George Kirov

List of Publications by Year in Descending Order

184 papers	20,319 citations	62 h-index	142 g-index
200 ext. papers	27,190 ext. citations	10.4 avg, IF	5.82 L-index

#	Paper	IF	Citations
184	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. *Biological Psychiatry*, **2022**, 91, 102-117	7.9	11
183	Reduced reproductive success is associated with selective constraint on human genes.. *Nature*, **2022**, 603, 858-863	50.4	0
182	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. *Nature*, **2022**,	50.4	35
181	Rare coding variants in ten genes confer substantial risk for schizophrenia.. *Nature*, **2022**,	50.4	16
180	Comorbid Medical Issues in X-Linked Ichthyosis.. *JID Innovations*, **2022**, 2, 100109		0
179	Genetic association of FMRP targets with psychiatric disorders. *Molecular Psychiatry*, **2021**, 26, 2977-2990	15.1	7
178	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. *Nature Genetics*, **2021**, 53, 817-829	36.3	83
177	Electroconvulsive therapy (ECT) and schizophrenia. *BJPsych Open*, **2021**, 7, S332-S332	5	
176	Copy number variation and neuropsychiatric illness. *Current Opinion in Genetics and Development*, **2021**, 68, 57-63	4.9	8
175	Effects of genomic copy number variants penetrant for schizophrenia on cortical thickness and surface area in healthy individuals: analysis of the UK Biobank. *British Journal of Psychiatry*, **2021**, 218, 104-111	5.4	2
174	Electroconvulsive therapy for depression: 80 years of progress.. *British Journal of Psychiatry*, **2021**, 219, 594-597	5.4	7
173	Rare Copy Number Variants Are Associated With Poorer Cognition in Schizophrenia. *Biological Psychiatry*, **2021**, 90, 28-34	7.9	5
172	Characterisation of age and polarity at onset in bipolar disorder.. *British Journal of Psychiatry*, **2021**, 219, 659-669	5.4	2
171	Analysis of Diffusion Tensor Imaging Data From the UK Biobank Confirms Dosage Effect of 15q11.2 Copy Number Variation on White Matter and Shows Association With Cognition. *Biological Psychiatry*, **2021**, 90, 307-316	7.9	1
170	Schizophrenia, autism spectrum disorders and developmental disorders share specific disruptive coding mutations. *Nature Communications*, **2021**, 12, 5353	17.4	8
169	A brief report: de novo copy number variants in children with attention deficit hyperactivity disorder. *Translational Psychiatry*, **2020**, 10, 135	8.6	5
168	The mutational constraint spectrum quantified from variation in 141,456 humans. *Nature*, **2020**, 581, 434-443	50.4	2278

167	A structural variation reference for medical and population genetics. *Nature*, **2020**, 581, 444-451	50.4	223
166	Transcript expression-aware annotation improves rare variant interpretation. *Nature*, **2020**, 581, 452-458	50.4	55
165	Contribution of de novo and inherited rare CNVs to very preterm birth. *Journal of Medical Genetics*, **2020**, 57, 552-557	5.8	2
164	De novo mutations identified by exome sequencing implicate rare missense variants in SLC6A1 in schizophrenia. *Nature Neuroscience*, **2020**, 23, 179-184	25.5	47
163	Medical and neurobehavioural phenotypes in carriers of X-linked ichthyosis-associated genetic deletions in the UK Biobank. *Journal of Medical Genetics*, **2020**, 57, 692-698	5.8	11
162	Characterization of Single Gene Copy Number Variants in Schizophrenia. *Biological Psychiatry*, **2020**, 87, 736-744	7.9	8
161	International Consortium on the Genetics of Electroconvulsive Therapy and Severe Depressive Disorders (Gen-ECT-ic). *European Archives of Psychiatry and Clinical Neuroscience*, **2020**, 270, 921-932	5.1	8
160	Molecular pathology in neuropsychiatric disorders **2020**, 487-495		
159	Impact of schizophrenia genetic liability on the association between schizophrenia and physical illness: data-linkage study. *BJPsych Open*, **2020**, 6, e139	5	1
158	Medical and neurobehavioural phenotypes in male and female carriers of Xp22.31 duplications in the UK Biobank. *Human Molecular Genetics*, **2020**, 29, 2872-2881	5.6	4
157	Genetic liability to schizophrenia is negatively associated with educational attainment in UK Biobank. *Molecular Psychiatry*, **2020**, 25, 703-705	15.1	11
156	Schizophrenia-associated genomic copy number variants and subcortical brain volumes in the UK Biobank. *Molecular Psychiatry*, **2020**, 25, 854-862	15.1	30
155	Clinical indicators of treatment-resistant psychosis. *British Journal of Psychiatry*, **2020**, 216, 259-266	5.4	19
154	Association of Genetic Liability to Psychotic Experiences With Neuropsychotic Disorders and Traits. *JAMA Psychiatry*, **2019**, 76, 1256-1265	14.5	58
153	Association of Rare Copy Number Variants With Risk of Depression. *JAMA Psychiatry*, **2019**, 76, 818-825	14.5	50
152	Genome-wide association study identifies 30 loci associated with bipolar disorder. *Nature Genetics*, **2019**, 51, 793-803	36.3	662
151	De novo single-nucleotide and copy number variation in discordant monozygotic twins reveals disease-related genes. *European Journal of Human Genetics*, **2019**, 27, 1121-1133	5.3	24
150	Cognitive performance and functional outcomes of carriers of pathogenic copy number variants: analysis of the UK Biobank. *British Journal of Psychiatry*, **2019**, 214, 297-304	5.4	46

149	The Relationship Between Common Variant Schizophrenia Liability and Number of Offspring in the UK Biobank: Response to Lawn et al. *American Journal of Psychiatry*, **2019**, 176, 574-575	11.9	5
148	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. *Cell*, **2019**, 179, 1469-1482.e11	56.2	402
147	Targeted Sequencing of 10,198 Samples Confirms Abnormalities in Neuronal Activity and Implicates Voltage-Gated Sodium Channels in Schizophrenia Pathogenesis. *Biological Psychiatry*, **2019**, 85, 554-562	7.9	21
146	The Relationship Between Common Variant Schizophrenia Liability and Number of Offspring in the UK Biobank. *American Journal of Psychiatry*, **2019**, 176, 661-666	11.9	6
145	Medical consequences of pathogenic CNVs in adults: analysis of the UK Biobank. *Journal of Medical Genetics*, **2019**, 56, 131-138	5.8	56
144	Polygenic risk for schizophrenia and season of birth within the UK Biobank cohort. *Psychological Medicine*, **2019**, 49, 2499-2504	6.9	12
143	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. *Nature Genetics*, **2018**, 50, 381-389	36.3	787
142	Association of copy number variation across the genome with neuropsychiatric traits in the general population. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2018**, 177, 489-502	3.5	17
141	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. *Cell*, **2018**, 173, 1705-1715.e16	56.2	360
140	An InDel in Phospholipase-C-B-1 Is Linked with Euthyroid Multinodular Goiter. *Thyroid*, **2018**, 28, 891-901	6.2	6
139	Effects of pathogenic CNVs on physical traits in participants of the UK Biobank. *BMC Genomics*, **2018**, 19, 867	4.5	19
138	Genome-wide significant locus for Research Diagnostic Criteria Schizoaffective Disorder Bipolar type. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2017**, 174, 767-771	3.5	
137	Cognitive Performance Among Carriers of Pathogenic Copy Number Variants: Analysis of 152,000 UK Biobank Subjects. *Biological Psychiatry*, **2017**, 82, 103-110	7.9	91
136	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. *Nature Genetics*, **2017**, 49, 27-35	36.3	530
135	Analysis of Intellectual Disability Copy Number Variants for Association With Schizophrenia. *JAMA Psychiatry*, **2016**, 73, 963-969	14.5	75
134	Authors' reply. *British Journal of Psychiatry*, **2016**, 209, 84-5	5.4	
133	Pathogenic copy number variants and SCN1A mutations in patients with intellectual disability and childhood-onset epilepsy. *BMC Medical Genetics*, **2016**, 17, 34	2.1	16
132	Genome-wide association study reveals greater polygenic loading for schizophrenia in cases with a family history of illness. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2016**, 171B, 276-89	3.5	23

131	Mutation screening of SCN2A in schizophrenia and identification of a novel loss-of-function mutation. *Psychiatric Genetics*, **2016**, 26, 60-5	2.9	33
130	Parental Origin of Interstitial Duplications at 15q11.2-q13.3 in Schizophrenia and Neurodevelopmental Disorders. *PLoS Genetics*, **2016**, 12, e1005993	6	38
129	Gender differences in CNV burden do not confound schizophrenia CNV associations. *Scientific Reports*, **2016**, 6, 25986	4.9	6
128	A Method to Exploit the Structure of Genetic Ancestry Space to Enhance Case-Control Studies. *American Journal of Human Genetics*, **2016**, 98, 857-868	11	14
127	CNVs in neuropsychiatric disorders. *Human Molecular Genetics*, **2015**, 24, R45-9	5.6	101
126	What a psychiatrist needs to know about copy number variants. *BJ Psych Advances*, **2015**, 21, 157-163	0.8	11
125	Evidence of Mitochondrial Dysfunction within the Complex Genetic Etiology of Schizophrenia. *Molecular Neuropsychiatry*, **2015**, 1, 201-19	4.9	50
124	No Evidence for Enrichment in Schizophrenia for Common Allelic Associations at Imprinted Loci. *PLoS ONE*, **2015**, 10, e0144172	3.7	3
123	Novel Findings from CNVs Implicate Inhibitory and Excitatory Signaling Complexes in Schizophrenia. *Neuron*, **2015**, 86, 1203-14	13.9	119
122	De novo mutations in schizophrenia implicate synaptic networks. *Nature*, **2014**, 506, 179-84	50.4	1163
121	Sequence analysis of 17 NRXN1 deletions. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2014**, 165B, 52-61	3.5	10
120	Authors' reply. *British Journal of Psychiatry*, **2014**, 205, 78	5.4	1
119	SGCE and myoclonus dystonia: motor characteristics, diagnostic criteria and clinical predictors of genotype. *Journal of Neurology*, **2014**, 261, 2296-304	5.5	42
118	Analysis of copy number variations at 15 schizophrenia-associated loci. *British Journal of Psychiatry*, **2014**, 204, 108-14	5.4	283
117	An inherited duplication at the gene p21 Protein-Activated Kinase 7 (PAK7) is a risk factor for psychosis. *Human Molecular Genetics*, **2014**, 23, 3316-26	5.6	32
116	CNV analysis in a large schizophrenia sample implicates deletions at 16p12.1 and SLC1A1 and duplications at 1p36.33 and CGNL1. *Human Molecular Genetics*, **2014**, 23, 1669-76	5.6	61
115	De novo CNVs in bipolar affective disorder and schizophrenia. *Human Molecular Genetics*, **2014**, 23, 6677-83	5.6	57
114	Reciprocal duplication of the Williams-Beuren syndrome deletion on chromosome 7q11.23 is associated with schizophrenia. *Biological Psychiatry*, **2014**, 75, 371-7	7.9	54

113	The penetrance of copy number variations for schizophrenia and developmental delay. *Biological Psychiatry*, **2014**, 75, 378-85	7.9	236
112	Rare exonic deletions implicate the synaptic organizer Gephyrin (GPHN) in risk for autism, schizophrenia and seizures. *Human Molecular Genetics*, **2013**, 22, 2055-66	5.6	110
111	Implication of a rare deletion at distal 16p11.2 in schizophrenia. *JAMA Psychiatry*, **2013**, 70, 253-60	14.5	56
110	Mosaic copy number variation in schizophrenia. *European Journal of Human Genetics*, **2013**, 21, 1007-11	5.3	10
109	Schizophrenia two-hit hypothesis in velo-cardio facial syndrome. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2013**, 162B, 177-82	3.5	19
108	Non-random mating, parent-of-origin, and maternal-fetal incompatibility effects in schizophrenia. *Schizophrenia Research*, **2013**, 143, 11-7	3.6	1
107	A comprehensive family-based replication study of schizophrenia genes. *JAMA Psychiatry*, **2013**, 70, 573-81	14.5	115
106	SGCE mutations cause psychiatric disorders: clinical and genetic characterization. *Brain*, **2013**, 136, 294-303	10.2	72
105	Whole-genome-wide association study in the Bulgarian population reveals HHAT as schizophrenia susceptibility gene. *Psychiatric Genetics*, **2013**, 23, 11-9	2.9	12
104	Reduced burden of very large and rare CNVs in bipolar affective disorder. *Bipolar Disorders*, **2013**, 15, 893-8	3.8	22
103	A genome-wide study shows a limited contribution of rare copy number variants to Alzheimer's disease risk. *Human Molecular Genetics*, **2013**, 22, 816-24	5.6	26
102	De novo CNV analysis implicates specific abnormalities of postsynaptic signalling complexes in the pathogenesis of schizophrenia. *Molecular Psychiatry*, **2012**, 17, 142-53	15.1	611
101	Discovery and statistical genotyping of copy-number variation from whole-exome sequencing depth. *American Journal of Human Genetics*, **2012**, 91, 597-607	11	391
100	Independent estimation of the frequency of rare CNVs in the UK population confirms their role in schizophrenia. *Schizophrenia Research*, **2012**, 135, 1-7	3.6	62
99	Absence of de novo point mutations in exons of GRIN2B in a large schizophrenia trio sample. *Schizophrenia Research*, **2012**, 141, 274-6	3.6	4
98	De novo mutation in schizophrenia. *Schizophrenia Bulletin*, **2012**, 38, 377-81	1.3	40
97	Genome-wide pooling approach identifies SPATA5 as a new susceptibility locus for alopecia areata. *European Journal of Human Genetics*, **2012**, 20, 326-32	5.3	37
96	Response to Boot et al. Letter. *American Journal of Psychiatry*, **2012**, 169, 97-97	11.9	4

95	Creativity and mental disorder. *British Journal of Psychiatry*, **2012**, 200, 347; author reply 348	5.4	
94	De novo rates and selection of schizophrenia-associated copy number variants. *Biological Psychiatry*, **2011**, 70, 1109-14	7.9	69
93	Analysis of neurogranin (NRGN) in schizophrenia. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2011**, 156B, 532-5	3.5	10
92	Maternally derived microduplications at 15q11-q13: implication of imprinted genes in psychotic illness. *American Journal of Psychiatry*, **2011**, 168, 408-17	11.9	84
91	Population structure and genome-wide patterns of variation in Ireland and Britain. *European Journal of Human Genetics*, **2010**, 18, 1248-54	5.3	36
90	Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. *Nature*, **2010**, 464, 713-20	50.4	639
89	Genetic differences between five European populations. *Human Heredity*, **2010**, 70, 141-9	1.1	24
88	Copy number variation in schizophrenia in the Japanese population. *Biological Psychiatry*, **2010**, 67, 283-6	7.9	95
87	The role of copy number variation in schizophrenia. *Expert Review of Neurotherapeutics*, **2010**, 10, 25-32	4.3	50
86	Rare copy number variants: a point of rarity in genetic risk for bipolar disorder and schizophrenia. *Archives of General Psychiatry*, **2010**, 67, 318-27		154
85	Effects of chronic lithium treatment on renal function. *Journal of Affective Disorders*, **2010**, 126, 436-40	6.6	67
84	Fine-mapping reveals novel alternative splicing of the dopamine transporter. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2010**, 153B, 1434-47	3.5	16
83	An international collaborative family-based whole-genome linkage scan for high-grade myopia **2009**, 50, 3116-27		60
82	Whole genome association study in a homogenous population in Shandong peninsula of China reveals JARID2 as a susceptibility gene for schizophrenia. *Journal of Biomedicine and Biotechnology*, **2009**, 2009, 536918		26
81	Case-control association study of 59 candidate genes reveals the DRD2 SNP rs6277 (C957T) as the only susceptibility factor for schizophrenia in the Bulgarian population. *Journal of Human Genetics*, **2009**, 54, 98-107	4.3	88
80	Neurexin 1 (NRXN1) deletions in schizophrenia. *Schizophrenia Bulletin*, **2009**, 35, 851-4	1.3	183
79	Support for the involvement of large copy number variants in the pathogenesis of schizophrenia. *Human Molecular Genetics*, **2009**, 18, 1497-503	5.6	346
78	Age-at-onset in bipolar-I disorder: mixture analysis of 1369 cases identifies three distinct clinical sub-groups. *Journal of Affective Disorders*, **2009**, 116, 23-9	6.6	65

77	Case-control association study of 65 candidate genes revealed a possible association of a SNP of HTR5A to be a factor susceptible to bipolar disease in Bulgarian population. *Journal of Affective Disorders*, **2009**, 117, 87-97	6.6	32
76	Convergent patterns of association between phenylalanine hydroxylase variants and schizophrenia in four independent samples. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics* , **2009**, 150B, 560-9	3.5	12
75	P2RX7: A bipolar and unipolar disorder candidate susceptibility gene?. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2009**, 150B, 1063-9	3.5	47
74	A genome-wide association study in 574 schizophrenia trios using DNA pooling. *Molecular Psychiatry*, **2009**, 14, 796-803	15.1	117
73	Microduplications of 16p11.2 are associated with schizophrenia. *Nature Genetics*, **2009**, 41, 1223-7	36.3	550
72	Mood-incongruent psychosis in bipolar disorder: conditional linkage analysis shows genome-wide suggestive linkage at 1q32.3, 7p13 and 20q13.31. *Bipolar Disorders*, **2009**, 11, 610-20	3.8	20
71	Four years of successful maintenance electroconvulsive therapy. *Journal of ECT*, **2009**, 25, 219-20	2	9
70	Identification of loci associated with schizophrenia by genome-wide association and follow-up. *Nature Genetics*, **2008**, 40, 1053-5	36.3	877
69	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. *Nature Genetics*, **2008**, 40, 1056-8	36.3	949
68	Support for neuregulin 1 as a susceptibility gene for bipolar disorder and schizophrenia. *Biological Psychiatry*, **2008**, 64, 419-27	7.9	95
67	Strong evidence that GNB1L is associated with schizophrenia. *Human Molecular Genetics*, **2008**, 17, 555-566	5.6	58
66	Quick recovery of orientation after magnetic seizure therapy for major depressive disorder. *British Journal of Psychiatry*, **2008**, 193, 152-5	5.4	70
65	The effectiveness of electroconvulsive therapy in treatment-resistant depression: a naturalistic study. *Journal of ECT*, **2008**, 24, 141-5	2	30
64	Association study in the 5q31-32 linkage region for schizophrenia using pooled DNA genotyping. *BMC Psychiatry*, **2008**, 8, 11	4.2	24
63	A genome-wide association study for late-onset Alzheimer's disease using DNA pooling. *BMC Medical Genomics*, **2008**, 1, 44	3.7	133
62	A network of dopaminergic gene variations implicated as risk factors for schizophrenia. *Human Molecular Genetics*, **2008**, 17, 747-58	5.6	120
61	Comparative genome hybridization suggests a role for NRXN1 and APBA2 in schizophrenia. *Human Molecular Genetics*, **2008**, 17, 458-65	5.6	315
60	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. *Nature Genetics*, **2007**, 39, 1329-37	36.3	1130

59	Relative suppression of magical thinking: a transcranial magnetic stimulation study. *Cortex*, **2007**, 43, 551-7	3.8	14
58	Pooled DNA genotyping on Affymetrix SNP genotyping arrays. *BMC Genomics*, **2006**, 7, 27	4.5	69
57	Evidence that interaction between neuregulin 1 and its receptor erbB4 increases susceptibility to schizophrenia. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2006**, 141B, 96-101	3.5	143
56	Convergent evidence that oligodendrocyte lineage transcription factor 2 (OLIG2) and interacting genes influence susceptibility to schizophrenia. *Proceedings of the National Academy of Sciences of the United States of America*, **2006**, 103, 12469-74	11.5	101
55	Evaluation of a susceptibility gene for schizophrenia: genotype based meta-analysis of RGS4 polymorphisms from thirteen independent samples. *Biological Psychiatry*, **2006**, 60, 152-62	7.9	80
54	Analysis of ProDH, COMT and ZDHHC8 risk variants does not support individual or interactive effects on schizophrenia susceptibility. *Schizophrenia Research*, **2006**, 87, 21-7	3.6	36
53	Genetic variation of brain-derived neurotrophic factor (BDNF) in bipolar disorder: case-control study of over 3000 individuals from the UK. *British Journal of Psychiatry*, **2006**, 188, 21-5	5.4	119
52	The effects of etomidate on seizure duration and electrical stimulus dose in seizure-resistant patients during electroconvulsive therapy. *Journal of ECT*, **2006**, 22, 184-8	2	22
51	Variation at the DAOA/G30 locus influences susceptibility to major mood episodes but not psychosis in schizophrenia and bipolar disorder. *Archives of General Psychiatry*, **2006**, 63, 366-73		124
50	Bipolar disorder and polymorphisms in the dysbindin gene (DTNBP1). *Biological Psychiatry*, **2005**, 57, 696-701	7.9	110
49	No association between the putative functional ZDHHC8 single nucleotide polymorphism rs175174 and schizophrenia in large European samples. *Biological Psychiatry*, **2005**, 58, 78-80	7.9	38
48	Operation of the schizophrenia susceptibility gene, neuregulin 1, across traditional diagnostic boundaries to increase risk for bipolar disorder. *Archives of General Psychiatry*, **2005**, 62, 642-8		209
47	A cross-sectional and a prospective study of thyroid disorders in lithium-treated patients. *Journal of Affective Disorders*, **2005**, 87, 313-7	6.6	56
46	Add-on topiramate reduces weight in overweight patients with affective disorders: a clinical case series. *BMC Psychiatry*, **2005**, 5, 19	4.2	17
45	Linkage disequilibrium mapping of bipolar affective disorder at 12q23-q24 provides evidence for association at CUX2 and FLJ32356. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2005**, 132B, 38-45	3.5	19
44	Genomewide linkage scan in schizoaffective disorder: significant evidence for linkage at 1q42 close to DISC1, and suggestive evidence at 22q11 and 19p13. *Archives of General Psychiatry*, **2005**, 62, 1081-8		164
43	Localization of bipolar susceptibility locus by molecular genetic analysis of the chromosome 12q23-q24 region in two pedigrees with bipolar disorder and Darier's disease. *American Journal of Psychiatry*, **2005**, 162, 35-42	11.9	50
42	Finding schizophrenia genes. *Journal of Clinical Investigation*, **2005**, 115, 1440-8	15.9	91

41	Family aggregation of high myopia: estimation of the sibling recurrence risk ratio. *Investigative Ophthalmology and Visual Science*, **2004**, 45, 2873-8		77
40	Linkage analysis of the genetic loci for high myopia on 18p, 12q, and 17q in 51 U.K. families. *Investigative Ophthalmology and Visual Science*, **2004**, 45, 2879-85		64
39	Lack of support for a genetic association of the XBP1 promoter polymorphism with bipolar disorder in probands of European origin. *Nature Genetics*, **2004**, 36, 783-4; author reply 784-5	36.3	54
38	The Bipolar Affective Disorder Dimension Scale (BADDS)--a dimensional scale for rating lifetime psychopathology in bipolar spectrum disorders. *BMC Psychiatry*, **2004**, 4, 19	4.2	59
37	Support for RGS4 as a susceptibility gene for schizophrenia. *Biological Psychiatry*, **2004**, 55, 192-5	7.9	125
36	Strong evidence for association between the dystrobrevin binding protein 1 gene (DTNBP1) and schizophrenia in 488 parent-offspring trios from Bulgaria. *Biological Psychiatry*, **2004**, 55, 971-5	7.9	133
35	. *Psychiatric Genetics*, **2003**, 13, 103-106	2.9	
34	Association analysis of the HOPA12bp polymorphism in schizophrenia and manic depressive illness. *American Journal of Medical Genetics Part A*, **2003**, 118B, 16-9		8
33	Variation in the protocadherin gamma A gene cluster. *Genomics*, **2003**, 82, 433-40	4.3	12
32	Schizophrenia and functional polymorphisms in the MAOA and COMT genes: no evidence for association or epistasis. *American Journal of Medical Genetics Part A*, **2002**, 114, 491-6		64
31	Universal, robust, highly quantitative SNP allele frequency measurement in DNA pools. *Human Genetics*, **2002**, 110, 471-8	6.3	152
30	Major psychiatric disorders and the serotonin transporter gene (SLC6A4): family-based association studies. *Psychiatric Genetics*, **2002**, 12, 137-41	2.9	10
29	Association analysis of NOTCH4 loci in schizophrenia using family and population-based controls. *Nature Genetics*, **2001**, 28, 126-8	36.3	57
28	Screening ABCG1, the human homologue of the Drosophila white gene, for polymorphisms and association with bipolar affective disorder. *Molecular Psychiatry*, **2001**, 6, 671-7	15.1	18
27	No evidence of association from transmission disequilibrium analysis of the hKCa3 gene in bipolar disorder. *Bipolar Disorders*, **2000**, 2, 328-31	3.8	2
26	Cheap, accurate and rapid allele frequency estimation of single nucleotide polymorphisms by primer extension and DHPLC in DNA pools. *Human Genetics*, **2000**, 107, 488-93	6.3	152
25	Tryptophan hydroxylase gene and manic-depressive illness. *Archives of General Psychiatry*, **1999**, 56, 98-9		29
24	Association analysis between dopamine receptor genes and bipolar affective disorder. *Psychiatry Research*, **1999**, 86, 193-201	9.9	45

23	A functional polymorphism in the promoter of monoamine oxidase A gene and bipolar affective disorder. *International Journal of Neuropsychopharmacology*, **1999**, 2, 293-298	5.8	17
22	Thyroid disorders in lithium-treated patients. *Journal of Affective Disorders*, **1998**, 50, 33-40	6.6	42
21	Apolipoprotein E: depressive illness, depressive symptoms, and Alzheimer's disease. *Biological Psychiatry*, **1998**, 43, 159-64	7.9	39
20	Randomised controlled trial of compliance therapy. 18-month follow-up. *British Journal of Psychiatry*, **1998**, 172, 413-9	5.4	412
19	Religious faith after psychotic illness. *Psychopathology*, **1998**, 31, 234-45	3.4	74
18	No evidence for an association of affective disorders with high- or low-activity allele of catechol-o-methyltransferase gene. *Biological Psychiatry*, **1997**, 42, 282-5	7.9	97
17	5-HT2A receptor and bipolar affective disorder: association studies in affected patients. *Neuroscience Letters*, **1997**, 224, 95-8	3.3	52
16	No association between bipolar disorder and alleles at a functional polymorphism in the COMT gene. Biomed European Bipolar Collaborative Group. *British Journal of Psychiatry*, **1997**, 170, 526-8	5.4	46
15	The molecular genetics of schizophrenia: progress so far. *Trends in Molecular Medicine*, **1997**, 3, 124-30		10
14	Do obstetric complications cause the earlier age at onset in male than female schizophrenics?. *Schizophrenia Research*, **1996**, 20, 117-24	3.6	45
13	The serotonin transporter is a potential susceptibility factor for bipolar affective disorder. *NeuroReport*, **1996**, 7, 1675-9	1.7	170
12	Characterization of single gene copy number variants in schizophrenia		1
11	Sex-biased reduction in reproductive success drives selective constraint on human genes		1
10	A brief report: de novo copy number variants in children with attention deficit hyperactivity disorder		1
9	Common schizophrenia alleles are enriched in mutation-intolerant genes and maintained by background selection		20
8	Genome-wide association study identifies 30 Loci Associated with Bipolar Disorder		28
7	Schizophrenia, autism spectrum disorders and developmental disorders share specific disruptive coding mutations		1
6	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology		11

5	Association between schizophrenia and both loss of function and missense mutations in paralog conserved sites of voltage-gated sodium channels	2
4	The role of rare copy number variants in depression	1
3	Analyses of rare and common alleles in parent-proband trios implicate rare missense variants in SLC6A1 in schizophrenia and confirm the involvement of loss of function intolerant and neurodevelopmental disorder genes	2
2	Effects of pathogenic CNVs on biochemical markers: a study on the UK Biobank	2
1	The rate of de novo CNVs in healthy controls	2