

T Conrad Gilliam

List of Publications by Year in descending order

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35
papers

11,389
citations

172457

29
h-index

361022

35
g-index

35
all docs

35
docs citations

35
times ranked

12254
citing authors

#	ARTICLE	IF	CITATIONS
1	Strong Association of De Novo Copy Number Mutations with Autism. <i>Science</i> , 2007, 316, 445-449.	12.6	2,497
2	Large-Scale Copy Number Polymorphism in the Human Genome. <i>Science</i> , 2004, 305, 525-528.	12.6	2,293
3	Mapping autism risk loci using genetic linkage and chromosomal rearrangements. <i>Nature Genetics</i> , 2007, 39, 319-328.	21.4	1,272
4	Recurrent 16p11.2 microdeletions in autism. <i>Human Molecular Genetics</i> , 2007, 17, 628-638.	2.9	614
5	Mutations in <i>LG11</i> cause autosomal-dominant partial epilepsy with auditory features. <i>Nature Genetics</i> , 2002, 30, 335-341.	21.4	555
6	Identification and Analysis of Mutations in the Wilson Disease Gene (<i>ATP7B</i>): Population Frequencies, Genotype-Phenotype Correlation, and Functional Analyses. <i>American Journal of Human Genetics</i> , 1997, 61, 317-328.	6.2	346
7	Differential <i>SMN2</i> expression associated with SMA severity. <i>Nature Genetics</i> , 1998, 20, 230-231.	21.4	304
8	A possible vulnerability locus for bipolar affective disorder on chromosome 21q22.3. <i>Nature Genetics</i> , 1994, 8, 291-296.	21.4	297
9	A Genomewide Screen for Autism Susceptibility Loci. <i>American Journal of Human Genetics</i> , 2001, 69, 327-340.	6.2	287
10	The neuronal ceroid lipofuscinoses in human <i>EPMR</i> and <i>mnd</i> mutant mice are associated with mutations in <i>CLN8</i> . <i>Nature Genetics</i> , 1999, 23, 233-236.	21.4	277
11	Novel Submicroscopic Chromosomal Abnormalities Detected in Autism Spectrum Disorder. <i>Biological Psychiatry</i> , 2008, 63, 1111-1117.	1.3	268
12	Evidence for a Language Quantitative Trait Locus on Chromosome 7q in Multiplex Autism Families. <i>American Journal of Human Genetics</i> , 2002, 70, 60-71.	6.2	253
13	A Genomewide Screen of 345 Families for Autism-Susceptibility Loci. <i>American Journal of Human Genetics</i> , 2003, 73, 886-897.	6.2	247
14	Catechol-O-Methyltransferase (<i>COMT</i>) Genotypes and Working Memory: Associations with Differing Cognitive Operations. <i>Biological Psychiatry</i> , 2005, 58, 901-907.	1.3	237
15	Linkage of Tunisian autosomal recessive Duchenneâ€“like muscular dystrophy to the pericentromeric region of chromosome 13q. <i>Nature Genetics</i> , 1992, 2, 315-317.	21.4	186
16	Null Mutation of the Murine <i>ATP7B</i> (Wilson Disease) Gene Results in Intracellular Copper Accumulation and Late-Onset Hepatic Nodular Transformation. <i>Human Molecular Genetics</i> , 1999, 8, 1665-1671.	2.9	186
17	Evidence for Sex-Specific Risk Alleles in Autism Spectrum Disorder. <i>American Journal of Human Genetics</i> , 2004, 75, 1117-1123.	6.2	165
18	Diminished support for linkage between manic depressive illness and Xâ€“chromosome markers in three Israeli pedigrees. <i>Nature Genetics</i> , 1993, 3, 49-55.	21.4	158

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19	TULP1 mutation in two extended Dominican kindreds with autosomal recessive Retinitis pigmentosa. <i>Nature Genetics</i> , 1998, 18, 177-179.	21.4	151
20	Genomewide Scan for Linkage Reveals Evidence of Several Susceptibility Loci for Alopecia Areata. <i>American Journal of Human Genetics</i> , 2007, 80, 316-328.	6.2	132
21	A Comprehensive Linkage Analysis of Chromosome 21q22 Supports Prior Evidence for a Putative Bipolar Affective Disorder Locus. <i>American Journal of Human Genetics</i> , 1999, 64, 210-217.	6.2	104
22	Genomewide Linkage Analysis of Celiac Disease in Finnish Families. <i>American Journal of Human Genetics</i> , 2002, 70, 51-59.	6.2	90
23	DRD2 C957T polymorphism interacts with the COMT Val158Met polymorphism in human working memory ability. <i>Schizophrenia Research</i> , 2007, 90, 104-107.	2.0	71
24	Population-based study of SR-BI genetic variation and lipid profile. <i>Atherosclerosis</i> , 2004, 175, 159-168.	0.8	69
25	Genetic-linkage mapping of complex hereditary disorders to a whole-genome molecular-interaction network. <i>Genome Research</i> , 2008, 18, 1150-1162.	5.5	63
26	Four New Families with Autosomal Dominant Partial Epilepsy with Auditory Features: Clinical Description and Linkage to Chromosome 10q24. <i>Epilepsia</i> , 2002, 43, 60-67.	5.1	59
27	A follow-up linkage study supports evidence for a bipolar affective disorder locus on chromosome 21q22. <i>American Journal of Medical Genetics Part A</i> , 2001, 105, 189-194.	2.4	43
28	No Evidence for Significant Linkage between Bipolar Affective Disorder and Chromosome 18 Pericentromeric Markers in a Large Series of Multiplex Extended Pedigrees. <i>American Journal of Human Genetics</i> , 1998, 62, 916-924.	6.2	36
29	Association of extreme blood lipid profile phenotypic variation with 11 reverse cholesterol transport genes and 10 non-genetic cardiovascular disease risk factors. <i>Human Molecular Genetics</i> , 2003, 12, 2733-2743.	2.9	34
30	High-Resolution Mapping and Transcript Identification at the Progressive Epilepsy with Mental Retardation Locus on Chromosome 8p. <i>Genome Research</i> , 1997, 7, 887-896.	5.5	26
31	Association between lipoprotein lipase (LPL) gene and blood lipids: A common variant for a common trait?. <i>Genetic Epidemiology</i> , 2003, 24, 309-321.	1.3	25
32	Extensive DNA deletion associated with severe disease alleles on spinal muscular atrophy homologues. <i>Annals of Neurology</i> , 1997, 42, 41-49.	5.3	17
33	Arm tremor secondary to Wilson's disease. <i>Movement Disorders</i> , 1998, 13, 351-353.	3.9	11
34	Genetic disorders of copper metabolism. <i>Current Opinion in Pediatrics</i> , 1994, 6, 698-701.	2.0	9
35	Is the Spinal Muscular Atrophy gene found?. <i>Nature Medicine</i> , 1995, 1, 124-127.	30.7	7