Michael J Gandal

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

 74
 7,203
 36
 84

 papers
 citations
 h-index
 g-index

 98
 10,639
 15.5
 5.53

 ext. papers
 ext. citations
 avg, IF
 L-index

| # | Paper | IF | Citations |
|----------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 74 | Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. <i>Nature Genetics</i> , 2019 , 51, 63-75 | 36.3 | 826 |
| 73 | Shared molecular neuropathology across major psychiatric disorders parallels polygenic overlap. <i>Science</i> , 2018 , 359, 693-697 | 33.3 | 547 |
| 7 2 | Transcriptome-wide isoform-level dysregulation in ASD, schizophrenia, and bipolar disorder. <i>Science</i> , 2018 , 362, | 33.3 | 434 |
| 71 | Human Gut Microbiota from Autism Spectrum Disorder Promote Behavioral Symptoms in Mice. <i>Cell</i> , 2019 , 177, 1600-1618.e17 | 56.2 | 379 |
| 70 | Genome-wide changes in lncRNA, splicing, and regional gene expression patterns in autism. <i>Nature</i> , 2016 , 540, 423-427 | 50.4 | 362 |
| 69 | Chromosome conformation elucidates regulatory relationships in developing human brain. <i>Nature</i> , 2016 , 538, 523-527 | 50.4 | 334 |
| 68 | Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , 2018 , 362, | 33.3 | 319 |
| 67 | Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018 , 362, | 33.3 | 277 |
| 66 | Systems biology and gene networks in neurodevelopmental and neurodegenerative disorders. <i>Nature Reviews Genetics</i> , 2015 , 16, 441-58 | 30.1 | 270 |
| 65 | Deletion of vanilloid receptor 1-expressing primary afferent neurons for pain control. <i>Journal of Clinical Investigation</i> , 2004 , 113, 1344-52 | 15.9 | 252 |
| 64 | Validating ©scillations and delayed auditory responses as translational biomarkers of autism. <i>Biological Psychiatry</i> , 2010 , 68, 1100-6 | 7.9 | 218 |
| 63 | Gamma synchrony: towards a translational biomarker for the treatment-resistant symptoms of schizophrenia. <i>Neuropharmacology</i> , 2012 , 62, 1504-18 | 5.5 | 206 |
| 62 | MEG detection of delayed auditory evoked responses in autism spectrum disorders: towards an imaging biomarker for autism. <i>Autism Research</i> , 2010 , 3, 8-18 | 5.1 | 172 |
| 61 | Ketamine modulates theta and gamma oscillations. <i>Journal of Cognitive Neuroscience</i> , 2010 , 22, 1452-64 | 43.1 | 164 |
| 60 | GABAB-mediated rescue of altered excitatory-inhibitory balance, gamma synchrony and behavioral deficits following constitutive NMDAR-hypofunction. <i>Translational Psychiatry</i> , 2012 , 2, e142 | 8.6 | 144 |
| 59 | Inherited and De Novo Genetic Risk for Autism Impacts Shared Networks. <i>Cell</i> , 2019 , 178, 850-866.e26 | 56.2 | 142 |
| 58 | Transcriptome and epigenome landscape of human cortical development modeled in organoids. <i>Science</i> , 2018 , 362, | 33.3 | 142 |

(2012-2009)

| 57 | N-methyl-d-aspartic acid receptor antagonist-induced frequency oscillations in mice recreate pattern of electrophysiological deficits in schizophrenia. <i>Neuroscience</i> , 2009 , 158, 705-12 | 3.9 | 135 |
|----|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----|
| 56 | The contribution of rare variants to risk of schizophrenia in individuals with and without intellectual disability. <i>Nature Genetics</i> , 2017 , 49, 1167-1173 | 36.3 | 132 |
| 55 | The road to precision psychiatry: translating genetics into disease mechanisms. <i>Nature Neuroscience</i> , 2016 , 19, 1397-1407 | 25.5 | 131 |
| 54 | Mouse behavioral endophenotypes for schizophrenia. <i>Brain Research Bulletin</i> , 2010 , 83, 147-61 | 3.9 | 127 |
| 53 | mGluR5-antagonist mediated reversal of elevated stereotyped, repetitive behaviors in the VPA model of autism. <i>PLoS ONE</i> , 2011 , 6, e26077 | 3.7 | 117 |
| 52 | A genome-wide association study of shared risk across psychiatric disorders implicates gene regulation during fetal neurodevelopment. <i>Nature Neuroscience</i> , 2019 , 22, 353-361 | 25.5 | 93 |
| 51 | Mice with reduced NMDA receptor expression: more consistent with autism than schizophrenia?. <i>Genes, Brain and Behavior</i> , 2012 , 11, 740-50 | 3.6 | 88 |
| 50 | Dysbindin-1 mutant mice implicate reduced fast-phasic inhibition as a final common disease mechanism in schizophrenia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, E962-70 | 11.5 | 83 |
| 49 | Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021 , 53, 817-829 | 36.3 | 83 |
| 48 | Parvalbumin cell ablation of NMDA-R1 causes increased resting network excitability with associated social and self-care deficits. <i>Neuropsychopharmacology</i> , 2014 , 39, 1603-13 | 8.7 | 81 |
| 47 | Pyramidal cell selective ablation of N-methyl-D-aspartate receptor 1 causes increase in cellular and network excitability. <i>Biological Psychiatry</i> , 2015 , 77, 556-68 | 7.9 | 75 |
| 46 | In vitro-in vivo correlations of scalable PLGA-risperidone implants for the treatment of schizophrenia. <i>Pharmaceutical Research</i> , 2010 , 27, 1730-7 | 4.5 | 75 |
| 45 | Genetic Control of Expression and Splicing in Developing Human Brain Informs Disease Mechanisms. <i>Cell</i> , 2019 , 179, 750-771.e22 | 56.2 | 70 |
| 44 | A novel electrophysiological model of chemotherapy-induced cognitive impairments in mice. <i>Neuroscience</i> , 2008 , 157, 95-104 | 3.9 | 64 |
| 43 | NMDA antagonists recreate signal-to-noise ratio and timing perturbations present in schizophrenia. <i>Neurobiology of Disease</i> , 2012 , 46, 93-100 | 7.5 | 63 |
| 42 | Discovery of the first genome-wide significant risk loci for ADHD | | 62 |
| 41 | Cannabis-induced psychosis associated with high potency "wax dabs". <i>Schizophrenia Research</i> , 2016 , 172, 211-2 | 3.6 | 56 |
| 40 | Measuring the maturity of the fast-spiking interneuron transcriptional program in autism, schizophrenia, and bipolar disorder. <i>PLoS ONE</i> , 2012 , 7, e41215 | 3.7 | 49 |

| 39 | MeCP2+/- mouse model of RTT reproduces auditory phenotypes associated with Rett syndrome and replicate select EEG endophenotypes of autism spectrum disorder. <i>Neurobiology of Disease</i> , 2012 , 46, 88-92 | 7.5 | 40 |
|----|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|----|
| 38 | Mapping genomic loci implicates genes and synaptic biology in schizophrenia <i>Nature</i> , 2022 , | 50.4 | 35 |
| 37 | Strong correlation of downregulated genes related to synaptic transmission and mitochondria in post-mortem autism cerebral cortex. <i>Journal of Neurodevelopmental Disorders</i> , 2018 , 10, 18 | 4.6 | 31 |
| 36 | NMDA antagonist MK801 recreates auditory electrophysiology disruption present in autism and other neurodevelopmental disorders. <i>Behavioural Brain Research</i> , 2012 , 234, 233-7 | 3.4 | 30 |
| 35 | Convergence of circuit dysfunction in ASD: a common bridge between diverse genetic and environmental risk factors and common clinical electrophysiology. <i>Frontiers in Cellular Neuroscience</i> , 2014 , 8, 414 | 6.1 | 25 |
| 34 | Integrative genomics identifies a convergent molecular subtype that links epigenomic with transcriptomic differences in autism. <i>Nature Communications</i> , 2020 , 11, 4873 | 17.4 | 23 |
| 33 | Synaptic and Gene Regulatory Mechanisms in Schizophrenia, Autism, and 22q11.2 Copy Number Variant-Mediated Risk for Neuropsychiatric Disorders. <i>Biological Psychiatry</i> , 2020 , 87, 150-163 | 7.9 | 23 |
| 32 | A Robust Method Uncovers Significant Context-Specific Heritability in Diverse Complex Traits. <i>American Journal of Human Genetics</i> , 2020 , 106, 71-91 | 11 | 21 |
| 31 | Identification of neural oscillations and epileptiform changes in human brain organoids. <i>Nature Neuroscience</i> , 2021 , 24, 1488-1500 | 25.5 | 20 |
| 30 | Profiling allele-specific gene expression in brains from individuals with autism spectrum disorder reveals preferential minor allele usage. <i>Nature Neuroscience</i> , 2019 , 22, 1521-1532 | 25.5 | 17 |
| 29 | Integrative network analysis reveals biological pathways associated with Williams syndrome. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2019 , 60, 585-598 | 7.9 | 14 |
| 28 | Quantum computing at the frontiers of biological sciences. <i>Nature Methods</i> , 2021 , 18, 701-709 | 21.6 | 14 |
| 27 | Transcriptomic Insight Into the Polygenic Mechanisms Underlying Psychiatric Disorders. <i>Biological Psychiatry</i> , 2021 , 89, 54-64 | 7.9 | 12 |
| 26 | Mechanisms of Neuronal Alternative Splicing and Strategies for Therapeutic Interventions. <i>Journal of Neuroscience</i> , 2019 , 39, 8193-8199 | 6.6 | 11 |
| 25 | Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology | | 11 |
| 24 | Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2021 , | 7.9 | 11 |
| 23 | Spatial fine-mapping for gene-by-environment effects identifies risk hot spots for schizophrenia. <i>Nature Communications</i> , 2018 , 9, 5296 | 17.4 | 10 |
| 22 | Nicotine normalizes event related potentials in COMT-Val-tg mice and increases gamma and theta spectral density. <i>Behavioral Neuroscience</i> , 2012 , 126, 332-43 | 2.1 | 9 |

| 21 | Brain gene co-expression networks link complement signaling with convergent synaptic pathology in schizophrenia. <i>Nature Neuroscience</i> , 2021 , 24, 799-809 | 25.5 | 9 |
|--------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------|---------------------------------------------------|
| 20 | Alterations in Retrotransposition, Synaptic Connectivity, and Myelination Implicated by Transcriptomic Changes Following Maternal Immune Activation in Nonhuman Primates. <i>Biological Psychiatry</i> , 2021 , 89, 896-910 | 7.9 | 7 |
| 19 | Shared Molecular Neuropathology Across Major Psychiatric Disorders Parallels Polygenic Overlap. <i>Focus (American Psychiatric Publishing)</i> , 2019 , 17, 66-72 | 1.1 | 6 |
| 18 | Full-length transcript sequencing of human and mouse cerebral cortex identifies widespread isoform diversity and alternative splicing. <i>Cell Reports</i> , 2021 , 37, 110022 | 10.6 | 5 |
| 17 | Whole genome sequencing in multiplex families reveals novel inherited and de novo genetic risk in auti | ism | 5 |
| 16 | Maternal Immune Activation during Pregnancy Alters Postnatal Brain Growth and Cognitive Development in Nonhuman Primate Offspring. <i>Journal of Neuroscience</i> , 2021 , 41, 9971-9987 | 6.6 | 3 |
| 15 | Genetic control of gene expression and splicing in the developing human brain | | 3 |
| 14 | Shared molecular neuropathology across major psychiatric disorders parallels polygenic overlap | | 3 |
| 13 | TGFBuperfamily signaling regulates the state of human stem cell pluripotency and competency to create telencephalic organoids | | 3 |
| | | | |
| 12 | Broad transcriptomic dysregulation across the cerebral cortex in ASD | | 3 |
| 12 11 | Broad transcriptomic dysregulation across the cerebral cortex in ASD Polygenicity in Psychiatry-Like It or Not, We Have to Understand It. <i>Biological Psychiatry</i> , 2021 , 89, 2-4 | 7.9 | 3 |
| | | | |
| 11 | Polygenicity in Psychiatry-Like It or Not, We Have to Understand It. <i>Biological Psychiatry</i> , 2021 , 89, 2-4 | | 3 |
| 11 | Polygenicity in Psychiatry-Like It or Not, We Have to Understand It. <i>Biological Psychiatry</i> , 2021 , 89, 2-4 The Genetics-Driven Revival in Neuropsychiatric Drug Development. <i>Biological Psychiatry</i> , 2016 , 79, 628 Human Astrocytes Exhibit Tumor Microenvironment-, Age-, and Sex-Related Transcriptomic | 3- 3 @ | 3 |
| 11 10 9 | Polygenicity in Psychiatry-Like It or Not, We Have to Understand It. <i>Biological Psychiatry</i> , 2021 , 89, 2-4 The Genetics-Driven Revival in Neuropsychiatric Drug Development. <i>Biological Psychiatry</i> , 2016 , 79, 628 Human Astrocytes Exhibit Tumor Microenvironment-, Age-, and Sex-Related Transcriptomic Signatures <i>Journal of Neuroscience</i> , 2022 , Brain gene co-expression networks link complement signaling with convergent synaptic pathology | 3- 3 @ | 3222 |
| 11 10 9 | Polygenicity in Psychiatry-Like It or Not, We Have to Understand It. <i>Biological Psychiatry</i> , 2021 , 89, 2-4 The Genetics-Driven Revival in Neuropsychiatric Drug Development. <i>Biological Psychiatry</i> , 2016 , 79, 628 Human Astrocytes Exhibit Tumor Microenvironment-, Age-, and Sex-Related Transcriptomic Signatures <i>Journal of Neuroscience</i> , 2022 , Brain gene co-expression networks link complement signaling with convergent synaptic pathology in schizophrenia Neuronal and glial 3D chromatin architecture informs the cellular etiology of brain disorders. | 6.6 | 2 2 2 |
| 11 10 9 8 | Polygenicity in Psychiatry-Like It or Not, We Have to Understand It. <i>Biological Psychiatry</i> , 2021 , 89, 2-4 The Genetics-Driven Revival in Neuropsychiatric Drug Development. <i>Biological Psychiatry</i> , 2016 , 79, 628 Human Astrocytes Exhibit Tumor Microenvironment-, Age-, and Sex-Related Transcriptomic Signatures <i>Journal of Neuroscience</i> , 2022 , Brain gene co-expression networks link complement signaling with convergent synaptic pathology in schizophrenia Neuronal and glial 3D chromatin architecture informs the cellular etiology of brain disorders. <i>Nature Communications</i> , 2021 , 12, 3968 Associations between patterns in comorbid diagnostic trajectories of individuals with schizophrenia | 6.6 17.4 | 2 2 2 |

| 2 | Postnatal immune activation causes social deficits in a mouse model of tuberous sclerosis: Role of microglia and clinical implications. <i>Science Advances</i> , 2021 , 7, eabf2073 | 14.3 1 |
|---|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------|
| 1 | Banking on Polygenicity to Disentangle Psychiatric Comorbidity. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2018 , 3, 577-578 | 3.4 |

Human Astrocytes Exhibit Tumor Microenvironment-, Age-, and Sex-Related Transcriptomic Signatures