

# A Cecile J W Janssens

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

156  
papers

10,996  
citations

45  
h-index

103  
g-index

168  
ext. papers

12,615  
ext. citations

8.4  
avg, IF

5.88  
L-index

#	Paper	IF	Citations
156	Improving reporting standards for polygenic scores in risk prediction studies. <i>Nature</i> , <b>2021</b> , 591, 211-219	50.4	70
155	Personalizing Breast Cancer Screening Based on Polygenic Risk and Family History. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 434-442	9.7	10
154	Do we measure or compute polygenic risk scores? Why language matters. <i>Human Genetics</i> , <b>2021</b> , 1	6.3	1
153	Critical examination of current response shift methods and proposal for advancing new methods. <i>Quality of Life Research</i> , <b>2021</b> , 30, 3325-3342	3.7	5
152	ROC curves for clinical prediction models part 2. The ROC plot: the picture that could be worth a 1000 words. <i>Journal of Clinical Epidemiology</i> , <b>2020</b> , 126, 217-219	5.7	4
151	ROC curves for clinical prediction models part 4. Selection of the risk threshold-once chosen, always the same?. <i>Journal of Clinical Epidemiology</i> , <b>2020</b> , 126, 224-225	5.7	
150	Novel citation-based search method for scientific literature: a validation study. <i>BMC Medical Research Methodology</i> , <b>2020</b> , 20, 25	4.7	12
149	Reflection on modern methods: Revisiting the area under the ROC Curve. <i>International Journal of Epidemiology</i> , <b>2020</b> , 49, 1397-1403	7.8	52
148	It is time to get real when trying to predict educational performance. <i>ELife</i> , <b>2020</b> , 9,	8.9	0
147	Cost-Effectiveness of Risk-Stratified Colorectal Cancer Screening Based on Polygenic Risk: Current Status and Future Potential. <i>JNCI Cancer Spectrum</i> , <b>2020</b> , 4, pkz086	4.6	16
146	Validity of polygenic risk scores: are we measuring what we think we are?. <i>Human Molecular Genetics</i> , <b>2019</b> , 28, R143-R150	5.6	35
145	Proprietary Algorithms for Polygenic Risk: Protecting Scientific Innovation or Hiding the Lack of It?. <i>Genes</i> , <b>2019</b> , 10,	4.2	6
144	How the Intended Use of Polygenic Risk Scores Guides the Design and Evaluation of Prediction Studies. <i>Current Epidemiology Reports</i> , <b>2019</b> , 6, 184-190	2.9	5
143	Direct-to-Consumer Genetic Testing <b>2019</b> , 89-101		1
142	Polygenic Risk Scores That Predict Common Diseases Using Millions of Single Nucleotide Polymorphisms: Is More, Better?. <i>Clinical Chemistry</i> , <b>2019</b> , 65, 609-611	5.5	21
141	Evaluation of polygenic risk models using multiple performance measures: a critical assessment of discordant results. <i>Genetics in Medicine</i> , <b>2019</b> , 21, 391-397	8.1	12
140	Defining Evidence for Precision Medicine: A Patient Is More Than a Set of Covariates. <i>Epidemiology</i> , <b>2019</b> , 30, 342-344	3.1	

139	A collaborative translational research framework for evaluating and implementing the appropriate use of human genome sequencing to improve health. <i>PLoS Medicine</i> , <b>2018</b> , 15, e1002631	11.6	25
138	Cost Effectiveness of Age-Specific Screening Intervals for People With Family Histories of Colorectal Cancer. <i>Gastroenterology</i> , <b>2018</b> , 154, 105-116.e20	13.3	14
137	Finding Wealth in Waste: Irreplicability Re-Examined. <i>BioEssays</i> , <b>2018</b> , 40, e1800173	4.1	2
136	Lowering the P Value Threshold. <i>JAMA - Journal of the American Medical Association</i> , <b>2018</b> , 320, 936-937	27.4	
135	Predicting Polygenic Obesity Using Genetic Information. <i>Cell Metabolism</i> , <b>2017</b> , 25, 535-543	24.6	57
134	Uninformed consent in nutrigenomic research. <i>European Journal of Human Genetics</i> , <b>2017</b> , 25, 789-790	5.3	4
133	Response to letter Dr Gail. <i>Journal of Clinical Epidemiology</i> , <b>2017</b> , 85, 70-71	5.7	
132	A critical evaluation of the algorithm behind the Relative Citation Ratio (RCR). <i>PLoS Biology</i> , <b>2017</b> , 15, e2002536	9.7	22
131	Small improvement in the area under the receiver operating characteristic curve indicated small changes in predicted risks. <i>Journal of Clinical Epidemiology</i> , <b>2016</b> , 79, 159-164	5.7	8
130	The Value of Genetic Variation in the Prediction of Obesity <b>2016</b> , 441-462		
129	External validation is only needed when prediction models are worth it (Letter commenting on: J Clin Epidemiol. 2015;68:25-34). <i>Journal of Clinical Epidemiology</i> , <b>2016</b> , 69, 249-50	5.7	5
128	Risk Analysis of Prostate Cancer in PRACTICAL Consortium-Letter. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 222	4	1
127	Prediction impact curve is a new measure integrating intervention effects in the evaluation of risk models. <i>Journal of Clinical Epidemiology</i> , <b>2016</b> , 69, 89-95	5.7	8
126	Constructing Hypothetical Risk Data from the Area under the ROC Curve: Modelling Distributions of Polygenic Risk. <i>PLoS ONE</i> , <b>2016</b> , 11, e0152359	3.7	9
125	Designing babies through gene editing: science or science fiction?. <i>Genetics in Medicine</i> , <b>2016</b> , 18, 1186-1187	8.7	8
124	An Electrocardiogram-Based Risk Equation for Incident Cardiovascular Disease From the National Health and Nutrition Examination Survey. <i>JAMA Cardiology</i> , <b>2016</b> , 1, 779-786	16.2	11
123	Family history and the natural history of colorectal cancer: systematic review. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 702-12	8.1	70
122	The hidden harm behind the return of results from personal genome services: a need for rigorous and responsible evaluation. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 621-2	8.1	13

121	Returning pharmacogenetic secondary findings from genome sequencing: let's not put the cart before the horse. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 854-6	8.1	3
120	A systematic review and critical assessment of 11 discordant meta-analyses on reduced-function CYP2C19 genotype and risk of adverse clinical outcomes in clopidogrel users. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 3-11	8.1	35
119	Personal utility in genomic testing: is there such a thing?. <i>Journal of Medical Ethics</i> , <b>2015</b> , 41, 322-6	2.5	68
118	Novel citation-based search method for scientific literature: application to meta-analyses. <i>BMC Medical Research Methodology</i> , <b>2015</b> , 15, 84	4.7	21
117	Predictive or not predictive: understanding the mixed messages from the patient's DNA sequence. <i>Journal of Clinical Nursing</i> , <b>2015</b> , 24, 3730-5	3.2	0
116	Scientific reporting is suboptimal for aspects that characterize genetic risk prediction studies: a review of published articles based on the Genetic Risk Prediction Studies statement. <i>Journal of Clinical Epidemiology</i> , <b>2014</b> , 67, 487-99	5.7	7
115	Quality criteria for health checks: development of a European consensus agreement. <i>Preventive Medicine</i> , <b>2014</b> , 67, 238-41	4.3	3
114	Estimating the predictive ability of genetic risk models in simulated data based on published results from genome-wide association studies. <i>Frontiers in Genetics</i> , <b>2014</b> , 5, 179	4.5	11
113	Raw data: access to inaccuracy. <i>Science</i> , <b>2014</b> , 343, 968	33.3	2
112	Informed consent in direct-to-consumer personal genome testing: the outline of a model between specific and generic consent. <i>Bioethics</i> , <b>2014</b> , 28, 343-51	2	24
111	B-type natriuretic peptide and C-reactive protein in the prediction of atrial fibrillation risk: the CHARGE-AF Consortium of community-based cohort studies. <i>Europace</i> , <b>2014</b> , 16, 1426-33	3.9	112
110	Variations in predicted risks in personal genome testing for common complex diseases. <i>Genetics in Medicine</i> , <b>2014</b> , 16, 85-91	8.1	51
109	Ethnicity, educational level and attitudes contribute to parental intentions about genetic testing for child obesity. <i>Journal of Community Genetics</i> , <b>2013</b> , 4, 243-50	2.5	3
108	Heritability of dietary food intake patterns. <i>Acta Diabetologica</i> , <b>2013</b> , 50, 721-6	3.9	12
107	Refining genome-wide linkage intervals using a meta-analysis of genome-wide association studies identifies loci influencing personality dimensions. <i>European Journal of Human Genetics</i> , <b>2013</b> , 21, 876-82	5.3	17
106	Prediction of age-related macular degeneration in the general population: the Three Continent AMD Consortium. <i>Ophthalmology</i> , <b>2013</b> , 120, 2644-2655	7.3	69
105	Simple risk model predicts incidence of atrial fibrillation in a racially and geographically diverse population: the CHARGE-AF consortium. <i>Journal of the American Heart Association</i> , <b>2013</b> , 2, e000102	6	425
104	Incremental value of rare genetic variants for the prediction of multifactorial diseases. <i>Genome Medicine</i> , <b>2013</b> , 5, 76	14.4	8

103	Migraine is not associated with enhanced atherosclerosis. <i>Cephalalgia</i> , <b>2013</b> , 33, 228-35	6.1	47
102	How can polygenic inheritance be used in population screening for common diseases?. <i>Genetics in Medicine</i> , <b>2013</b> , 15, 437-43	8.1	40
101	A tiered-layered-staged model for informed consent in personal genome testing. <i>European Journal of Human Genetics</i> , <b>2013</b> , 21, 596-601	5.3	72
100	Longevity candidate genes and their association with personality traits in the elderly. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2012</b> , 159B, 192-200	3.5	11
99	Plasma $\beta$ -amyloid and the risk of Alzheimer's disease in Down syndrome. <i>Neurobiology of Aging</i> , <b>2012</b> , 33, 1988-94	5.6	63
98	Accuracy of self-reported family history is strongly influenced by the accuracy of self-reported personal health status of relatives. <i>Journal of Clinical Epidemiology</i> , <b>2012</b> , 65, 82-9	5.7	18
97	Ethnic differences and parental beliefs are important for overweight prevention and management in children: a cross-sectional study in the Netherlands. <i>BMC Public Health</i> , <b>2012</b> , 12, 867	4.1	12
96	Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. <i>Nature Genetics</i> , <b>2012</b> , 44, 260-8	36.3	243
95	The role of disease characteristics in the ethical debate on personal genome testing. <i>BMC Medical Genomics</i> , <b>2012</b> , 5, 4	3.7	24
94	Research conducted using data obtained through online communities: ethical implications of methodological limitations. <i>PLoS Medicine</i> , <b>2012</b> , 9, e1001328	11.6	43
93	Genome-wide profiling of blood pressure in adults and children. <i>Hypertension</i> , <b>2012</b> , 59, 241-7	8.5	28
92	Genome-wide association study identifies novel loci associated with circulating phospho- and sphingolipid concentrations. <i>PLoS Genetics</i> , <b>2012</b> , 8, e1002490	6	145
91	Analytical and simulation methods for estimating the potential predictive ability of genetic profiling: a comparison of methods and results. <i>European Journal of Human Genetics</i> , <b>2012</b> , 20, 1270-4	5.3	4
90	Interpreting incremental value of markers added to risk prediction models. <i>American Journal of Epidemiology</i> , <b>2012</b> , 176, 473-81	3.8	312
89	Personal genome testing: test characteristics to clarify the discourse on ethical, legal and societal issues. <i>BMC Medical Ethics</i> , <b>2011</b> , 12, 11	2.9	46
88	Predictive genetic testing for the identification of high-risk groups: a simulation study on the impact of predictive ability. <i>Genome Medicine</i> , <b>2011</b> , 3, 51	14.4	8
87	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. <i>Genome Medicine</i> , <b>2011</b> , 3, 16	14.4	5
86	Strengthening the reporting of Genetic Risk Prediction Studies (GRIPS): explanation and elaboration. <i>Journal of Clinical Epidemiology</i> , <b>2011</b> , 64, e1-e22	5.7	8

85	Strengthening the reporting of Genetic Risk Prediction Studies: the GRIPS Statement. <i>Journal of Clinical Epidemiology</i> , <b>2011</b> , 64, 843-7	5.7	2
84	Clinical implications of old and new genes for open-angle glaucoma. <i>Ophthalmology</i> , <b>2011</b> , 118, 2389-97	7.3	26
83	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. <i>European Journal of Clinical Investigation</i> , <b>2011</b> , 41, 1010-35	4.6	19
82	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. <i>European Journal of Clinical Investigation</i> , <b>2011</b> , 41, 1004-9	4.6	22
81	Discriminative accuracy of genomic profiling comparing multiplicative and additive risk models. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 180-5	5.3	10
80	Genetic testing and common disorders in a public health framework: how to assess relevance and possibilities. Background Document to the ESHG recommendations on genetic testing and common disorders. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19 Suppl 1, S6-44	5.3	60
79	Genetic architecture of circulating lipid levels. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 813-9	5.3	19
78	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 833-6	5.3	5
77	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 18 p preceding 494	5.3	4
76	Meta-analysis of genome-wide association for migraine in six population-based European cohorts. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 901-7	5.3	70
75	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. <i>European Journal of Epidemiology</i> , <b>2011</b> , 26, 313-37	12.1	10
74	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. <i>European Journal of Epidemiology</i> , <b>2011</b> , 26, 255-9	12.1	9
73	PredictABEL: an R package for the assessment of risk prediction models. <i>European Journal of Epidemiology</i> , <b>2011</b> , 26, 261-4	12.1	175
72	A methodological perspective on genetic risk prediction studies in type 2 diabetes: recommendations for future research. <i>Current Diabetes Reports</i> , <b>2011</b> , 11, 511-8	5.6	26
71	Strengthening the reporting of Genetic Risk Prediction Studies: the GRIPS statement. <i>Genetics in Medicine</i> , <b>2011</b> , 13, 453-6	8.1	11
70	Racial/ethnic variation in the association of lipid-related genetic variants with blood lipids in the US adult population. <i>Circulation: Cardiovascular Genetics</i> , <b>2011</b> , 4, 523-33		40
69	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. <i>Circulation: Cardiovascular Genetics</i> , <b>2011</b> , 4, 206-9		1
68	Two Authors Reply. <i>American Journal of Epidemiology</i> , <b>2011</b> , 173, 714-715	3.8	

67	Carriage of reduced-function CYP2C19 allele among patients treated with clopidogrel. <i>JAMA - Journal of the American Medical Association</i> , <b>2011</b> , 305, 467-8; author reply 468	27.4	2
66	Genetic architecture of open angle glaucoma and related determinants. <i>Journal of Medical Genetics</i> , <b>2011</b> , 48, 190-6	5.8	18
65	Genome-based prediction of breast cancer risk in the general population: a modeling study based on meta-analyses of genetic associations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2011</b> , 20, 9-22 <sup>4</sup>		28
64	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. <i>Nature Genetics</i> , <b>2011</b> , 43, 1005-11	36.3	338
63	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. <i>BMJ, The</i> , <b>2011</b> , 342, d631	5.9	14
62	Genetic risk profiling for prediction of type 2 diabetes. <i>PLOS Currents</i> , <b>2011</b> , 3, RRN1208		28
61	Strengthening the reporting of Genetic Risk Prediction Studies: the GRIPS Statement. <i>PLoS Medicine</i> , <b>2011</b> , 8, e1000420	11.6	40
60	Perspectives on the use of multiple sclerosis risk genes for prediction. <i>PLoS ONE</i> , <b>2011</b> , 6, e26493	3.7	14
59	Comparison of participant information and informed consent forms of five European studies in genetic isolated populations. <i>European Journal of Human Genetics</i> , <b>2010</b> , 18, 296-302	5.3	26
58	Biological, clinical and population relevance of 95 loci for blood lipids. <i>Nature</i> , <b>2010</b> , 466, 707-13	50.4	2742
57	Early age at menopause is associated with increased risk of dementia and mortality in women with Down syndrome. <i>Journal of Alzheimer's Disease</i> , <b>2010</b> , 19, 545-50	4.3	46
56	Genome-wide analysis of genetic loci associated with Alzheimer disease. <i>JAMA - Journal of the American Medical Association</i> , <b>2010</b> , 303, 1832-40	27.4	888
55	Improvement of risk prediction by genomic profiling: reclassification measures versus the area under the receiver operating characteristic curve. <i>American Journal of Epidemiology</i> , <b>2010</b> , 172, 353-61	3.8	53
54	Assessment of improved prediction beyond traditional risk factors: when does a difference make a difference?. <i>Circulation: Cardiovascular Genetics</i> , <b>2010</b> , 3, 3-5		24
53	A genome-wide screen for depression in two independent Dutch populations. <i>Biological Psychiatry</i> , <b>2010</b> , 68, 187-96	7.9	20
52	The apolipoprotein E gene and its age-specific effects on cognitive function. <i>Neurobiology of Aging</i> , <b>2010</b> , 31, 1831-3	5.6	51
51	Translational research in genomics of Alzheimer's disease: a review of current practice and future perspectives. <i>Journal of Alzheimer's Disease</i> , <b>2010</b> , 20, 967-80	4.3	14
50	Use of genomic profiling to assess risk for cardiovascular disease and identify individualized prevention strategies--a targeted evidence-based review. <i>Genetics in Medicine</i> , <b>2010</b> , 12, 772-84	8.1	29



49	Reply to Disease risks derived from genetic variants need clinical context <i>Genetics in Medicine</i> , <b>2010</b> , 12, 56-57	8.1	
48	HFE gene mutations increase the risk of coronary heart disease in women. <i>European Journal of Epidemiology</i> , <b>2010</b> , 25, 643-9	12.1	11
47	Using family history information to promote healthy lifestyles and prevent diseases; a discussion of the evidence. <i>BMC Public Health</i> , <b>2010</b> , 10, 248	4.1	74
46	Shared genetic factors in the co-occurrence of symptoms of depression and cardiovascular risk factors. <i>Journal of Affective Disorders</i> , <b>2010</b> , 122, 247-52	6.6	17
45	An epidemiological perspective on the future of direct-to-consumer personal genome testing. <i>Investigative Genetics</i> , <b>2010</b> , 1, 10		34
44	Impact of communicating familial risk of diabetes on illness perceptions and self-reported behavioral outcomes: a randomized controlled trial. <i>Diabetes Care</i> , <b>2009</b> , 32, 597-9	14.6	53
43	Evaluation of risk prediction updates from commercial genome-wide scans. <i>Genetics in Medicine</i> , <b>2009</b> , 11, 588-94	8.1	63
42	An empirical comparison of meta-analyses of published gene-disease associations versus consortium analyses. <i>Genetics in Medicine</i> , <b>2009</b> , 11, 153-62	8.1	9
41	How attitudes research contributes to overoptimistic expectations of personal genome testing. <i>American Journal of Bioethics</i> , <b>2009</b> , 9, 23-5	1.1	6
40	Eye color and the prediction of complex phenotypes from genotypes. <i>Current Biology</i> , <b>2009</b> , 19, R192-3	6.3	190
39	Usefulness of genetic polymorphisms and conventional risk factors to predict coronary heart disease in patients with familial hypercholesterolemia. <i>American Journal of Cardiology</i> , <b>2009</b> , 103, 375-80		27
38	Genetic scoring analysis: a way forward in genome wide association studies?. <i>European Journal of Epidemiology</i> , <b>2009</b> , 24, 585-7	12.1	16
37	Predicting human height by Victorian and genomic methods. <i>European Journal of Human Genetics</i> , <b>2009</b> , 17, 1070-5	5.3	91
36	Loci influencing lipid levels and coronary heart disease risk in 16 European population cohorts. <i>Nature Genetics</i> , <b>2009</b> , 41, 47-55	36.3	708
35	Value of genetic profiling for the prediction of coronary heart disease. <i>American Heart Journal</i> , <b>2009</b> , 158, 105-10	4.9	38
34	Genome-based prediction of common diseases: methodological considerations for future research. <i>Genome Medicine</i> , <b>2009</b> , 1, 20	14.4	65
33	The Scientific Foundation for personal genomics: recommendations from a National Institutes of Health-Centers for Disease Control and Prevention multidisciplinary workshop. <i>Genetics in Medicine</i> , <b>2009</b> , 11, 559-67	8.1	186
32	Replication of CD58 and CLEC16A as genome-wide significant risk genes for multiple sclerosis. <i>Journal of Human Genetics</i> , <b>2009</b> , 54, 676-80	4.3	57



31	A study of the SORL1 gene in Alzheimer's disease and cognitive function. <i>Journal of Alzheimer's Disease</i> , <b>2009</b> , 18, 51-64	4.3	32
30	Role of shared genetic and environmental factors in symptoms of depression and body composition. <i>Psychiatric Genetics</i> , <b>2009</b> , 19, 32-8	2.9	13
29	Genetic factors influence the clustering of depression among individuals with lower socioeconomic status. <i>PLoS ONE</i> , <b>2009</b> , 4, e5069	3.7	10
28	Cox proportional hazards models have more statistical power than logistic regression models in cross-sectional genetic association studies. <i>European Journal of Human Genetics</i> , <b>2008</b> , 16, 1111-6	5.3	33
27	Survival in elderly persons with Down syndrome. <i>Journal of the American Geriatrics Society</i> , <b>2008</b> , 56, 2311-6	5.6	90
26	Prenatal exposure to the 1944-45 Dutch 'hunger winter' and addiction later in life. <i>Addiction</i> , <b>2008</b> , 103, 433-8	4.6	78
25	Genome-based prediction of common diseases: advances and prospects. <i>Human Molecular Genetics</i> , <b>2008</b> , 17, R166-73	5.6	250
24	Predicting type 2 diabetes based on polymorphisms from genome-wide association studies: a population-based study. <i>Diabetes</i> , <b>2008</b> , 57, 3122-8	0.9	231
23	Angiotensinogen M235T polymorphism and symptoms of depression in a population-based study and a family-based study. <i>Psychiatric Genetics</i> , <b>2008</b> , 18, 162-6	2.9	9
22	Apolipoprotein E gene is related to mortality only in normal weight individuals: the Rotterdam Study. <i>European Journal of Epidemiology</i> , <b>2008</b> , 23, 135-42	12.1	9
21	Is the time right for translation research in genomics?. <i>European Journal of Epidemiology</i> , <b>2008</b> , 23, 707-10	12.1	12
20	Three genome-wide association studies and a linkage analysis identify HERC2 as a human iris color gene. <i>American Journal of Human Genetics</i> , <b>2008</b> , 82, 411-23	11	183
19	A critical appraisal of the scientific basis of commercial genomic profiles used to assess health risks and personalize health interventions. <i>American Journal of Human Genetics</i> , <b>2008</b> , 82, 593-9	11	224
18	Three Genome-wide Association Studies and a Linkage Analysis Identify HERC2 as a Human Iris Color Gene. <i>American Journal of Human Genetics</i> , <b>2008</b> , 82, 801	11	4
17	Reply to Stephan et al.. <i>American Journal of Human Genetics</i> , <b>2008</b> , 83, 131	11	2
16	Five genetic variants associated with prostate cancer. <i>New England Journal of Medicine</i> , <b>2008</b> , 358, 2739; author reply 2741	59.2	4
15	The impact of genotype frequencies on the clinical validity of genomic profiling for predicting common chronic diseases. <i>Genetics in Medicine</i> , <b>2007</b> , 9, 528-35	8.1	112
14	Patients with Barrett's esophagus perceive their risk of developing esophageal adenocarcinoma as low. <i>Gastrointestinal Endoscopy</i> , <b>2007</b> , 65, 26-30	5.2	24

13	Cholesteryl ester transfer protein gene and coronary heart disease mortality: the Rotterdam study. <i>Journal of the American Geriatrics Society</i> , <b>2007</b> , 55, 1483-4	5.6	1
12	Fatigue: an important feature of late-onset Pompe disease. <i>Journal of Neurology</i> , <b>2007</b> , 254, 941-5	5.5	46
11	Does genetic testing really improve the prediction of future type 2 diabetes?. <i>PLoS Medicine</i> , <b>2006</b> , 3, e114; author reply e127	11.6	26
10	Predictive testing for complex diseases using multiple genes: fact or fiction?. <i>Genetics in Medicine</i> , <b>2006</b> , 8, 395-400	8.1	181
9	No association between the angiotensin-converting enzyme gene and major depression: a case-control study and meta-analysis. <i>Psychiatric Genetics</i> , <b>2006</b> , 16, 225-6	2.9	13
8	Towards predictive genetic testing of complex diseases. <i>European Journal of Epidemiology</i> , <b>2006</b> , 21, 869-70	12.1	8
7	Value of the HLA-DRB1 shared epitope for predicting radiographic damage in rheumatoid arthritis depends on the individual patient risk profile. <i>Journal of Rheumatology</i> , <b>2006</b> , 33, 2383-9	4.1	6
6	A new logistic regression approach for the evaluation of diagnostic test results. <i>Medical Decision Making</i> , <b>2005</b> , 25, 168-77	2.5	35
5	Divorce and unemployment in multiple sclerosis: recalculation of the data of Morales-González J M, Benito-Leon J, Rivera-Navarro J and Mitchell A J. <i>Multiple Sclerosis Journal</i> , <b>2004</b> , 10, 716; author reply 717	5	2
4	'It might happen or it might not': how patients with multiple sclerosis explain their perception of prognostic risk. <i>Social Science and Medicine</i> , <b>2004</b> , 59, 861-8	5.1	37
3	Perception of prognostic risk in patients with multiple sclerosis: the relationship with anxiety, depression, and disease-related distress. <i>Journal of Clinical Epidemiology</i> , <b>2004</b> , 57, 180-6	5.7	35
2	Revisiting the clinical validity of multiplex genetic testing in complex diseases. <i>American Journal of Human Genetics</i> , <b>2004</b> , 74, 585-8; author reply 588-9	11	62
1	Improving reporting standards for polygenic scores in risk prediction studies		4