

Matthew J Gazzellone

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

1,677
citations

623734

14
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

4195
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | De novo and rare inherited copy-number variations in the hemiplegic form of cerebral palsy. <i>Genetics in Medicine</i> , 2018, 20, 172-180. | 2.4 | 82 |
| 2 | Genome-wide copy number variation analysis identifies novel candidate loci associated with pediatric obesity. <i>European Journal of Human Genetics</i> , 2018, 26, 1588-1596. | 2.8 | 23 |
| 3 | Rare Genome-Wide Copy Number Variation and Expression of Schizophrenia in 22q11.2 Deletion Syndrome. <i>American Journal of Psychiatry</i> , 2017, 174, 1054-1063. | 7.2 | 77 |
| 4 | Microduplications at the pseudoautosomal <i>SHOX</i> locus in autism spectrum disorders and related neurodevelopmental conditions. <i>Journal of Medical Genetics</i> , 2016, 53, 536-547. | 3.2 | 26 |
| 5 | Uncovering obsessive-compulsive disorder risk genes in a pediatric cohort by high-resolution analysis of copy number variation. <i>Journal of Neurodevelopmental Disorders</i> , 2016, 8, 36. | 3.1 | 55 |
| 6 | Indexing Effects of Copy Number Variation on Genes Involved in Developmental Delay. <i>Scientific Reports</i> , 2016, 6, 28663. | 3.3 | 35 |
| 7 | Whole-Genome Sequencing Suggests Schizophrenia Risk Mechanisms in Humans with 22q11.2 Deletion Syndrome. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 2453-2461. | 1.8 | 43 |
| 8 | Whole-genome sequencing of quartet families with autism spectrum disorder. <i>Nature Medicine</i> , 2015, 21, 185-191. | 30.7 | 457 |
| 9 | A high-resolution copy-number variation resource for clinical and population genetics. <i>Genetics in Medicine</i> , 2015, 17, 747-752. | 2.4 | 73 |
| 10 | Clinically relevant copy number variations detected in cerebral palsy. <i>Nature Communications</i> , 2015, 6, 7949. | 12.8 | 120 |
| 11 | Molecular Diagnostic Yield of Chromosomal Microarray Analysis and Whole-Exome Sequencing in Children With Autism Spectrum Disorder. <i>JAMA - Journal of the American Medical Association</i> , 2015, 314, 895. | 7.4 | 352 |
| 12 | Disruption of the <i>ASTN2/TRIM32</i> locus at 9q33.1 is a risk factor in males for autism spectrum disorders, ADHD and other neurodevelopmental phenotypes. <i>Human Molecular Genetics</i> , 2014, 23, 2752-2768. | 2.9 | 140 |
| 13 | Copy number variation in Han Chinese individuals with autism spectrum disorder. <i>Journal of Neurodevelopmental Disorders</i> , 2014, 6, 34. | 3.1 | 55 |
| 14 | Rare exonic deletions implicate the synaptic organizer Gephyrin (<i>GPHN</i>) in risk for autism, schizophrenia and seizures. <i>Human Molecular Genetics</i> , 2013, 22, 2055-2066. | 2.9 | 139 |