Stefan E Johansson

List of Publications by Year
in descending order

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135 papers	11,318 citations	50276 46 h-index	34986 98 g-index
144 all docs	144 docs citations	144 times ranked	20453 citing authors

#	Article	IF	Citations
1	Placental weight centiles adjusted for age, parity and fetal sex. Placenta, 2022, 117, 87-94.	1.5	14
2	Two New Mutations in the CEL Gene Causing Diabetes and Hereditary Pancreatitis: How to Correctly Identify MODY8 Cases. Journal of Clinical Endocrinology and Metabolism, 2022, 107, e1455-e1466.	3.6	12
3	Association of sweetened carbonated beverage consumption during pregnancy and ADHD symptoms in the offspring: a study from the Norwegian Mother, Father and Child Cohort Study (MoBa). European Journal of Nutrition, 2022, 61, 2153-2166.	3.9	3
4	Characterization of the genetic architecture of infant and early childhood body mass index. Nature Metabolism, 2022, 4, 344-358.	11.9	26
5	Population prevalence and inheritance pattern of recurrent CNVs associated with neurodevelopmental disorders in 12,252 newborns and their parents. European Journal of Human Genetics, 2021, 29, 205-215.	2.8	40
6	The position of single-base deletions in the VNTR sequence of the carboxyl ester lipase (CEL) gene determines proteotoxicity. Journal of Biological Chemistry, 2021, 296, 100661.	3.4	13
7	SeeCiTe: a method to assess CNV calls from SNP arrays using trio data. Bioinformatics, 2021, 37, 1876-1883.	4.1	4
8	GWAS for autoimmune Addisonâ€™s disease identifies multiple risk loci and highlights AIRE in disease susceptibility. Nature Communications, 2021, 12, 959.	12.8	33
9	1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. Translational Psychiatry, 2021, 11, 182.	4.8	24
10	Genome-Wide Association Study Links Autoimmune Addisonâ€™s Disease to Break of Central Tolerance. Journal of the Endocrine Society, 2021, 5, A167-A168.	0.2	0
11	Genetic Dominant Variants in STUB1, Segregating in Families with SCA48, Display In Vitro Functional Impairments Indistinctive from Recessive Variants Associated with SCAR16. International Journal of Molecular Sciences, 2021, 22, 5870.	4.1	10
12	Genetic association study of childhood aggression across raters, instruments, and age. Translational Psychiatry, 2021, 11, 413.	4.8	31
13	Transcriptional Changes in Regulatory T Cells From Patients With Autoimmune Polyendocrine Syndrome Type 1 Suggest Functional Impairment of Lipid Metabolism and Gut Homing. Frontiers in Immunology, 2021, 12, 722860.	4.8	3
14	Continuity of Genetic Risk for Aggressive Behavior Across the Life-Course. Behavior Genetics, 2021, 51, 592-606.	2.1	13
15	Chip Protein U-Box Domain Truncation Affects Purkinje Neuron Morphology and Leads to Behavioral Changes in Zebrafish. Frontiers in Molecular Neuroscience, 2021, 14, 723912.	2.9	0
16	Double paternal uniparental isodisomy 7 and 15 presenting with Beckwithâ€“Wiedemann spectrum features. Journal of Physical Education and Sports Management, 2021, 7, a006113.	1.2	5
17	Autozygosity mapping and time-to-spontaneous delivery in Norwegian parent-offspring trios. Human Molecular Genetics, 2021, 29, 3845-3858.	2.9	1
18	Comprehensive characterization of copy number variation (CNV) called from array, long- and short-read data. BMC Genomics, 2021, 22, 826.	2.8	7

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19	Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. Molecular Psychiatry, 2020, 25, 584-602.	7.9	49
20	Identification of ADHD risk genes in extended pedigrees by combining linkage analysis and whole-exome sequencing. Molecular Psychiatry, 2020, 25, 2047-2057.	7.9	17
21	Introducing M-GCTA a Software Package to Estimate Maternal (or Paternal) Genetic Effects on Offspring Phenotypes. Behavior Genetics, 2020, 50, 51-66.	2.1	18
22	Association of Copy Number Variation of the 15q11.2 BP1-BP2 Region With Cortical and Subcortical Morphology and Cognition. JAMA Psychiatry, 2020, 77, 420.	11.0	54
23	Novel loci for childhood body mass index and shared heritability with adult cardiometabolic traits. PLoS Genetics, 2020, 16, e1008718.	3.5	95
24	The genetic architecture of sporadic and multiple consecutive miscarriage. Nature Communications, 2020, 11, 5980.	12.8	52
25	Characterization of CEL-DUP2: Complete duplication of the carboxyl ester lipase gene is unlikely to influence risk of chronic pancreatitis. Pancreatology, 2020, 20, 377-384.	1.1	5
26	Shared genetic background between children and adults with attention deficit/hyperactivity disorder. Neuropsychopharmacology, 2020, 45, 1617-1626.	5.4	72
27	A trans-ancestral meta-analysis of genome-wide association studies reveals loci associated with childhood obesity. Human Molecular Genetics, 2019, 28, 3327-3338.	2.9	76
28	PathwayMatcher: proteoform-centric network construction enables fine-granularity multiomics pathway mapping. GigaScience, 2019, 8, .	6.4	4
29	Attention-deficit hyperactivity disorder shares copy number variant risk with schizophrenia and autism spectrum disorder. Translational Psychiatry, 2019, 9, 258.	4.8	75
30	Genome-wide association study reveals dynamic role of genetic variation in infant and early childhood growth. Nature Communications, 2019, 10, 4448.	12.8	61
31	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. Nature Communications, 2019, 10, 3927.	12.8	49
32	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	21.4	402
33	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	21.4	1,594
34	Further delineation of an entity caused by CREBBP and EP300 mutations but not resembling Rubinsteinâ€“Taybi syndrome. American Journal of Medical Genetics, Part A, 2018, 176, 862-876.	1.2	52
35	No evidence for rare TRAP1 mutations influencing the risk of idiopathic Parkinsonâ€™s disease. Brain, 2018, 141, e16-e16.	7.6	4
36	Novel Loci Associated With Attention-Deficit/Hyperactivity Disorder Are Revealed by Leveraging Polygenic Overlap With Educational Attainment. Journal of the American Academy of Child and Adolescent Psychiatry, 2018, 57, 86-95.	0.5	30

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37	The role of the carboxyl ester lipase (CEL) gene in pancreatic disease. Pancreatology, 2018, 18, 12-19.	1.1	60
38	Moderating effect of mode of delivery on the genetics of intelligence: Explorative genomeâ€wide analyses in ALSPAC. Brain and Behavior, 2018, 8, e01144.	2.2	6
39	Rare genetic variation in mitochondrial pathways influences the risk for Parkinson's disease. Movement Disorders, 2018, 33, 1591-1600.	3.9	51
40	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	12.6	1,085
41	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41.	21.4	286
42	PNKP Mutations Identified by Whole-Exome Sequencing in a Norwegian Patient with Sporadic Ataxia and Edema. Cerebellum, 2017, 16, 272-275.	2.5	17
43	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	27.8	544
44	Mutations in MAPKBP1 Cause Juvenile or Late-Onset Cilia-Independent Nephronophthisis. American Journal of Human Genetics, 2017, 100, 323-333.	6.2	29
45	<scp>SLC</scp>2A3 singleâ€nucleotide polymorphism and duplication influence cognitive processing and populationâ€specific risk for attentionâ€deficit/hyperactivity disorder. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2017, 58, 798-809.	5.2	25
46	In vitro characterization of six STUB1 variants in spinocerebellar ataxia 16 reveals altered structural properties for the encoded CHIP proteins. Bioscience Reports, 2017, 37, .	2.4	27
47	Functional Investigations of HNF1A Identify Rare Variants as Risk Factors for Type 2 Diabetes in the General Population. Diabetes, 2017, 66, 335-346.	0.6	54
48	Targeted next-generation sequencing reveals MODY in up to 6.5% of antibody-negative diabetes cases listed in the Norwegian Childhood Diabetes Registry. Diabetologia, 2017, 60, 625-635.	6.3	106
49	Associations between <scp>ABO</scp> blood groups and pancreatic ductal adenocarcinoma: influence on resection status and survival. Cancer Medicine, 2017, 6, 1531-1540.	2.8	26
50	The HNF1A mutant Ala180Val: Clinical challenges in determining causality of a rare HNF1A variant in familial diabetes. Diabetes Research and Clinical Practice, 2017, 133, 142-149.	2.8	6
51	Copy number variants and VNTR length polymorphisms of the carboxyl-ester lipase (CEL) gene as risk factors in pancreatic cancer. Pancreatology, 2017, 17, 83-88.	1.1	33
52	GBA2 Mutations Cause a Marinesco-SjÃ¶gren-Like Syndrome: Genetic and Biochemical Studies. PLoS ONE, 2017, 12, e0169309.	2.5	17
53	<scp>ADCK</scp>3 mutations with epilepsy, strokeâ€like episodes and ataxia: a <scp>POLG</scp> mimic?. European Journal of Neurology, 2016, 23, 1188-1194.	3.3	42
54	Novel <scp>NALCN</scp> variant: altered respiratory and circadian rhythm, anesthetic sensitivity. Annals of Clinical and Translational Neurology, 2016, 3, 876-883.	3.7	20

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55	Common and rare forms of diabetes mellitus: towards a continuum of diabetes subtypes. Nature Reviews Endocrinology, 2016, 12, 394-406.	9.6	112
56	Meta-analysis of the DRD5 VNTR in persistent ADHD. European Neuropsychopharmacology, 2016, 26, 1527-1532.	0.7	4
57	Genomeâ€wide analyses of aggressiveness in attentionâ€deficit hyperactivity disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 733-747.	1.7	40
58	Exome chip analyses in adult attention deficit hyperactivity disorder. Translational Psychiatry, 2016, 6, e923-e923.	4.8	27
59	Defective <scp>PITRM</scp> 1 mitochondrial peptidase is associated with AÎ² amyloidotic neurodegeneration. EMBO Molecular Medicine, 2016, 8, 176-190.	6.9	60
60	Hereditary hypophosphatemia in Norway: a retrospective population-based study of genotypes, phenotypes, and treatment complications. European Journal of Endocrinology, 2016, 174, 125-136.	3.7	116
61	B vitamin treatments modify the risk of myocardial infarction associated with a MTHFD1 polymorphism in patients with stable angina pectoris. Nutrition, Metabolism and Cardiovascular Diseases, 2016, 26, 495-501.	2.6	7
62	Novel SLC19A3 Promoter Deletion and Allelic Silencing in Biotin-Thiamine-Responsive Basal Ganglia Encephalopathy. PLoS ONE, 2016, 11, e0149055.	2.5	18
63	Length of Variable Numbers of Tandem Repeats in the Carboxyl Ester Lipase (CEL) Gene May Confer Susceptibility to Alcoholic Liver Cirrhosis but Not Alcoholic Chronic Pancreatitis. PLoS ONE, 2016, 11, e0165567.	2.5	16
64	On the role of NOS1 ex1fâ€VNTR in ADHDâ€”allelic, subgroup, and metaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 445-458.	1.7	20
65	Epistatic and gene wide effects in YWHA and aromatic amino hydroxylase genes across ADHD and other common neuropsychiatric disorders: Association with YWHAE. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 423-432.	1.7	21
66	Expanding the toolbox of ADHD genetics. How can we make sense of parent of origin effects in ADHD and related behavioral phenotypes?. Behavioral and Brain Functions, 2015, 11, 33.	3.3	10
67	The Chromosome 9p21 CVD- and T2D-Associated Regions in a Norwegian Population (The HUNT2 Survey). International Journal of Endocrinology, 2015, 2015, 1-9.	1.5	5
68	Common variants in the ARC gene are not associated withÂcognitive abilities. Brain and Behavior, 2015, 5, e00376.	2.2	7
69	Glycogenin-2 Is Dispensable for Liver Glycogen Synthesis and Glucagon-Stimulated Glucose Release. Journal of Clinical Endocrinology and Metabolism, 2015, 100, E767-E775.	3.6	11
70	Genome wide association study identifies variants in NBEA associated with migraine in bipolar disorder. Journal of Affective Disorders, 2015, 172, 453-461.	4.1	15
71	Dominant Mutations in the Autoimmune Regulator AIRE Are Associated with Common Organ-Specific Autoimmune Diseases. Immunity, 2015, 42, 1185-1196.	14.3	246
72	High Myopiaâ€“Excavated Optic Disc Anomaly Associated With a Frameshift Mutation in the MYC-Binding Protein 2 Gene (MYCBP2). American Journal of Ophthalmology, 2015, 159, 973-979.e2.	3.3	13

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73	A recombined allele of the lipase gene CEL and its pseudogene CELP confers susceptibility to chronic pancreatitis. Nature Genetics, 2015, 47, 518-522.	21.4	157
74	HTRA2 p.G399S in Parkinson disease, essential tremor, and tremulous cervical dystonia. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E2268.	7.1	14
75	Caseâ€“Control Genome-Wide Association Study of Persistent Attention-Deficit Hyperactivity Disorder Identifies FBXO33 as a Novel Susceptibility Gene for the Disorder. Neuropsychopharmacology, 2015, 40, 915-926.	5.4	59
76	Assessing the Causal Relationship of Maternal Height on Birth Size and Gestational Age at Birth: A Mendelian Randomization Analysis. PLoS Medicine, 2015, 12, e1001865.	8.4	121
77	Genome-Wide Analysis of Attention Deficit Hyperactivity Disorder in Norway. PLoS ONE, 2015, 10, e0122501.	2.5	71
78	STUB1 mutations in autosomal recessive ataxias â€“ evidence for mutation-specific clinical heterogeneity. Orphanet Journal of Rare Diseases, 2014, 9, 146.	2.7	63
79	Long-term clinical outcome and phenotypic variability in hyperphosphatemic familial tumoral calcinosis and hyperphosphatemic hyperostosis syndrome caused by a novel GALNT3mutation; case report and review of the literature. BMC Genetics, 2014, 15, 98.	2.7	44
80	Endocytosis of Secreted Carboxyl Ester Lipase in a Syndrome of Diabetes and Pancreatic Exocrine Dysfunction. Journal of Biological Chemistry, 2014, 289, 29097-29111.	3.4	39
81	Haploinsufficiency of MEIS2 is associated with orofacial clefting and learning disability. American Journal of Medical Genetics, Part A, 2014, 164, 1622-1626.	1.2	35
82	Loss-of-function mutations in SLC30A8 protect against type 2 diabetes. Nature Genetics, 2014, 46, 357-363.	21.4	428
83	Monoallelic and Biallelic Mutations in MAB21L2 Cause a Spectrum of Major Eye Malformations. American Journal of Human Genetics, 2014, 94, 915-923.	6.2	79
84	Unusual StÃ¼veâ€“Wiedemann syndrome with complete maternal chromosome 5 isodisomy. Annals of Clinical and Translational Neurology, 2014, 1, 926-932.	3.7	14
85	MRI characterisation of adult onset alpha-methylacyl-coA racemase deficiency diagnosed by exome sequencing. Orphanet Journal of Rare Diseases, 2013, 8, 1.	2.7	102
86	Assessing the phenotypic effects in the general population of rare variants in genes for a dominant Mendelian form of diabetes. Nature Genetics, 2013, 45, 1380-1385.	21.4	129
87	Exome sequencing reveals FAM20c mutations associated with fibroblast growth factor 23â€“related hypophosphatemia, dental anomalies, and ectopic calcification. Journal of Bone and Mineral Research, 2013, 28, 1378-1385.	2.8	144
88	Variants of Anterior Segment Dysgenesis and Cerebral Involvement in a Large Family With a Novel COL4A1 Mutation. American Journal of Ophthalmology, 2013, 155, 946-953.e2.	3.3	27
89	The number of tandem repeats in the carboxyl-ester lipase (CEL) gene as a risk factor in alcoholic and idiopathic chronic pancreatitis. Pancreatology, 2013, 13, 29-32.	1.1	38
90	SHORT Syndrome with Partial Lipodystrophy Due to Impaired Phosphatidylinositol 3 Kinase Signaling. American Journal of Human Genetics, 2013, 93, 150-157.	6.2	117

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91	DISC1 in adult ADHD patients: An association study in two European samples. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 227-234.	1.7	16
92	A novel <scp>GATA</scp>6 mutation in a child with congenital heart malformation and neonatal diabetes. Clinical Case Reports (discontinued), 2013, 1, 86-90.	0.5	10
93	Functional Properties of Rare Missense Variants of Human CDH13 Found in Adult Attention Deficit/Hyperactivity Disorder (ADHD) Patients. PLoS ONE, 2013, 8, e71445.	2.5	29
94	Novel SACS Mutations Identified by Whole Exome Sequencing in a Norwegian Family with Autosomal Recessive Spastic Ataxia of Charlevoix-Saguenay. PLoS ONE, 2013, 8, e66145.	2.5	19
95	The genetics of attention deficit/hyperactivity disorder in adults, a review. Molecular Psychiatry, 2012, 17, 960-987.	7.9	317
96	Familial Diarrhea Syndrome Caused by an ActivatingGUCY2CMutation. New England Journal of Medicine, 2012, 366, 1586-1595.	27.0	175
97	Exome Sequencing and Genetic Testing for MODY. PLoS ONE, 2012, 7, e38050.	2.5	91
98	DCLK1 Variants Are Associated across Schizophrenia and Attention Deficit/Hyperactivity Disorder. PLoS ONE, 2012, 7, e35424.	2.5	30
99	Maternal genotypes as predictors of offspring mental health: the next frontier of genomic medicine?. Future Neurology, 2011, 6, 731-743.	0.5	7
100	Bipolar disorder risk alleles in adult ADHD patients. Genes, Brain and Behavior, 2011, 10, 418-423.	2.2	17
101	No association between the serotonin transporter gene polymorphism 5-HTTLPR and cyclothymic temperament as measured by TEMPS-A. Journal of Affective Disorders, 2011, 129, 308-312.	4.1	11
102	Glutamate cysteine ligase (GCL) and self reported depression: An association study from the HUNT. Journal of Affective Disorders, 2011, 131, 207-213.	4.1	12
103	Evaluation of four novel genetic variants affecting hemoglobin A1c levels in a population-based type 2 diabetes cohort (the HUNT2 study). BMC Medical Genetics, 2011, 12, 20.	2.1	13
104	Exploring DRD4 and its interaction with SLC6A3 as possible risk factors for adult ADHD: A metaâ€analysis in four European populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 600-612.	1.7	22
105	FTO, Type 2 Diabetes, and Weight Gain Throughout Adult Life. Diabetes, 2011, 60, 1637-1644.	0.6	120
106	Diabetes and Pancreatic Exocrine Dysfunction Due to Mutations in the Carboxyl Ester Lipase Gene-Maturity Onset Diabetes of the Young (CEL-MODY). Journal of Biological Chemistry, 2011, 286, 34593-34605.	3.4	80
107	DIRAS2 is Associated with Adult ADHD, Related Traits, and Co-Morbid Disorders. Neuropsychopharmacology, 2011, 36, 2318-2327.	5.4	49
108	Physical Activity Attenuates the Influence of FTO Variants on Obesity Risk: A Meta-Analysis of 218,166 Adults and 19,268 Children. PLoS Medicine, 2011, 8, e1001116.	8.4	446

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109	Mutations in ABHD12 Cause the Neurodegenerative Disease PHARC: An Inborn Error of Endocannabinoid Metabolism. American Journal of Human Genetics, 2010, 87, 410-417.	6.2	188
110	Mutations in the VNTR of the carboxyl-ester lipase gene (CEL) are a rare cause of monogenic diabetes. Human Genetics, 2010, 127, 55-64.	3.8	61
111	Metaâ€analysis of brainâ€derived neurotrophic factor p.Val66Met in adult ADHD in four European populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 512-523.	1.7	55
112	Common variants in the TPH1 and TPH2 regions are not associated with persistent ADHD in a combined sample of 1,636 adult cases and 1,923 controls from four European populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 1008-1015.	1.7	18
113	An international multicenter association study of the serotonin transporter gene in persistent ADHD. Genes, Brain and Behavior, 2010, 9, 449-458.	2.2	55
114	A genomeâ€wide association study of bipolar disorder and comorbid migraine. Genes, Brain and Behavior, 2010, 9, 673-680.	2.2	40
115	A common variant of the latrophilin 3 gene, LPHN3, confers susceptibility to ADHD and predicts effectiveness of stimulant medication. Molecular Psychiatry, 2010, 15, 1053-1066.	7.9	245
116	Polygenic Risk Variants for Type 2 Diabetes Susceptibility Modify Age at Diagnosis in Monogenic HNF1A Diabetes. Diabetes, 2010, 59, 266-271.	0.6	37
117	Multicenter Analysis of the SLC6A3/DAT1 VNTR Haplotype in Persistent ADHD Suggests Differential Involvement of the Gene in Childhood and Persistent ADHD. Neuropsychopharmacology, 2010, 35, 656-664.	5.4	180
118	Attention-Deficit/Hyperactivity Disorder Symptoms in Offspring of Mothers With Impaired Serotonin Production. Archives of General Psychiatry, 2010, 67, 1033-1043.	12.3	47
119	Association between Catechol Oâ€methyltransferase (COMT) haplotypes and severity of hyperactivity symptoms in Adults. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 403-410.	1.7	55
120	A coding polymorphism in NALP1 confers risk for autoimmune Addison's disease and type 1 diabetes. Genes and Immunity, 2009, 10, 120-124.	4.1	167
121	Case-Control Study of Six Genes Asymmetrically Expressed in the Two Cerebral Hemispheres: Association of BAIAP2 with Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2009, 66, 926-934.	1.3	59
122	Genetic analysis of recently identified type 2 diabetes loci in 1,638 unselected patients with type 2 diabetes and 1,858 control participants from a Norwegian population-based cohort (the HUNT study). Diabetologia, 2008, 51, 971-977.	6.3	86
123	Genetic analyses of dopamine related genes in adult ADHD patients suggest an association with the DRD5â€microsatellite repeat, but not with DRD4 or SLC6A3 VNTRs. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 1470-1475.	1.7	72
124	AIRE variations in Addison's disease and autoimmune polyendocrine syndromes (APS): partial gene deletions contribute to APS I. Genes and Immunity, 2008, 9, 130-136.	4.1	36
125	A loss-of-function mutation in tryptophan hydroxylase 2 segregating with attention-deficit/hyperactivity disorder. Molecular Psychiatry, 2008, 13, 365-367.	7.9	41
126	Prevalence of HNF1A (MODY3) mutations in a Norwegian population (the HUNT2 Study). Diabetic Medicine, 2008, 25, 775-781.	2.3	40

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127	Pancreatic Exocrine Dysfunction in Maturity-Onset Diabetes of the Young Type 3. Diabetes Care, 2008, 31, 306-310.	8.6	25
128	Studies in 3,523 Norwegians and Meta-Analysis in 11,571 Subjects Indicate That Variants in the Hepatocyte Nuclear Factor 4Â (HNF4A) P2 Region Are Associated With Type 2 Diabetes in Scandinavians. Diabetes, 2007, 56, 3112-3117.	0.6	46
129	Mutations in the CEL VNTR cause a syndrome of diabetes and pancreatic exocrine dysfunction. Nature Genetics, 2006, 38, 54-62.	21.4	296
130	Linkage disequilibrium and haplotype blocks in the MHC vary in an HLA haplotype specific manner assessed mainly by DRB1*03 and DRB1*04 haplotypes. Genes and Immunity, 2006, 7, 130-140.	4.1	42
131	A Hepatocyte Nuclear Factor-4Â Gene (HNF4A) P2 Promoter Haplotype Linked With Late-Onset Diabetes: Studies of HNF4A Variants in the Norwegian MODY Registry. Diabetes, 2006, 55, 1899-1903.	0.6	33
132	HLA associations in type 1 diabetes: DPB1 alleles may act as markers of other HLA-complex susceptibility genes. Tissue Antigens, 2003, 61, 344-351.	1.0	9
133	Evidence of at least two type 1 diabetes susceptibility genes in the HLA complex distinct from HLA-DQB1, -DQA1 and â€“DRB1. Genes and Immunity, 2003, 4, 46-53.	4.1	62
134	No evidence of type 1 diabetes susceptibility genes in the region centromeric of the HLA complex. Human Immunology, 2003, 64, 951-959.	2.4	3
135	The polymorphism in the 3â€² untranslated region of IL12B has a negligible effect on the susceptibility to develop type 1 diabetes in Norway. Immunogenetics, 2001, 53, 603-605.	2.4	20