

Raphael A Bernier

List of Publications by Year in descending order

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Version: 2024-02-01

85
papers

8,156
citations

101543

36
h-index

58581

82
g-index

90
all docs

90
docs citations

90
times ranked

12207
citing authors

#	ARTICLE	IF	CITATIONS
1	Impact of autism genetic risk on brain connectivity: a mechanism for the female protective effect. <i>Brain</i> , 2022, 145, 378-387.	7.6	9
2	Brief Report: Associations Between Self-injurious Behaviors and Abdominal Pain Among Individuals with ASD-Associated Disruptive Mutations. <i>Journal of Autism and Developmental Disorders</i> , 2021, 51, 3365-3373.	2.7	5
3	Brief Report: Can a Composite Heart Rate Variability Biomarker Shed New Insights About Autism Spectrum Disorder in School-Aged Children?. <i>Journal of Autism and Developmental Disorders</i> , 2021, 51, 346-356.	2.7	11
4	The gap between IQ and adaptive functioning in autism spectrum disorder: Disentangling diagnostic and sex differences. <i>Autism</i> , 2021, 25, 1565-1579.	4.1	23
5	Clinical delineation of SETBP1 haploinsufficiency disorder. <i>European Journal of Human Genetics</i> , 2021, 29, 1198-1205.	2.8	12
6	A neurogenetic analysis of female autism. <i>Brain</i> , 2021, 144, 1911-1926.	7.6	24
7	Sleep Problems in Children with ASD and Gene Disrupting Mutations. <i>Journal of Genetic Psychology</i> , 2021, 182, 317-334.	1.2	3
8	Reflections on the genetics-first approach to advancements in molecular genetic and neurobiological research on neurodevelopmental disorders. <i>Journal of Neurodevelopmental Disorders</i> , 2021, 13, 24.	3.1	12
9	The CHD8/CHD7/Kismet family links blood-brain barrier glia and serotonin to ASD-associated sleep defects. <i>Science Advances</i> , 2021, 7, .	10.3	24
10	Recent ultra-rare inherited variants implicate new autism candidate risk genes. <i>Nature Genetics</i> , 2021, 53, 1125-1134.	21.4	68
11	Coding and noncoding variants in EBF3 are involved in HADDs and simplex autism. <i>Human Genomics</i> , 2021, 15, 44.	2.9	16
12	Pathogenic SPTBN1 variants cause an autosomal dominant neurodevelopmental syndrome. <i>Nature Genetics</i> , 2021, 53, 1006-1021.	21.4	44
13	Clinical Phenotypes of Carriers of Mutations in CHD8 or Its Conserved Target Genes. <i>Biological Psychiatry</i> , 2020, 87, 123-131.	1.3	22
14	Language characterization in 16p11.2 deletion and duplication syndromes. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 380-391.	1.7	16
15	Late fMRI Response Components Are Altered in Autism Spectrum Disorder. <i>Frontiers in Human Neuroscience</i> , 2020, 14, 241.	2.0	5
16	Episignatures Stratifying Helsmoortel-Van Der Aa Syndrome Show Modest Correlation with Phenotype. <i>American Journal of Human Genetics</i> , 2020, 107, 555-563.	6.2	32
17	NCKAP1 Disruptive Variants Lead to a Neurodevelopmental Disorder with Core Features of Autism. <i>American Journal of Human Genetics</i> , 2020, 107, 963-976.	6.2	18
18	Neural responsivity to social rewards in autistic female youth. <i>Translational Psychiatry</i> , 2020, 10, 178.	4.8	22

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19	Weaker neural suppression in autism. <i>Nature Communications</i> , 2020, 11, 2675.	12.8	28
20	Imaging-genetics of sex differences in ASD: distinct effects of OXTR variants on brain connectivity. <i>Translational Psychiatry</i> , 2020, 10, 82.	4.8	31
21	Evaluating heterogeneity in <sc>ASD</sc> symptomatology, cognitive ability, and adaptive functioning among 16p11.2 <sc>CNV</sc> carriers. <i>Autism Research</i> , 2020, 13, 1300-1310.	3.8	23
22	Response Dissociation in Hierarchical Cortical Circuits: a Unique Feature of Autism Spectrum Disorder. <i>Journal of Neuroscience</i> , 2020, 40, 2269-2281.	3.6	13
23	Day-to-Day Test-Retest Reliability of EEG Profiles in Children With Autism Spectrum Disorder and Typical Development. <i>Frontiers in Integrative Neuroscience</i> , 2020, 14, 21.	2.1	32
24	Sex Differences in Functional Connectivity of the Salience, Default Mode, and Central Executive Networks in Youth with ASD. <i>Cerebral Cortex</i> , 2020, 30, 5107-5120.	2.9	46
25	Co-occurring medical conditions among individuals with ASD-associated disruptive mutations. <i>Children's Health Care</i> , 2020, 49, 361-384.	0.9	3
26	Concentrations of Cortical <sc>GABA</sc> and Glutamate in Young Adults With Autism Spectrum Disorder. <i>Autism Research</i> , 2020, 13, 1111-1129.	3.8	38
27	A framework for an evidence-based gene list relevant to autism spectrum disorder. <i>Nature Reviews Genetics</i> , 2020, 21, 367-376.	16.3	83
28	Large-scale targeted sequencing identifies risk genes for neurodevelopmental disorders. <i>Nature Communications</i> , 2020, 11, 4932.	12.8	105
29	The state of research on the genetics of autism spectrum disorder: methodological, clinical and conceptual progress. <i>Current Opinion in Psychology</i> , 2019, 27, 1-5.	4.9	27
30	AMPA receptor GluA2 subunit defects are a cause of neurodevelopmental disorders. <i>Nature Communications</i> , 2019, 10, 3094.	12.8	150
31	Disruptive mutations in TANC2 define a neurodevelopmental syndrome associated with psychiatric disorders. <i>Nature Communications</i> , 2019, 10, 4679.	12.8	43
32	The Human-Specific BOLA2 Duplication Modifies Iron Homeostasis and Anemia Predisposition in Chromosome 16p11.2 Autism Individuals. <i>American Journal of Human Genetics</i> , 2019, 105, 947-958.	6.2	30
33	Disruptive variants of <i>CSDE1</i> associate with autism and interfere with neuronal development and synaptic transmission. <i>Science Advances</i> , 2019, 5, eaax2166.	10.3	35
34	Psychiatric disorders in children with 16p11.2 deletion and duplication. <i>Translational Psychiatry</i> , 2019, 9, 8.	4.8	93
35	Î±7 Nicotinic Acetylcholine Receptor Signaling Modulates Ovine Fetal Brain Astrocytes Transcriptome in Response to Endotoxin. <i>Frontiers in Immunology</i> , 2019, 10, 1063.	4.8	18
36	Methodological considerations in the use of Noldus EthoVision XT video tracking of children with autism in multi-site studies. <i>Biological Psychology</i> , 2019, 146, 107712.	2.2	10

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37	Habituation Learning Is a Widely Affected Mechanism in Drosophila Models of Intellectual Disability and Autism Spectrum Disorders. <i>Biological Psychiatry</i> , 2019, 86, 294-305.	1.3	39
38	Sex-Based Analysis of De Novo Variants in Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , 2019, 105, 1274-1285.	6.2	84
39	Neurodevelopmental disease genes implicated by de novo mutation and copy number variation morbidity. <i>Nature Genetics</i> , 2019, 51, 106-116.	21.4	231
40	Glutamatergic facilitation of neural responses in MT enhances motion perception in humans. <i>NeuroImage</i> , 2019, 184, 925-931.	4.2	16
41	Genome sequencing identifies multiple deleterious variants in autism patients with more severe phenotypes. <i>Genetics in Medicine</i> , 2019, 21, 1611-1620.	2.4	88
42	Clinical Presentation of a Complex Neurodevelopmental Disorder Caused by Mutations in ADNP. <i>Biological Psychiatry</i> , 2019, 85, 287-297.	1.3	108
43	Progress in Understanding and Treating SCN2A-Mediated Disorders. <i>Trends in Neurosciences</i> , 2018, 41, 442-456.	8.6	210
44	Longitudinal report of child with de novo 16p11.2 triplication. <i>Clinical Case Reports (discontinued)</i> , 2018, 6, 147-154.	0.5	5
45	Interdisciplinary Team Evaluation: An Effective Method for the Diagnostic Assessment of Autism Spectrum Disorder. <i>Journal of Developmental and Behavioral Pediatrics</i> , 2018, 39, 271-281.	1.1	18
46	Child and family characteristics moderate agreement between caregiver and clinician report of autism symptoms. <i>Autism Research</i> , 2018, 11, 476-487.	3.8	19
47	Truncating Variants in NAA15 Are Associated with Variable Levels of Intellectual Disability, Autism Spectrum Disorder, and Congenital Anomalies. <i>American Journal of Human Genetics</i> , 2018, 102, 985-994.	6.2	59
48	Early enhanced processing and delayed habituation to deviance sounds in autism spectrum disorder. <i>Brain and Cognition</i> , 2018, 123, 110-119.	1.8	53
49	Comorbid symptoms of inattention, autism, and executive cognition in youth with putative genetic risk. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2018, 59, 268-276.	5.2	8
50	A genotype-first approach identifies an intellectual disability-overweight syndrome caused by PHIP haploinsufficiency. <i>European Journal of Human Genetics</i> , 2018, 26, 54-63.	2.8	32
51	Inherited and multiple de novo mutations in autism/developmental delay risk genes suggest a multifactorial model. <i>Molecular Autism</i> , 2018, 9, 64.	4.9	114
52	Auditory perception is associated with implicit language learning and receptive language ability in autism spectrum disorder. <i>Brain and Language</i> , 2018, 187, 1-8.	1.6	9
53	Suppression and facilitation of human neural responses. <i>ELife</i> , 2018, 7, .	6.0	48
54	Sex Differences in Visual Motion Processing. <i>Current Biology</i> , 2018, 28, 2794-2799.e3.	3.9	35

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55	The autism spectrum phenotype in ADNP syndrome. <i>Autism Research</i> , 2018, 11, 1300-1310.	3.8	49
56	Reduced auditory cortical adaptation in autism spectrum disorder. <i>ELife</i> , 2018, 7, .	6.0	41
57	Targeted sequencing identifies 91 neurodevelopmental-disorder risk genes with autism and developmental-disability biases. <i>Nature Genetics</i> , 2017, 49, 515-526.	21.4	443
58	Hotspots of missense mutation identify neurodevelopmental disorder genes and functional domains. <i>Nature Neuroscience</i> , 2017, 20, 1043-1051.	14.8	152
59	Developmental trajectories for young children with 16p11.2 copy number variation. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 367-380.	1.7	42
60	denovo-db: a compendium of human<i>de novo</i>variants. <i>Nucleic Acids Research</i> , 2017, 45, D804-D811.	14.5	173
61	Gene Disrupting Mutations Associated with Regression in Autism Spectrum Disorder. <i>Journal of Autism and Developmental Disorders</i> , 2017, 47, 3600-3607.	2.7	26
62	Exploring the heterogeneity of neural social indices for genetically distinct etiologies of autism. <i>Journal of Neurodevelopmental Disorders</i> , 2017, 9, 24.	3.1	19
63	Associations between Familial Rates of Psychiatric Disorders and De Novo Genetic Mutations in Autism. <i>Autism Research & Treatment</i> , 2017, 2017, 1-9.	0.5	4
64	Prospective investigation of FOXP1 syndrome. <i>Molecular Autism</i> , 2017, 8, 57.	4.9	65
65	Clinical phenotype of ASD-associated DYRK1A haploinsufficiency. <i>Molecular Autism</i> , 2017, 8, 54.	4.9	55
66	Autism Spectrum Disorder, Developmental and Psychiatric Features in 16p11.2 Duplication. <i>Journal of Autism and Developmental Disorders</i> , 2016, 46, 2734-2748.	2.7	47
67	Children with Autism Show Altered Autonomic Adaptation to Novel and Familiar Social Partners. <i>Autism Research</i> , 2016, 9, 579-591.	3.8	41
68	Infants's™ grip strength predicts mu rhythm attenuation during observation of lifting actions with weighted blocks. <i>Developmental Science</i> , 2016, 19, 195-207.	2.4	10
69	The Number of Genomic Copies at the 16p11.2 Locus Modulates Language, Verbal Memory, and Inhibition. <i>Biological Psychiatry</i> , 2016, 80, 129-139.	1.3	78
70	Disruption of POGZ Is Associated with Intellectual Disability and Autism Spectrum Disorders. <i>American Journal of Human Genetics</i> , 2016, 98, 541-552.	6.2	132
71	Maternal Modifiers and Parent-of-Origin Bias of the Autism-Associated 16p11.2 CNV. <i>American Journal of Human Genetics</i> , 2016, 98, 45-57.	6.2	55
72	Face processing among twins with and without autism: social correlates and twin concordance. <i>Social Cognitive and Affective Neuroscience</i> , 2016, 11, 44-54.	3.0	24

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73	Defining the Effect of the 16p11.2 Duplication on Cognition, Behavior, and Medical Comorbidities. <i>JAMA Psychiatry</i> , 2016, 73, 20.	11.0	195
74	Clinical phenotype of the recurrent 1q21.1 copy-number variant. <i>Genetics in Medicine</i> , 2016, 18, 341-349.	2.4	134
75	Modulation of mu attenuation to social stimuli in children and adults with 16p11.2 deletions and duplications. <i>Journal of Neurodevelopmental Disorders</i> , 2015, 7, 25.	3.1	12
76	Electrodermal Response to Reward and Non-Reward Among Children With Autism. <i>Autism Research</i> , 2015, 8, 357-370.	3.8	22
77	The Role of Parental Cognitive, Behavioral, and Motor Profiles in Clinical Variability in Individuals With Chromosome 16p11.2 Deletions. <i>JAMA Psychiatry</i> , 2015, 72, 119.	11.0	112
78	Excess of rare, inherited truncating mutations in autism. <i>Nature Genetics</i> , 2015, 47, 582-588.	21.4	531
79	The Cognitive and Behavioral Phenotype of the 16p11.2 Deletion in a Clinically Ascertained Population. <i>Biological Psychiatry</i> , 2015, 77, 785-793.	1.3	198
80	The transcriptional regulator <i>ADNP</i> links the BAF (SWI/SNF) complexes with autism. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2014, 166, 315-326.	1.6	68
81	De novo TBR1 mutations in sporadic autism disrupt protein functions. <i>Nature Communications</i> , 2014, 5, 4954.	12.8	109
82	Disruptive CHD8 Mutations Define a Subtype of Autism Early in Development. <i>Cell</i> , 2014, 158, 263-276.	28.9	637
83	A 600-kb deletion syndrome at 16p11.2 leads to energy imbalance and neuropsychiatric disorders. <i>Journal of Medical Genetics</i> , 2012, 49, 660-668.	3.2	251
84	Sporadic autism exomes reveal a highly interconnected protein network of de novo mutations. <i>Nature</i> , 2012, 485, 246-250.	27.8	1,960
85	The Broader Autism Phenotype and Its Implications on the Etiology and Treatment of Autism Spectrum Disorders. <i>Autism Research & Treatment</i> , 2011, 2011, 1-19.	0.5	103