84 |
| 114 | Linkage-disequilibrium mapping without genotyping. <i>Nature Genetics</i> , 1998, 18, 225-230. | 9.4 | 82 |
| 115 | Loss of CHSY1, a Secreted FRINGE Enzyme, Causes Syndromic Brachydactyly in Humans via Increased NOTCH Signaling. <i>American Journal of Human Genetics</i> , 2010, 87, 768-778. | 2.6 | 82 |
| 116 | Collaborative analysis of DRD4 and DAT genotypes in population-defined ADHD subtypes. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2005, 46, 1067-1073. | 3.1 | 79 |
| 117 | Mutations in DYNC2LI1 disrupt cilia function and cause short rib polydactyly syndrome. <i>Nature Communications</i> , 2015, 6, 7092. | 5.8 | 79 |
| 118 | Familial migraine with vertigo: No mutations found in CACNA1A. , 1998, 79, 148-151. | | 78 |
| 119 | SeqWare Query Engine: storing and searching sequence data in the cloud. <i>BMC Bioinformatics</i> , 2010, 11, S2. | 1.2 | 78 |
| 120 | Klinefelter's syndrome as a model of anomalous cerebral laterality: Testing gene dosage in the X chromosome pseudoautosomal region using a DNA microarray. , 1998, 23, 215-229. | | 77 |
| 121 | Dantrolene Enhances Antisense-Mediated Exon Skipping in Human and Mouse Models of Duchenne Muscular Dystrophy. <i>Science Translational Medicine</i> , 2012, 4, 164ra160. | 5.8 | 77 |
| 122 | Demographically-Based Evaluation of Genomic Regions under Selection in Domestic Dogs. <i>PLoS Genetics</i> , 2016, 12, e1005851. | 1.5 | 77 |
| 123 | Sequence variant in the laminin β 1 (LAMC1) gene associated with familial pelvic organ prolapse. <i>Human Genetics</i> , 2007, 120, 847-856. | 1.8 | 76 |
| 124 | Proteinase and Growth Factor Alterations Revealed by Gene Microarray Analysis of Human Diabetic Corneas. , 2005, 46, 3604. | | 75 |
| 125 | Identification of allele-specific alternative mRNA processing via transcriptome sequencing. <i>Nucleic Acids Research</i> , 2012, 40, e104-e104. | 6.5 | 74 |
| 126 | Loss of function of SLC25A46 causes lethal congenital pontocerebellar hypoplasia. <i>Brain</i> , 2016, 139, 2877-2890. | 3.7 | 74 |

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|-----|---|-----|-----------|
| 127 | De novo mutation in CACNA1A caused acetazolamide-responsive episodic ataxia. , 1998, 77, 298-301. | | 73 |
| 128 | Genome-wide scan of reading ability in affected sibling pairs with attention-deficit/hyperactivity disorder: unique and shared genetic effects. Molecular Psychiatry, 2004, 9, 485-493. | 4.1 | 72 |
| 129 | Targeted massively parallel sequencing provides comprehensive genetic diagnosis for patients with disorders of sex development. Clinical Genetics, 2013, 83, 35-43. | 1.0 | 72 |
| 130 | IRF2BPL Is Associated with Neurological Phenotypes. American Journal of Human Genetics, 2018, 103, 245-260. | 2.6 | 69 |
| 131 | Loss of annexin A1 expression in human breast cancer detected by multiple high-throughput analyses. Biochemical and Biophysical Research Communications, 2004, 326, 218-227. | 1.0 | 68 |
| 132 | Mutations in IRX5 impair craniofacial development and germ cell migration via SDF1. Nature Genetics, 2012, 44, 709-713. | 9.4 | 68 |
| 133 | KAT6A Syndrome: genotypeâ€“phenotype correlation in 76 patients with pathogenic KAT6A variants. Genetics in Medicine, 2019, 21, 850-860. | 1.1 | 68 |
| 134 | High-density SNP association study of the 17q21 chromosomal region linked to autism identifies CACNA1G as a novel candidate gene. Molecular Psychiatry, 2010, 15, 996-1005. | 4.1 | 67 |
| 135 | Association of Sex With Recurrence of Autism Spectrum Disorder Among Siblings. JAMA Pediatrics, 2017, 171, 1107. | 3.3 | 66 |
| 136 | A New Episodic Ataxia Syndrome With Linkage to Chromosome 19q13. Archives of Neurology, 2007, 64, 749. | 4.9 | 65 |
| 137 | Expanding the phenotype of mutations in DICER1: mosaic missense mutations in the RNase IIIb domain of <i>DICER1</i> cause GLOW syndrome. Journal of Medical Genetics, 2014, 51, 294-302. | 1.5 | 65 |
| 138 | <i>DMD</i> genotype correlations from the Duchenne Registry: Endogenous exon skipping is a factor in prolonged ambulation for individuals with a defined mutation subtype. Human Mutation, 2018, 39, 1193-1202. | 1.1 | 65 |
| 139 | CBCL Pediatric Bipolar Disorder Profile and ADHD: Comorbidity and Quantitative Trait Loci Analysis. Journal of the American Academy of Child and Adolescent Psychiatry, 2008, 47, 1151-1157. | 0.3 | 64 |
| 140 | Differential Gene Expression in Glioblastoma Defined by ADC Histogram Analysis: Relationship to Extracellular Matrix Molecules and Survival. American Journal of Neuroradiology, 2012, 33, 1059-1064. | 1.2 | 64 |
| 141 | Pleiotropic Mechanisms Indicated for Sex Differences in Autism. PLoS Genetics, 2016, 12, e1006425. | 1.5 | 64 |
| 142 | Emerging genetic therapies to treat Duchenne muscular dystrophy. Current Opinion in Neurology, 2009, 22, 532-538. | 1.8 | 63 |
| 143 | Homozygous mutation in <i>NUP107</i> leads to microcephaly with steroid-resistant nephrotic condition similar to Galloway-Mowat syndrome. Journal of Medical Genetics, 2017, 54, 399-403. | 1.5 | 62 |
| 144 | Improved variant discovery through local re-alignment of short-read next-generation sequencing data using SRMA. Genome Biology, 2010, 11, R99. | 3.8 | 61 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|------|-----------|
| 145 | Detection of a MicroRNA Signal in an In Vivo Expression Set of mRNAs. PLoS ONE, 2007, 2, e804. | 1.1 | 61 |
| 146 | A comprehensive iterative approach is highly effective in diagnosing individuals who are exome negative. Genetics in Medicine, 2019, 21, 161-172. | 1.1 | 60 |
| 147 | Expanding the Spectrum of BAF-Related Disorders: De Novo Variants in SMARCC2 Cause a Syndrome with Intellectual Disability and Developmental Delay. American Journal of Human Genetics, 2019, 104, 164-178. | 2.6 | 59 |
| 148 | Expanding the mutational spectrum of LZTR1 in schwannomatosis. European Journal of Human Genetics, 2015, 23, 963-968. | 1.4 | 58 |
| 149 | A diagnostic ceiling for exome sequencing in cerebellar ataxia and related neurological disorders. Human Mutation, 2020, 41, 487-501. | 1.1 | 58 |
| 150 | Molecular Analysis of Glioblastoma: Pathway Profiling and Its Implications for Patient Therapy. Cancer Biology and Therapy, 2003, 2, 242-247. | 1.5 | 57 |
| 151 | High density SNP association study of a major autism linkage region on chromosome 17. Human Molecular Genetics, 2007, 16, 704-715. | 1.4 | 57 |
| 152 | Celsius: a community resource for Affymetrix microarray data. Genome Biology, 2007, 8, R112. | 13.9 | 57 |
| 153 | A genome-wide linkage scan of familial benign recurrent vertigo: linkage to 22q12 with evidence of heterogeneity. Human Molecular Genetics, 2006, 15, 251-258. | 1.4 | 56 |
| 154 | Identification of Retinol Binding Protein 1 Promoter Hypermethylation in Isocitrate Dehydrogenase 1 and 2 Mutant Gliomas. Journal of the National Cancer Institute, 2012, 104, 1458-1469. | 3.0 | 56 |
| 155 | Characterization of transformation related genes in oral cancer cells. Oncogene, 1998, 16, 1921-1930. | 2.6 | 55 |
| 156 | Pooled genome-wide linkage data on 424 ADHD ASPs suggests genetic heterogeneity and a common risk locus at 5p13. Molecular Psychiatry, 2006, 11, 5-8. | 4.1 | 55 |
| 157 | Rare Genomic Variants Link Bipolar Disorder with Anxiety Disorders to CREB-Regulated Intracellular Signaling Pathways. Frontiers in Psychiatry, 2013, 4, 154. | 1.3 | 54 |
| 158 | A comparison of gene expression profiles produced by SAGE, long SAGE, and oligonucleotide chips. Genomics, 2004, 84, 631-636. | 1.3 | 53 |
| 159 | A homozygous loss-of-function CAMK2A mutation causes growth delay, frequent seizures and severe intellectual disability. ELife, 2018, 7, . | 2.8 | 53 |
| 160 | Spectrum of neurodevelopmental disease associated with the GNAO1 guanosine triphosphate-binding region. Epilepsia, 2019, 60, 406-418. | 2.6 | 53 |
| 161 | Ribosomal Proteins RPS11 and RPS20, Two Stress-Response Markers of Glioblastoma Stem Cells, Are Novel Predictors of Poor Prognosis in Glioblastoma Patients. PLoS ONE, 2015, 10, e0141334. | 1.1 | 52 |
| 162 | ADHD Candidate Gene Study in a Population-Based Birth Cohort: Association with DBH and DRD2. Journal of the American Academy of Child and Adolescent Psychiatry, 2007, 46, 1614-1621. | 0.3 | 50 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 163 | De Novo variants in the KMT2A (MLL) gene causing atypical Wiedemann-Steiner syndrome in two unrelated individuals identified by clinical exome sequencing. BMC Medical Genetics, 2014, 15, 49. | 2.1 | 49 |
| 164 | Familial Clustering and DRD4 Effects on Electroencephalogram Measures in Multiplex Families With Attention Deficit/Hyperactivity Disorder. Journal of the American Academy of Child and Adolescent Psychiatry, 2010, 49, 368-377. | 0.3 | 48 |
| 165 | Protective Properties of Radio-Chemoresistant Glioblastoma Stem Cell Clones Are Associated with Metabolic Adaptation to Reduced Glucose Dependence. PLoS ONE, 2013, 8, e80397. | 1.1 | 48 |
| 166 | A novel mutation in KCNA1 causes episodic ataxia without myokymia. Human Mutation, 2004, 24, 536-536. | 1.1 | 47 |
| 167 | Repetitive sequence environment distinguishes housekeeping genes. Gene, 2007, 390, 153-165. | 1.0 | 47 |
| 168 | Phenotypic and Genetic Analysis of a Large Family With Migraine-Associated Vertigo. Headache, 2008, 48, 1460-1467. | 1.8 | 46 |
| 169 | Partial Loss of USP9X Function Leads to a Male Neurodevelopmental and Behavioral Disorder Converging on Transforming Growth Factor β Signaling. Biological Psychiatry, 2020, 87, 100-112. | 0.7 | 42 |
| 170 | Distinct gene expression profiles in adult mouse heart following targeted MAP kinase activation. Physiological Genomics, 2006, 25, 50-59. | 1.0 | 41 |
| 171 | Rich annotation of DNA sequencing variants by leveraging the Ensembl Variant Effect Predictor with plugins. Briefings in Bioinformatics, 2015, 16, 255-264. | 3.2 | 41 |
| 172 | Missense-depleted regions in population exomes implicate ras superfamily nucleotide-binding protein alteration in patients with brain malformation. Npj Genomic Medicine, 2016, 1, . | 1.7 | 41 |
| 173 | SLC35A2-CDG: Functional characterization, expanded molecular, clinical, and biochemical phenotypes of 30 unreported Individuals. Human Mutation, 2019, 40, 908-925. | 1.1 | 39 |
| 174 | Osteogenic oxysterol, 20(<i>S</i>)-hydroxycholesterol, induces notch target gene expression in bone marrow stromal cells. Journal of Bone and Mineral Research, 2010, 25, 782-795. | 3.1 | 38 |
| 175 | Decitabine immunosensitizes human gliomas to NY-ESO-1 specific T lymphocyte targeting through the Fas/Fas Ligand pathway. Journal of Translational Medicine, 2011, 9, 192. | 1.8 | 38 |
| 176 | Autocrine Endothelin-3/Endothelin Receptor B Signaling Maintains Cellular and Molecular Properties of Glioblastoma Stem Cells. Molecular Cancer Research, 2011, 9, 1668-1685. | 1.5 | 38 |
| 177 | Bone morphogenetic protein 7 sensitizes O6-methylguanine methyltransferase expressing-glioblastoma stem cells to clinically relevant dose of temozolomide. Molecular Cancer, 2015, 14, 189. | 7.9 | 38 |
| 178 | A novel ICK mutation causes ciliary disruption and lethal endocrine-cerebro-osteodysplasia syndrome. Cilia, 2016, 5, 8. | 1.8 | 37 |
| 179 | Familial clustering and DRD4 effects on electroencephalogram measures in multiplex families with attention deficit/hyperactivity disorder. Journal of the American Academy of Child and Adolescent Psychiatry, 2010, 49, 368-77. | 0.3 | 37 |
| 180 | Cognitive functioning in affected sibling pairs with ADHD: familial clustering and dopamine genes. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2008, 49, 950-957. | 3.1 | 36 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|------|-----------|
| 181 | Therapeutic Potential of HSP90 Inhibition for Neurofibromatosis Type 2. <i>Clinical Cancer Research</i> , 2013, 19, 3856-3870. | 3.2 | 36 |
| 182 | Disseminated Coccidioidomycosis Treated with Interferon- β and Dupilumab. <i>New England Journal of Medicine</i> , 2020, 382, 2337-2343. | 13.9 | 36 |
| 183 | Association of progesterone receptor with migraine-associated vertigo. <i>Neurogenetics</i> , 2007, 8, 195-200. | 0.7 | 35 |
| 184 | BMPER Mutation in Diaphanospondylodysostosis Identified by Ancestral Autozygosity Mapping and Targeted High-Throughput Sequencing. <i>American Journal of Human Genetics</i> , 2010, 87, 532-537. | 2.6 | 35 |
| 185 | A Recurrent De Novo Variant in NACC1 Causes a Syndrome Characterized by Infantile Epilepsy, Cataracts, and Profound Developmental Delay. <i>American Journal of Human Genetics</i> , 2017, 100, 343-351. | 2.6 | 35 |
| 186 | Improving the efficiency of genomic loci capture using oligonucleotide arrays for high throughput resequencing. <i>BMC Genomics</i> , 2009, 10, 646. | 1.2 | 34 |
| 187 | De novo truncating variants in the intronless IRF2BPL are responsible for developmental epileptic encephalopathy. <i>Genetics in Medicine</i> , 2019, 21, 1008-1014. | 1.1 | 34 |
| 188 | Expanding the genotypic and phenotypic spectrum in a diverse cohort of 104 individuals with Wiedemann-Steiner syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1649-1665. | 0.7 | 34 |
| 189 | Local alignment of two-base encoded DNA sequence. <i>BMC Bioinformatics</i> , 2009, 10, 175. | 1.2 | 32 |
| 190 | Genomic Mismatch Scanning Identifies Human Genomic DNA Shared Identical by Descent. <i>Genomics</i> , 1998, 47, 1-6. | 1.3 | 31 |
| 191 | Robustness of gene expression profiling in glioma specimen samplings and derived cell lines. <i>Molecular Brain Research</i> , 2005, 136, 99-103. | 2.5 | 31 |
| 192 | Loss of the scavenger mRNA decapping enzyme DCPS causes syndromic intellectual disability with neuromuscular defects. <i>Human Molecular Genetics</i> , 2015, 24, 3163-3171. | 1.4 | 31 |
| 193 | Loss of ADAM17 is associated with severe multiorgan dysfunction. <i>Human Pathology</i> , 2015, 46, 923-928. | 1.1 | 31 |
| 194 | Molecular diagnosis of putative Stargardt disease probands by exome sequencing. <i>BMC Medical Genetics</i> , 2012, 13, 67. | 2.1 | 30 |
| 195 | Reciprocal Regulation of the Cardiac Epigenome by Chromatin Structural Proteins Hmgb and Ctcf. <i>Journal of Biological Chemistry</i> , 2016, 291, 15428-15446. | 1.6 | 30 |
| 196 | Disease Gene Characterization through Large-Scale Co-Expression Analysis. <i>PLoS ONE</i> , 2009, 4, e8491. | 1.1 | 30 |
| 197 | Exome Sequencing Finds a Novel PCSK1 Mutation in a Child With Generalized Malabsorptive Diarrhea and Diabetes Insipidus. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2013, 57, 759-767. | 0.9 | 29 |
| 198 | Evaluation of techniques using amplified nucleic acid probes for gene expression profiling. <i>New Biotechnology</i> , 2003, 20, 97-106. | 2.7 | 28 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 199 | Genetically defined EWS/FLI1 model system suggests mesenchymal origin of Ewing's family tumors. <i>Laboratory Investigation</i> , 2008, 88, 1291-1302. | 1.7 | 28 |
| 200 | Rare deleterious mutations are associated with disease in bipolar disorder families. <i>Molecular Psychiatry</i> , 2017, 22, 1009-1014. | 4.1 | 28 |
| 201 | Ppp1r1b-lncRNA inhibits PRC2 at myogenic regulatory genes to promote cardiac and skeletal muscle development in mouse and human. <i>Rna</i> , 2020, 26, 481-491. | 1.6 | 28 |
| 202 | Deletion mapping in Xp21 for patients with complex glycerol kinase deficiency using SNP mapping arrays. <i>Human Mutation</i> , 2007, 28, 235-242. | 1.1 | 27 |
| 203 | CREB regulates Meis1 expression in normal and malignant hematopoietic cells. <i>Leukemia</i> , 2008, 22, 665-667. | 3.3 | 26 |
| 204 | Clinical aggressiveness of malignant gliomas is linked to augmented metabolism of amino acids. <i>Journal of Neuro-Oncology</i> , 2016, 128, 57-66. | 1.4 | 26 |
| 205 | Genomic Identification of Significant Targets in Ciliochoroidal Melanoma. , 2011, 52, 3018. | | 25 |
| 206 | X chromosome exome sequencing reveals a novel <i>ALG13</i> mutation in a nonsyndromic intellectual disability family with multiple affected male siblings. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 164-169. | 0.7 | 25 |
| 207 | Failure to up-regulate transcription of genes necessary for muscle adaptation underlies limb girdle muscular dystrophy 2A (calpainopathy). <i>Human Molecular Genetics</i> , 2016, 25, 2194-2207. | 1.4 | 25 |
| 208 | Online Self-Report Data for Duchenne Muscular Dystrophy Confirms Natural History and Can Be Used to Assess for Therapeutic Benefits. <i>PLOS Currents</i> , 2014, 6, . | 1.4 | 25 |
| 209 | Genome-Wide Association Study of Intelligence: Additive Effects of Novel Brain Expressed Genes. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2012, 51, 432-440.e2. | 0.3 | 24 |
| 210 | Identification of somatic and germline mutations using whole exome sequencing of congenital acute lymphoblastic leukemia. <i>BMC Cancer</i> , 2013, 13, 55. | 1.1 | 24 |
| 211 | Whole exome sequencing of pediatric gastric adenocarcinoma reveals an atypical presentation of Li-Fraumeni syndrome. <i>Pediatric Blood and Cancer</i> , 2013, 60, 570-574. | 0.8 | 24 |
| 212 | Variant in human POFUT1 reduces enzymatic activity and likely causes a recessive microcephaly, global developmental delay with cardiac and vascular features. <i>Glycobiology</i> , 2018, 28, 276-283. | 1.3 | 24 |
| 213 | Cardiac MRI biomarkers for Duchenne muscular dystrophy. <i>Biomarkers in Medicine</i> , 2018, 12, 1271-1289. | 0.6 | 24 |
| 214 | Evidence-Based Consensus and Systematic Review on Reducing the Time to Diagnosis of Duchenne Muscular Dystrophy. <i>Journal of Pediatrics</i> , 2019, 204, 305-313.e14. | 0.9 | 24 |
| 215 | COVID-19 drug repurposing: Summary statistics on current clinical trials and promising untested candidates. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 313-317. | 1.3 | 24 |
| 216 | Functional Consequences of a Novel Variant of PCSK1. <i>PLoS ONE</i> , 2013, 8, e55065. | 1.1 | 24 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 217 | Segmental uniparental isodisomy on 5q32-qter in a patient with childhood-onset schizophrenia. <i>Journal of Medical Genetics</i> , 2006, 43, 887-892. | 1.5 | 23 |
| 218 | Sex-specific influence of DRD 2 on ADHD-type temperament in a large population-based birth cohort. <i>Psychiatric Genetics</i> , 2012, 22, 197-201. | 0.6 | 23 |
| 219 | Clinical exome sequencing in neurogenetic and neuropsychiatric disorders. <i>Annals of the New York Academy of Sciences</i> , 2016, 1366, 49-60. | 1.8 | 23 |
| 220 | GATAD2B-associated neurodevelopmental disorder (GAND): clinical and molecular insights into a NuRD-related disorder. <i>Genetics in Medicine</i> , 2020, 22, 878-888. | 1.1 | 22 |
| 221 | A biotinylated MutS fusion protein and its use in a rapid mutation screening technique. <i>Genetic Analysis, Techniques and Applications</i> , 1996, 13, 105-111. | 1.5 | 21 |
| 222 | High-throughput sequencing of the DBA/2J mouse genome. <i>BMC Bioinformatics</i> , 2010, 11, . | 1.2 | 21 |
| 223 | Identification of Candidate Tumor Oncogenes by Integrative Molecular Analysis of Choroidal Melanoma Fine-Needle Aspiration Biopsy Specimens. <i>JAMA Ophthalmology</i> , 2010, 128, 1170. | 2.6 | 20 |
| 224 | A well-tolerated core needle muscle biopsy process suitable for children and adults. <i>Muscle and Nerve</i> , 2020, 62, 688-698. | 1.0 | 20 |
| 225 | Phenotype Sequencing: Identifying the Genes That Cause a Phenotype Directly from Pooled Sequencing of Independent Mutants. <i>PLoS ONE</i> , 2011, 6, e16517. | 1.1 | 20 |
| 226 | Cartilage-selective genes identified in genome-scale analysis of non-cartilage and cartilage gene expression. <i>BMC Genomics</i> , 2007, 8, 165. | 1.2 | 19 |
| 227 | Heterozygous variants in <i>MYBPC1</i> are associated with an expanded neuromuscular phenotype beyond arthrogryposis. <i>Human Mutation</i> , 2019, 40, 1115-1126. | 1.1 | 19 |
| 228 | Multimarker analysis and imputation of multiple platform pooling-based genome-wide association studies. <i>Bioinformatics</i> , 2008, 24, 1896-1902. | 1.8 | 18 |
| 229 | Truncating mutations in <i>APP</i> cause a distinct neurological phenotype. <i>Annals of Neurology</i> , 2016, 80, 456-460. | 2.8 | 18 |
| 230 | FDA Approval of Eteplirsen for Muscular Dystrophy. <i>JAMA - Journal of the American Medical Association</i> , 2017, 317, 1480. | 3.8 | 18 |
| 231 | Calpain 3 and CaMKII β signaling are required to induce HSP70 necessary for adaptive muscle growth after atrophy. <i>Human Molecular Genetics</i> , 2018, 27, 1642-1653. | 1.4 | 18 |
| 232 | Bi-allelic TMEM94 Truncating Variants Are Associated with Neurodevelopmental Delay, Congenital Heart Defects, and Distinct Facial Dysmorphism. <i>American Journal of Human Genetics</i> , 2018, 103, 948-967. | 2.6 | 18 |
| 233 | Efficient discovery of single-nucleotide polymorphisms in coding regions of human genes. <i>Pharmacogenomics Journal</i> , 2002, 2, 236-242. | 0.9 | 17 |
| 234 | Genomic predictors of remission to antidepressant treatment in geriatric depression using genome-wide expression analyses: a pilot study. <i>International Journal of Geriatric Psychiatry</i> , 2016, 31, 510-517. | 1.3 | 17 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 235 | ZBED4, a BED-Type Zinc-Finger Protein in the Cones of the Human Retina. , 2009, 50, 3580. | | 16 |
| 236 | Commonalities across computational workflows for uncovering explanatory variants in undiagnosed cases. <i>Genetics in Medicine</i> , 2021, 23, 1075-1085. | 1.1 | 16 |
| 237 | Genotype-phenotype correlations and novel molecular insights into the DHX30-associated neurodevelopmental disorders. <i>Genome Medicine</i> , 2021, 13, 90. | 3.6 | 16 |
| 238 | High Throughput Screening for Compounds That Alter Muscle Cell Glycosylation Identifies New Role for N-Glycans in Regulating Sarcolemmal Protein Abundance and Laminin Binding. <i>Journal of Biological Chemistry</i> , 2012, 287, 22759-22770. | 1.6 | 15 |
| 239 | The case for eteplirsen: Paving the way for precision medicine. <i>Molecular Genetics and Metabolism</i> , 2016, 118, 70-71. | 0.5 | 15 |
| 240 | Targeting RyR Activity Boosts Antisense Exon 44 and 45 Skipping in Human DMD Skeletal or Cardiac Muscle Culture Models. <i>Molecular Therapy - Nucleic Acids</i> , 2019, 18, 580-589. | 2.3 | 15 |
| 241 | Identification of novel candidate genes for 46,XY disorders of sex development (DSD) using a C57BL/6J-Y POS mouse model. <i>Biology of Sex Differences</i> , 2018, 9, 8. | 1.8 | 14 |
| 242 | Sex chromosome anomalies in childhood onset schizophrenia: an update. <i>Molecular Psychiatry</i> , 2008, 13, 910-911. | 4.1 | 13 |
| 243 | Calculating the statistical significance of rare variants causal for Mendelian and complex disorders. <i>BMC Medical Genomics</i> , 2018, 11, 53. | 0.7 | 13 |
| 244 | Quantitative immuno-mass spectrometry imaging of skeletal muscle dystrophin. <i>Scientific Reports</i> , 2021, 11, 1128. | 1.6 | 13 |
| 245 | A Path to Implement Precision Child Health Cardiovascular Medicine. <i>Frontiers in Cardiovascular Medicine</i> , 2017, 4, 36. | 1.1 | 12 |
| 246 | Loss of IRF2BPL impairs neuronal maintenance through excess Wnt signaling. <i>Science Advances</i> , 2022, 8, eabl5613. | 4.7 | 12 |
| 247 | Genomic Organization, 5' Flanking Enhancer Region, and Chromosomal Assignment of the Cell Cycle Gene, p53Cdc. <i>Molecular Genetics and Metabolism</i> , 1998, 64, 52-57. | 0.5 | 11 |
| 248 | Predicting the severity of Duchenne muscular dystrophy. <i>Neurology</i> , 2011, 76, 208-209. | 1.5 | 11 |
| 249 | A toolkit for genetics providers in follow-up of patients with non-diagnostic exome sequencing. <i>Journal of Genetic Counseling</i> , 2019, 28, 213-228. | 0.9 | 11 |
| 250 | Gene-environment regulatory circuits of right ventricular pathology in tetralogy of fallot. <i>Journal of Molecular Medicine</i> , 2019, 97, 1711-1722. | 1.7 | 11 |
| 251 | Repurposing Dantrolene for Long-Term Combination Therapy to Potentiate Antisense-Mediated DMD Exon Skipping in the mdx Mouse. <i>Molecular Therapy - Nucleic Acids</i> , 2018, 11, 180-191. | 2.3 | 10 |
| 252 | Mitchell-Riley syndrome iPSC exhibit reduced pancreatic endoderm differentiation due to an <i>RFX6</i> mutation. <i>Development (Cambridge)</i> , 2020, 147, . | 1.2 | 10 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 253 | Variants in PRKAR1B cause a neurodevelopmental disorder with autism spectrum disorder, apraxia, and insensitivity to pain. <i>Genetics in Medicine</i> , 2021, 23, 1465-1473. | 1.1 | 10 |
| 254 | Genomic mismatch scanning: Current progress and potential applications. <i>Electrophoresis</i> , 1995, 16, 279-285. | 1.3 | 9 |
| 255 | Whole exome sequencing detects homozygosity for ABCA4 p.Arg602Trp missense mutation in a pediatric patient with rapidly progressive retinal dystrophy. <i>BMC Medical Genetics</i> , 2014, 15, 11. | 2.1 | 9 |
| 256 | Exome Sequencing Identified a Splice Site Mutation in <i>FHL1</i> that Causes Uruguay Syndrome, an X-Linked Disorder With Skeletal Muscle Hypertrophy and Premature Cardiac Death. <i>Circulation: Cardiovascular Genetics</i> , 2016, 9, 130-135. | 5.1 | 8 |
| 257 | IgG4-related disease: Association with a rare gene variant expressed in cytotoxic T cells. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e686. | 0.6 | 8 |
| 258 | Congenital myasthenic syndrome caused by a frameshift insertion mutation in <i>GFPT1</i> . <i>Neurology: Genetics</i> , 2020, 6, e468. | 0.9 | 8 |
| 259 | Representational oligonucleotide microarray analysis (ROMA) and comparison of binning and change-point methods of analysis: application to detection of del22q11.2 (DiGeorge) Syndrome. <i>Human Mutation</i> , 2008, 29, 176-181. | 1.1 | 7 |
| 260 | Expansion of <i>NEUROD2</i> phenotypes to include developmental delay without seizures. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1076-1080. | 0.7 | 7 |
| 261 | Characterization of three cell lines derived from fine needle biopsy of choroidal melanoma with metastatic outcome. <i>Molecular Vision</i> , 2011, 17, 607-15. | 1.1 | 7 |
| 262 | What can Duchenne Connect teach us about treating Duchenne muscular dystrophy?. <i>Current Opinion in Neurology</i> , 2015, 28, 535-541. | 1.8 | 6 |
| 263 | Dusp6 is a genetic modifier of growth through enhanced ERK activity. <i>Human Molecular Genetics</i> , 2018, 28, 279-289. | 1.4 | 6 |
| 264 | Magnetic Resonance Imaging characteristics in case of TOR1AIP1 muscular dystrophy. <i>Clinical Imaging</i> , 2019, 58, 108-113. | 0.8 | 6 |
| 265 | Large in-frame 5â€² deletions in DMD associated with mild Duchenne muscular dystrophy: Two case reports and a review of the literature. <i>Neuromuscular Disorders</i> , 2019, 29, 863-873. | 0.3 | 6 |
| 266 | The frontiers of sequencing in undiagnosed neurodevelopmental diseases. <i>Current Opinion in Genetics and Development</i> , 2020, 65, 76-83. | 1.5 | 6 |
| 267 | Myopathy associated with homozygous <i>PYROXD1</i> pathogenic variants detected by genome sequencing. <i>Neuropathology</i> , 2020, 40, 302-307. | 0.7 | 6 |
| 268 | Mosaic de novo <i>SNRPN</i> gene variant associated with Prader-Willi syndrome. <i>Journal of Medical Genetics</i> , 2022, 59, 719-722. | 1.5 | 6 |
| 269 | Novel <i>NUDT2</i> variant causes intellectual disability and polyneuropathy. <i>Annals of Clinical and Translational Neurology</i> , 2020, 7, 2320-2325. | 1.7 | 5 |
| 270 | MPEG1/Perforin-2 Haploinsufficiency Associated Polymicrobial Skin Infections and Considerations for Interferon-Î³ Therapy. <i>Frontiers in Immunology</i> , 2020, 11, 601584. | 2.2 | 5 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 271 | EIF3F-related neurodevelopmental disorder: refining the phenotypic and expanding the molecular spectrum. <i>Orphanet Journal of Rare Diseases</i> , 2021, 16, 136. | 1.2 | 5 |
| 272 | Rethinking clinical practice: clinical implementation of exome sequencing. <i>Personalized Medicine</i> , 2012, 9, 785-787. | 0.8 | 4 |
| 273 | Validation and Detection of Exon Skipping Boosters in DMD Patient Cell Models and mdx Mouse. <i>Methods in Molecular Biology</i> , 2018, 1828, 309-326. | 0.4 | 4 |
| 274 | Genetic characterization and long-term management of severely affected siblings with intellectual developmental disorder with cardiac arrhythmia syndrome. <i>Molecular Genetics and Metabolism Reports</i> , 2020, 23, 100582. | 0.4 | 4 |
| 275 | Recessive ciliopathy mutations in primary endocardial fibroelastosis: a rare neonatal cardiomyopathy in a case of Alstrom syndrome. <i>Journal of Molecular Medicine</i> , 2021, 99, 1623-1638. | 1.7 | 4 |
| 276 | Modeling Patient-Specific Muscular Dystrophy Phenotypes and Therapeutic Responses in Reprogrammed Myotubes Engineered on Micromolded Gelatin Hydrogels. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 830415. | 1.8 | 4 |
| 277 | Mutation in TWINKLE in a Large Iranian Family with Progressive External Ophthalmoplegia, Myopathy, Dysphagia and Dysphonia, and Behavior Change. <i>Archives of Iranian Medicine</i> , 2016, 19, 87-91. | 0.2 | 4 |
| 278 | DNA Sequencing of Maternal Plasma to Detect Down Syndrome. <i>Obstetrical and Gynecological Survey</i> , 2012, 67, 86-88. | 0.2 | 3 |
| 279 | Leveraging ancestry to improve causal variant identification in exome sequencing for monogenic disorders. <i>European Journal of Human Genetics</i> , 2016, 24, 113-119. | 1.4 | 3 |
| 280 | In silico enhanced restriction enzyme based methylation analysis of the human glioblastoma genome using Agilent 244K CpG Island microarrays. <i>Frontiers in Neuroscience</i> , 2009, 3, 57. | 1.4 | 2 |
| 281 | Kcnd2 Mutation Associated with Autism and Epilepsy Impairs Inactivation Gating in Kv4.2 K+ Channels. <i>Biophysical Journal</i> , 2014, 106, 741a. | 0.2 | 2 |
| 282 | Familial Clustering and DRD4 Effects on Electroencephalogram Measures in Multiplex Families With Attention Deficit/Hyperactivity Disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2010, 49, 368-377. | 0.3 | 2 |
| 283 | Downregulation of SATB1 by miRNAs reduces megakaryocyte/erythroid progenitor expansion in preclinical models of Diamond-Blackfan anemia. <i>Experimental Hematology</i> , 2022, 111, 66-78. | 0.2 | 2 |
| 284 | Selective surface treatment of micro printing pin and its performance. <i>Applied Physics Letters</i> , 2006, 89, 083901. | 1.5 | 1 |
| 285 | Local alignment of generalized k-base encoded DNA sequence. <i>BMC Bioinformatics</i> , 2010, 11, 347. | 1.2 | 1 |
| 286 | Interpreting Whole-Genome Sequencing. <i>JAMA - Journal of the American Medical Association</i> , 2014, 312, 296. | 3.8 | 1 |
| 287 | Case Report: Whole Exome Sequencing Identifies Compound Heterozygous Variants in TSFM Gene Causing Juvenile Hypertrophic Cardiomyopathy. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 798985. | 1.1 | 1 |
| 288 | The dopamine receptor DRD4 gene. <i>Trends in Pharmacological Sciences</i> , 2001, 22, 56. | 4.0 | 0 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 289 | 735. Functional Restoration of Dystrophin Protein in HiPSC-Derived Skeletal Myotubes and Cardiomyocytes After CRISPR/Cas9-Mediated Deletion of 530-725kb of DMD. <i>Molecular Therapy</i> , 2016, 24, S290. | 3.7 | 0 |
| 290 | Gene-environment regulation of chamber-specific maturation during hypoxemic perinatal circulatory transition. <i>Journal of Molecular Medicine</i> , 2020, 98, 1009-1020. | 1.7 | 0 |
| 291 | Identification of Meis1 as a Target of CREB in Myeloid Leukemogenesis.. <i>Blood</i> , 2006, 108, 1945-1945. | 0.6 | 0 |
| 292 | Requirement of CREB in Normal and Malignant Hematopoiesis.. <i>Blood</i> , 2006, 108, 1168-1168. | 0.6 | 0 |
| 293 | Abstract 4749: Genetic signature of malignant glioma is associated with responsiveness to autologous tumor lysate-pulsed dendritic cell vaccination + TLR agonists. , 2010, , . | | 0 |
| 294 | Abstract LB-245: Loss of function of the TGF β RI receptor leads to the spontaneously regressing squamous carcinoma condition, multiple self-healing squamous epithelioma (Ferguson-Smith disease). , 2011, , . | | 0 |
| 295 | Abstract 447: The IDH1 mutation in human glioblastoma and its effects on epigenetic modification and cell fate selection. , 2014, , . | | 0 |
| 296 | Abstract 08: Clinical exome sequencing in the diagnosis of pediatric cancer predisposition. , 2014, , . | | 0 |
| 297 | Role of Mir-34 Upregulation in Disruption of c-Myc, c-Myb and NOTCH Signaling in Diamond-Blackfan Anemia. <i>Blood</i> , 2016, 128, 3895-3895. | 0.6 | 0 |
| 298 | RABENOSYN separation-of-function mutations uncouple endosomal recycling from lysosomal degradation, causing a distinct Mendelian Disorder. <i>Human Molecular Genetics</i> , 0, , . | 1.4 | 0 |