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List of Publications by Citations

815 papers	120,517 citations	146 h-index	332 g-index
946 ext. papers	142,149 ext. citations	10.2 avg, IF	8.92 L-index

#	Paper	IF	Citations
815	Genome-wide association study of 14,000 cases of seven common diseases and 3,000 shared controls. *Nature*, **2007**, 447, 661-78	50.4	7801
814	A novel gene containing a trinucleotide repeat that is expanded and unstable on Huntington's disease chromosomes. The Huntington's Disease Collaborative Research Group. *Cell*, **1993**, 72, 971-83	56.2	6854
813	Biological insights from 108 schizophrenia-associated genetic loci. *Nature*, **2014**, 511, 421-7	50.4	5249
812	Common polygenic variation contributes to risk of schizophrenia and bipolar disorder. *Nature*, **2009**, 460, 748-52	50.4	3568
811	Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. *Nature*, **2012**, 491, 119-24	50.4	3239
810	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. *Nature Genetics*, **2013**, 45, 1452-8	36.3	2714
809	Identification of risk loci with shared effects on five major psychiatric disorders: a genome-wide analysis. *Lancet, The*, **2013**, 381, 1371-1379	40	2112
808	LD Score regression distinguishes confounding from polygenicity in genome-wide association studies. *Nature Genetics*, **2015**, 47, 291-5	36.3	2096
807	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. *Nature Genetics*, **2009**, 41, 1088-93	36.3	2018
806	Replication of genome-wide association signals in UK samples reveals risk loci for type 2 diabetes. *Science*, **2007**, 316, 1336-41	33.3	1823
805	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. *Nature Genetics*, **2013**, 45, 984-94	36.3	1628
804	Synaptic, transcriptional and chromatin genes disrupted in autism. *Nature*, **2014**, 515, 209-15	50.4	1581
803	Genome-wide association study identifies five new schizophrenia loci. *Nature Genetics*, **2011**, 43, 969-76	36.3	1508
802	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. *Nature Genetics*, **2011**, 43, 429-35	36.3	1421
801	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. *Nature Genetics*, **2018**, 50, 668-681	36.3	1301
800	Rare chromosomal deletions and duplications increase risk of schizophrenia. *Nature*, **2008**, 455, 237-41	50.4	1251
799	De novo mutations in schizophrenia implicate synaptic networks. *Nature*, **2014**, 506, 179-84	50.4	1163

798	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. *Nature Genetics*, **2013**, 45, 1150-9	36.3	1153
797	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. *Nature Genetics*, **2007**, 39, 1329-37	36.3	1130
796	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. *Nature Genetics*, **2011**, 43, 977-83	36.3	1094
795	Genome scan meta-analysis of schizophrenia and bipolar disorder, part II: Schizophrenia. *American Journal of Human Genetics*, **2003**, 73, 34-48	11	985
794	Genome-wide association study identifies eight loci associated with blood pressure. *Nature Genetics*, **2009**, 41, 666-76	36.3	970
793	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. *Nature Genetics*, **2008**, 40, 1056-8	36.3	949
792	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A, tau, immunity and lipid processing. *Nature Genetics*, **2019**, 51, 414-430	36.3	917
791	Genome-wide analysis of genetic loci associated with Alzheimer disease. *JAMA - Journal of the American Medical Association*, **2010**, 303, 1832-40	27.4	888
790	Identification of loci associated with schizophrenia by genome-wide association and follow-up. *Nature Genetics*, **2008**, 40, 1053-5	36.3	877
789	Genetic architectures of psychiatric disorders: the emerging picture and its implications. *Nature Reviews Genetics*, **2012**, 13, 537-51	30.1	866
788	A mega-analysis of genome-wide association studies for major depressive disorder. *Molecular Psychiatry*, **2013**, 18, 497-511	15.1	853
787	Schizophrenia. *Lancet, The*, **2016**, 388, 86-97	40	836
786	High rates of schizophrenia in adults with velo-cardio-facial syndrome. *Archives of General Psychiatry*, **1999**, 56, 940-5		813
785	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
784	The UK10K project identifies rare variants in health and disease. *Nature*, **2015**, 526, 82-90	50.4	776
783	Analysis of shared heritability in common disorders of the brain. *Science*, **2018**, 360,	33.3	666
782	Genome-wide association study identifies 30 loci associated with bipolar disorder. *Nature Genetics*, **2019**, 51, 793-803	36.3	662
781	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. *American Journal of Human Genetics*, **2015**, 97, 576-92	11	649

780	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
779	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
778	Mutations in ATP2A2, encoding a Ca2+ pump, cause Darier disease. *Nature Genetics*, **1999**, 21, 271-7	36.3	601
777	Microduplications of 16p11.2 are associated with schizophrenia. *Nature Genetics*, **2009**, 41, 1223-7	36.3	550
776	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
775	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. *Nature Genetics*, **2017**, 49, 1373-1384	36.3	508
774	Genes for schizophrenia? Recent findings and their pathophysiological implications. *Lancet, The*, **2003**, 361, 417-9	40	488
773	DNA Pooling: a tool for large-scale association studies. *Nature Reviews Genetics*, **2002**, 3, 862-71	30.1	468
772	Meta-analysis shows significant association between dopamine system genes and attention deficit hyperactivity disorder (ADHD). *Human Molecular Genetics*, **2006**, 15, 2276-84	5.6	437
771	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. *Nature Genetics*, **2012**, 44, 1336-40	36.3	436
770	Schizophrenia. *Nature Reviews Disease Primers*, **2015**, 1, 15067	51.1	432
769	Localization of type 1 diabetes susceptibility to the MHC class I genes HLA-B and HLA-A. *Nature*, **2007**, 450, 887-92	50.4	421
768	Partitioning heritability of regulatory and cell-type-specific variants across 11 common diseases. *American Journal of Human Genetics*, **2014**, 95, 535-52	11	411
767	The Kraepelinian dichotomy - going, going... but still not gone. *British Journal of Psychiatry*, **2010**, 196, 92-5	5.4	411
766	The genetics of schizophrenia and bipolar disorder: dissecting psychosis. *Journal of Medical Genetics* , **2005**, 42, 193-204	5.8	406
765	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. *Cell*, **2019**, 179, 1469-1482.e11	56.2	402
764	Rare chromosomal deletions and duplications in attention-deficit hyperactivity disorder: a genome-wide analysis. *Lancet, The*, **2010**, 376, 1401-8	40	399
763	Discovery and statistical genotyping of copy-number variation from whole-exome sequencing depth. *American Journal of Human Genetics*, **2012**, 91, 597-607	11	391

762	The bipolar disorder risk allele at CACNA1C also confers risk of recurrent major depression and of schizophrenia. *Molecular Psychiatry*, **2010**, 15, 1016-22	15.1	389
761	Genes for schizophrenia and bipolar disorder? Implications for psychiatric nosology. *Schizophrenia Bulletin*, **2006**, 32, 9-16	1.3	385
760	Candidate single-nucleotide polymorphisms from a genomewide association study of Alzheimer disease. *Archives of Neurology*, **2008**, 65, 45-53		362
759	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. *Cell*, **2018**, 173, 1705-1715.e16	56.2	360
758	Meta-analysis of genome-wide association studies of attention-deficit/hyperactivity disorder. *Journal of the American Academy of Child and Adolescent Psychiatry*, **2010**, 49, 884-97	7.2	357
757	Genomewide association studies: history, rationale, and prospects for psychiatric disorders. *American Journal of Psychiatry*, **2009**, 166, 540-56	11.9	355
756	Support for the involvement of large copy number variants in the pathogenesis of schizophrenia. *Human Molecular Genetics*, **2009**, 18, 1497-503	5.6	346
755	Susceptibility locus for Alzheimer's disease on chromosome 10. *Science*, **2000**, 290, 2304-5	33.3	345
754	Neurodevelopmental hypothesis of schizophrenia. *British Journal of Psychiatry*, **2011**, 198, 173-5	5.4	338
753	Identifying relationships among genomic disease regions: predicting genes at pathogenic SNP associations and rare deletions. *PLoS Genetics*, **2009**, 5, e1000534	6	337
752	Gene ontology analysis of GWA study data sets provides insights into the biology of bipolar disorder. *American Journal of Human Genetics*, **2009**, 85, 13-24	11	333
751	Psychiatric Genomics: An Update and an Agenda. *American Journal of Psychiatry*, **2018**, 175, 15-27	11.9	328
750	Rheumatoid arthritis association at 6q23. *Nature Genetics*, **2007**, 39, 1431-3	36.3	328
749	Definition and description of schizophrenia in the DSM-5. *Schizophrenia Research*, **2013**, 150, 3-10	3.6	323
748	Comparative genome hybridization suggests a role for NRXN1 and APBA2 in schizophrenia. *Human Molecular Genetics*, **2008**, 17, 458-65	5.6	315
747	Convergent functional genomics of schizophrenia: from comprehensive understanding to genetic risk prediction. *Molecular Psychiatry*, **2012**, 17, 887-905	15.1	308
746	Blind analysis of denaturing high-performance liquid chromatography as a tool for mutation detection. *Genomics*, **1998**, 52, 44-9	4.3	303
745	A haplotype implicated in schizophrenia susceptibility is associated with reduced COMT expression in human brain. *American Journal of Human Genetics*, **2003**, 73, 152-61	11	301

744	Contrasting genetic architectures of schizophrenia and other complex diseases using fast variance-components analysis. *Nature Genetics*, **2015**, 47, 1385-92	36.3	299
743	Gene-wide analyses of genome-wide association data sets: evidence for multiple common risk alleles for schizophrenia and bipolar disorder and for overlap in genetic risk. *Molecular Psychiatry*, **2009**, 14, 252-60	15.1	296
742	Genetic identification of brain cell types underlying schizophrenia. *Nature Genetics*, **2018**, 50, 825-833	36.3	295
741	Improved detection of common variants associated with schizophrenia by leveraging pleiotropy with cardiovascular-disease risk factors. *American Journal of Human Genetics*, **2013**, 92, 197-209	11	293
740	The Role of Genes, Stress, and Dopamine in the Development of Schizophrenia. *Biological Psychiatry*, **2017**, 81, 9-20	7.9	289
739	Rare loss-of-function variants in SETD1A are associated with schizophrenia and developmental disorders. *Nature Neuroscience*, **2016**, 19, 571-7	25.5	284
738	Altering the course of schizophrenia: progress and perspectives. *Nature Reviews Drug Discovery*, **2016**, 15, 485-515	64.1	284
737	Analysis of copy number variations at 15 schizophrenia-associated loci. *British Journal of Psychiatry*, **2014**, 204, 108-14	5.4	283
736	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. *Science*, **2018**, 362,	33.3	277
735	Genetic evidence implicates the immune system and cholesterol metabolism in the aetiology of Alzheimer's disease. *PLoS ONE*, **2010**, 5, e13950	3.7	276
734	Genome-wide association analysis identifies 30 new susceptibility loci for schizophrenia. *Nature Genetics*, **2017**, 49, 1576-1583	36.3	272
733	Genome-wide association identifies a common variant in the reelin gene that increases the risk of schizophrenia only in women. *PLoS Genetics*, **2008**, 4, e28	6	270
732	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. *Molecular Psychiatry*, **2014**, 19, 1017-1024	15.1	258
731	The molecular genetics of schizophrenia: new findings promise new insights. *Molecular Psychiatry*, **2004**, 9, 14-27	15.1	257
730	Optimal Temperature Selection for Mutation Detection by Denaturing HPLC and Comparison to Single-Stranded Conformation Polymorphism and Heteroduplex Analysis. *Clinical Chemistry*, **1999**, 45, 1133-1140	5.5	252
729	Genome-wide analyses for personality traits identify six genomic loci and show correlations with psychiatric disorders. *Nature Genetics*, **2017**, 49, 152-156	36.3	251
728	Increased expression of BIN1 mediates Alzheimer genetic risk by modulating tau pathology. *Molecular Psychiatry*, **2013**, 18, 1225-34	15.1	251
727	Tbx1 haploinsufficiency is linked to behavioral disorders in mice and humans: implications for 22q11 deletion syndrome. *Proceedings of the National Academy of Sciences of the United States of America*, **2006**, 103, 7729-34	11.5	249

726	Schizophrenia: genes at last?. *Trends in Genetics*, **2005**, 21, 518-25	8.5	241
725	Predictors of developmental dyslexia in European orthographies with varying complexity. *Journal of Child Psychology and Psychiatry and Allied Disciplines*, **2013**, 54, 686-94	7.9	238
724	The penetrance of copy number variations for schizophrenia and developmental delay. *Biological Psychiatry*, **2014**, 75, 378-85	7.9	236
723	Variation in DCP1, encoding ACE, is associated with susceptibility to Alzheimer disease. *Nature Genetics*, **1999**, 21, 71-2	36.3	236
722	Strong evidence that KIAA0319 on chromosome 6p is a susceptibility gene for developmental dyslexia. *American Journal of Human Genetics*, **2005**, 76, 581-91	11	232
721	Methylation QTLs in the developing brain and their enrichment in schizophrenia risk loci. *Nature Neuroscience*, **2016**, 19, 48-54	25.5	227
720	Cognitive mechanisms underlying reading and spelling development in five European orthographies. *Learning and Instruction*, **2014**, 29, 65-77	5.8	223
719	Transcriptome-wide association study of schizophrenia and chromatin activity yields mechanistic disease insights. *Nature Genetics*, **2018**, 50, 538-548	36.3	222
718	Fine mapping of ZNF804A and genome-wide significant evidence for its involvement in schizophrenia and bipolar disorder. *Molecular Psychiatry*, **2011**, 16, 429-41	15.1	221
717	Genome-wide association study identifies genetic variation in neurocan as a susceptibility factor for bipolar disorder. *American Journal of Human Genetics*, **2011**, 88, 372-81	11	221
716	Evidence for novel susceptibility genes for late-onset Alzheimer's disease from a genome-wide association study of putative functional variants. *Human Molecular Genetics*, **2007**, 16, 865-73	5.6	221
715	Evaluating historical candidate genes for schizophrenia. *Molecular Psychiatry*, **2015**, 20, 555-62	15.1	219
714	The Machado-Joseph disease-associated mutant form of ataxin-3 regulates parkin ubiquitination and stability. *Human Molecular Genetics*, **2011**, 20, 141-54	5.6	217
713	The genetic deconstruction of psychosis. *Schizophrenia Bulletin*, **2007**, 33, 905-11	1.3	215
712	All SNPs are not created equal: genome-wide association studies reveal a consistent pattern of enrichment among functionally annotated SNPs. *PLoS Genetics*, **2013**, 9, e1003449	6	209
711	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
710	Support for genetic variation in neuregulin 1 and susceptibility to schizophrenia. *Molecular Psychiatry*, **2003**, 8, 485-7	15.1	206
709	Meta-analysis of 32 genome-wide linkage studies of schizophrenia. *Molecular Psychiatry*, **2009**, 14, 774-85	15.1	202

708	Expanded CAG repeats in schizophrenia and bipolar disorder. *Nature Genetics*, **1995**, 10, 380-1	36.3	199
707	Cis-acting variation in the expression of a high proportion of genes in human brain. *Human Genetics*, **2003**, 113, 149-53	6.3	198
706	Genetic overlap between autism, schizophrenia and bipolar disorder. *Genome Medicine*, **2009**, 1, 102	14.4	197
705	Genome-wide analysis of copy number variants in attention deficit hyperactivity disorder: the role of rare variants and duplications at 15q13.3. *American Journal of Psychiatry*, **2012**, 169, 195-204	11.9	195
704	Genetic risk for schizophrenia: convergence on synaptic pathways involved in plasticity. *Biological Psychiatry*, **2015**, 77, 52-8	7.9	192
703	Improved detection of common variants associated with schizophrenia and bipolar disorder using pleiotropy-informed conditional false discovery rate. *PLoS Genetics*, **2013**, 9, e1003455	6	192
702	Copy number variation in schizophrenia in Sweden. *Molecular Psychiatry*, **2014**, 19, 762-73	15.1	191
701	Methylomic trajectories across human fetal brain development. *Genome Research*, **2015**, 25, 338-52	9.7	188
700	Improved imputation of low-frequency and rare variants using the UK10K haplotype reference panel. *Nature Communications*, **2015**, 6, 8111	17.4	186
699	Identifying gene-environment interactions in schizophrenia: contemporary challenges for integrated, large-scale investigations. *Schizophrenia Bulletin*, **2014**, 40, 729-36	1.3	186
698	Comparative genetic architectures of schizophrenia in East Asian and European populations. *Nature Genetics*, **2019**, 51, 1670-1678	36.3	185
697	Genome-wide significant associations in schizophrenia to ITIH3/4, CACNA1C and SDCCAG8, and extensive replication of associations reported by the Schizophrenia PGC. *Molecular Psychiatry*, **2013**, 18, 708-12	15.1	184
696	Neurexin 1 (NRXN1) deletions in schizophrenia. *Schizophrenia Bulletin*, **2009**, 35, 851-4	1.3	183
695	Multicenter linkage study of schizophrenia candidate regions on chromosomes 5q, 6q, 10p, and 13q: schizophrenia linkage collaborative group III. *American Journal of Human Genetics*, **2000**, 67, 652-63	11	182
694	Genetics of psychosis; insights from views across the genome. *Human Genetics*, **2009**, 126, 3-12	6.3	180
693	Full genome screen for Alzheimer disease: stage II analysis. *American Journal of Medical Genetics Part A*, **2002**, 114, 235-44		179
692	Functional analysis of human promoter polymorphisms. *Human Molecular Genetics*, **2003**, 12, 2249-54	5.6	179
691	A novel Alzheimer disease locus located near the gene encoding tau protein. *Molecular Psychiatry*, **2016**, 21, 108-17	15.1	175

690	Molecular pathways involved in neuronal cell adhesion and membrane scaffolding contribute to schizophrenia and bipolar disorder susceptibility. *Molecular Psychiatry*, **2011**, 16, 286-92	15.1	175
689	Genome-wide analysis of over 106 000 individuals identifies 9 neuroticism-associated loci. *Molecular Psychiatry*, **2016**, 21, 749-57	15.1	175
688	Identification of pathways for bipolar disorder: a meta-analysis. *JAMA Psychiatry*, **2014**, 71, 657-64	14.5	172
687	Psychosis genetics: modeling the relationship between schizophrenia, bipolar disorder, and mixed (or "schizoaffective") psychoses. *Schizophrenia Bulletin*, **2009**, 35, 482-90	1.3	169
686	A replicated molecular genetic basis for subtyping antisocial behavior in children with attention-deficit/hyperactivity disorder. *Archives of General Psychiatry*, **2008**, 65, 203-10		168
685	The catechol-O-methyl transferase (COMT) gene as a candidate for psychiatric phenotypes: evidence and lessons. *Molecular Psychiatry*, **2006**, 11, 446-58	15.1	164
684	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
683	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. *Nature Neuroscience*, **2016**, 19, 420-431	25.5	163
682	Common variants at VRK2 and TCF4 conferring risk of schizophrenia. *Human Molecular Genetics*, **2011**, 20, 4076-81	5.6	162
681	Joint analysis of psychiatric disorders increases accuracy of risk prediction for schizophrenia, bipolar disorder, and major depressive disorder. *American Journal of Human Genetics*, **2015**, 96, 283-94	11	161
680	The genetics of attention deficit hyperactivity disorder. *Human Molecular Genetics*, **2005**, 14 Spec No. 2, R275-82	5.6	158
679	Haplotypes at the dystrobrevin binding protein 1 (DTNBP1) gene locus mediate risk for schizophrenia through reduced DTNBP1 expression. *Human Molecular Genetics*, **2005**, 14, 1947-54	5.6	156
678	Identification in 2 independent samples of a novel schizophrenia risk haplotype of the dystrobrevin binding protein gene (DTNBP1). *Archives of General Psychiatry*, **2004**, 61, 336-44		155
677	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
676	Universal, robust, highly quantitative SNP allele frequency measurement in DNA pools. *Human Genetics*, **2002**, 110, 471-8	6.3	152
675	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
674	Catechol O-methyltransferase gene variant and birth weight predict early-onset antisocial behavior in children with attention-deficit/hyperactivity disorder. *Archives of General Psychiatry*, **2005**, 62, 1275-8		150
673	Association of the dopamine D4 receptor gene 7-repeat allele with neuropsychological test performance of children with ADHD. *American Journal of Psychiatry*, **2004**, 161, 133-8	11.9	148

672	Investigating the contribution of common genetic variants to the risk and pathogenesis of ADHD. *American Journal of Psychiatry*, **2012**, 169, 186-94	11.9	147
671	Additional support for schizophrenia linkage on chromosomes 6 and 8: a multicenter study. Schizophrenia Linkage Collaborative Group for Chromosomes 3, 6 and 8. *American Journal of Medical Genetics Part A*, **1996**, 67, 580-94		147
670	Convergent genetic and expression data implicate immunity in Alzheimer's disease. *Alzheimer's and Dementia*, **2015**, 11, 658-71	1.2	146
669	Phenotypic Manifestation of Genetic Risk for Schizophrenia During Adolescence in the General Population. *JAMA Psychiatry*, **2016**, 73, 221-8	14.5	145
668	Genome-wide association study of schizophrenia in a Japanese population. *Biological Psychiatry*, **2011**, 69, 472-8	7.9	145
667	Genetic relationships between schizophrenia, bipolar disorder, and schizoaffective disorder. *Schizophrenia Bulletin*, **2014**, 40, 504-15	1.3	144
666	Genetic pleiotropy between multiple sclerosis and schizophrenia but not bipolar disorder: differential involvement of immune-related gene loci. *Molecular Psychiatry*, **2015**, 20, 207-14	15.1	144
665	Genotype effects of CHRNA7, CNR1 and COMT in schizophrenia: interactions with tobacco and cannabis use. *British Journal of Psychiatry*, **2007**, 191, 402-7	5.4	144
664	Genetic disruption of voltage-gated calcium channels in psychiatric and neurological disorders. *Progress in Neurobiology*, **2015**, 134, 36-54	10.9	143
663	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
662	Is COMT a susceptibility gene for schizophrenia?. *Schizophrenia Bulletin*, **2007**, 33, 635-41	1.3	143
661	Structural brain abnormalities associated with deletion at chromosome 22q11: quantitative neuroimaging study of adults with velo-cardio-facial syndrome. *British Journal of Psychiatry*, **2001**, 178, 412-9	5.4	143
660	Genomic insights into the overlap between psychiatric disorders: implications for research and clinical practice. *Genome Medicine*, **2014**, 6, 29	14.4	142
659	Schizophrenia and the neurodevelopmental continuum:evidence from genomics. *World Psychiatry*, **2017**, 16, 227-235	14.4	138
658	A scan of chromosome 10 identifies a novel locus showing strong association with late-onset Alzheimer disease. *American Journal of Human Genetics*, **2006**, 78, 78-88	11	137
657	Agreement between maternal report and antenatal records for a range of pre and peri-natal factors: the influence of maternal and child characteristics. *Early Human Development*, **2007**, 83, 497-504	2.2	135
656	A genome-wide association study for late-onset Alzheimer's disease using DNA pooling. *BMC Medical Genomics*, **2008**, 1, 44	3.7	133
655	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

654	Whole genome linkage scan of recurrent depressive disorder from the depression network study. *Human Molecular Genetics*, **2005**, 14, 3337-45	5.6	133
653	The contribution of rare variants to risk of schizophrenia in individuals with and without intellectual disability. *Nature Genetics*, **2017**, 49, 1167-1173	36.3	132
652	Logic and justification for dimensional assessment of symptoms and related clinical phenomena in psychosis: relevance to DSM-5. *Schizophrenia Research*, **2013**, 150, 15-20	3.6	132
651	Serotonergic system and attention deficit hyperactivity disorder (ADHD): a potential susceptibility locus at the 5-HT(1B) receptor gene in 273 nuclear families from a multi-centre sample. *Molecular Psychiatry*, **2002**, 7, 718-25	15.1	132
650	Evidence that duplications of 22q11.2 protect against schizophrenia. *Molecular Psychiatry*, **2014**, 19, 37-40	15.1	130
649	Meta-analysis of genome-wide association data of bipolar disorder and major depressive disorder. *Molecular Psychiatry*, **2011**, 16, 2-4	15.1	130
648	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. *Biological Psychiatry*, **2017**, 81, 325-335	7.9	129
647	Wake-up call for British psychiatry. *British Journal of Psychiatry*, **2008**, 193, 6-9	5.4	129
646	Paternal age and risk for schizophrenia. *British Journal of Psychiatry*, **2003**, 183, 405-8	5.4	128
645	Support for RGS4 as a susceptibility gene for schizophrenia. *Biological Psychiatry*, **2004**, 55, 192-5	7.9	125
644	Adolescent clinical outcomes for young people with attention-deficit hyperactivity disorder. *British Journal of Psychiatry*, **2010**, 196, 235-40	5.4	124
643	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
642	Functional gene group analysis identifies synaptic gene groups as risk factor for schizophrenia. *Molecular Psychiatry*, **2012**, 17, 996-1006	15.1	123
641	Defining the Effect of the 16p11.2 Duplication on Cognition, Behavior, and Medical Comorbidities. *JAMA Psychiatry*, **2016**, 73, 20-30	14.5	120
640	Psychosis susceptibility gene ZNF804A and cognitive performance in schizophrenia. *Archives of General Psychiatry*, **2010**, 67, 692-700		120
639	A network of dopaminergic gene variations implicated as risk factors for schizophrenia. *Human Molecular Genetics*, **2008**, 17, 747-58	5.6	120
638	Recent advances in the genetics of schizophrenia. *Human Molecular Genetics*, **2003**, 12 Spec No 2, R125-33	5.6	120
637	Novel Findings from CNVs Implicate Inhibitory and Excitatory Signaling Complexes in Schizophrenia. *Neuron*, **2015**, 86, 1203-14	13.9	119

636	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
635	Further evidence that the KIAA0319 gene confers susceptibility to developmental dyslexia. *Molecular Psychiatry*, **2006**, 11, 1085-91, 1061	15.1	119
634	Confirmation of association between expanded CAG/CTG repeats and both schizophrenia and bipolar disorder. *Psychological Medicine*, **1996**, 26, 1145-53	6.9	119
633	Accurately assessing the risk of schizophrenia conferred by rare copy-number variation affecting genes with brain function. *PLoS Genetics*, **2010**, 6, e1001097	6	118
632	A genome-wide association study in 574 schizophrenia trios using DNA pooling. *Molecular Psychiatry*, **2009**, 14, 796-803	15.1	117
631	Meta-analysis of COMT val158met in panic disorder: ethnic heterogeneity and gender specificity. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2007**, 144B, 667-73	3.5	117
630	A comprehensive family-based replication study of schizophrenia genes. *JAMA Psychiatry*, **2013**, 70, 573-81	14.5	115
629	Copy number variation in bipolar disorder. *Molecular Psychiatry*, **2016**, 21, 89-93	15.1	113
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