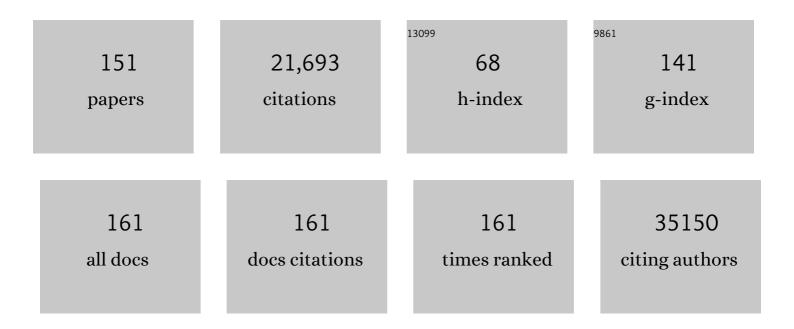
Thomas Lumley

List of Publications by Year in descending order

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THOMAS LUMIEV

#	Article	IF	CITATIONS
1	Systematic identification of trans eQTLs as putative drivers of known disease associations. Nature Genetics, 2013, 45, 1238-1243.	21.4	1,544
2	Genome-wide association study of blood pressure and hypertension. Nature Genetics, 2009, 41, 677-687.	21.4	1,224
3	The Importance of the Normality Assumption in Large Public Health Data Sets. Annual Review of Public Health, 2002, 23, 151-169.	17.4	1,192
4	Genome-wide Analysis of Genetic Loci Associated With Alzheimer Disease. JAMA - Journal of the American Medical Association, 2010, 303, 1832.	7.4	1,064
5	Network meta-analysis for indirect treatment comparisons. Statistics in Medicine, 2002, 21, 2313-2324.	1.6	985
6	Health Outcomes Associated With Various Antihypertensive Therapies Used as First-Line Agents. JAMA - Journal of the American Medical Association, 2003, 289, 2534.	7.4	869
7	New loci associated with kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 376-384.	21.4	710
8	Case???Crossover Analyses of Air Pollution Exposure Data. Epidemiology, 2005, 16, 717-726.	2.7	606
9	Multiple loci associated with indices of renal function and chronic kidney disease. Nature Genetics, 2009, 41, 712-717.	21.4	553
10	Meta-analyses of genome-wide association studies identify multiple loci associated with pulmonary function. Nature Genetics, 2010, 42, 45-52.	21.4	549
11	Genomewide Association Studies of Stroke. New England Journal of Medicine, 2009, 360, 1718-1728.	27.0	420
12	Quality control and quality assurance in genotypic data for genomeâ€wide association studies. Genetic Epidemiology, 2010, 34, 591-602.	1.3	389
13	Common variants at ten loci influence QT interval duration in the QTGEN Study. Nature Genetics, 2009, 41, 399-406.	21.4	386
14	Genome-wide association and large-scale follow up identifies 16 new loci influencing lung function. Nature Genetics, 2011, 43, 1082-1090.	21.4	367
15	Variants in ZFHX3 are associated with atrial fibrillation in individuals of European ancestry. Nature Genetics, 2009, 41, 879-881.	21.4	363
16	Association Between Blood Pressure Level and the Risk of Myocardial Infarction, Stroke, and Total Mortality. Archives of Internal Medicine, 2001, 161, 1183.	3.8	362
17	Multiple loci influence erythrocyte phenotypes in the CHARGE Consortium. Nature Genetics, 2009, 41, 1191-1198.	21.4	324
18	Genetic Loci Associated with Plasma Phospholipid n-3 Fatty Acids: A Meta-Analysis of Genome-Wide Association Studies from the CHARGE Consortium. PLoS Genetics. 2011. 7. e1002193.	3.5	324

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19	Novel Associations of Multiple Genetic Loci With Plasma Levels of Factor VII, Factor VIII, and von Willebrand Factor. Circulation, 2010, 121, 1382-1392.	1.6	311
20	Genetic Diversity and Association Studies in US Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. American Journal of Human Genetics, 2016, 98, 165-184.	6.2	266
21	Risk of New-Onset Atrial Fibrillation in Relation to Body Mass Index. Archives of Internal Medicine, 2006, 166, 2322.	3.8	258
22	Diabetes Mellitus, Glycemic Control, and Risk of Atrial Fibrillation. Journal of General Internal Medicine, 2010, 25, 853-858.	2.6	238
23	Bias in the case - crossover design: implications for studies of air pollution. Environmetrics, 2000, 11, 689-704.	1.4	222
24	Pharmacogenetic meta-analysis of genome-wide association studies of LDL cholesterol response to statins. Nature Communications, 2014, 5, 5068.	12.8	216
25	CUBN Is a Gene Locus for Albuminuria. Journal of the American Society of Nephrology: JASN, 2011, 22, 555-570.	6.1	208
26	Genomeâ€wide association studies of cerebral white matter lesion burden. Annals of Neurology, 2011, 69, 928-939.	5.3	201
27	Whole-Exome Sequencing Identifies Rare and Low-Frequency Coding Variants Associated with LDL Cholesterol. American Journal of Human Genetics, 2014, 94, 233-245.	6.2	193
28	Diuretic Therapy, the α-Adducin Gene Variant, and the Risk of Myocardial Infarction or Stroke in Persons With Treated Hypertension. JAMA - Journal of the American Medical Association, 2002, 287, 1680.	7.4	189
29	The Association Between Lipid Levels and the Risks of Incident Myocardial Infarction, Stroke, and Total Mortality: The Cardiovascular Health Study. Journal of the American Geriatrics Society, 2004, 52, 1639-1647.	2.6	186
30	Pulmonary Effects of Indoor- and Outdoor-Generated Particles in Children with Asthma. Environmental Health Perspectives, 2005, 113, 499-503.	6.0	183
31	Ambient Air Pollution and Asthma Exacerbations in Children: An Eight-City Analysis. American Journal of Epidemiology, 2006, 164, 505-517.	3.4	179
32	GWASTools: an R/Bioconductor package for quality control and analysis of genome-wide association studies. Bioinformatics, 2012, 28, 3329-3331.	4.1	177
33	Association of Genome-Wide Variation With the Risk of Incident Heart Failure in Adults of European and African Ancestry. Circulation: Cardiovascular Genetics, 2010, 3, 256-266.	5.1	176
34	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	27.8	173
35	Association of Genetic Variations With Nonfatal Venous Thrombosis in Postmenopausal Women. JAMA - Journal of the American Medical Association, 2007, 297, 489.	7.4	171
36	Association of Gene Variants With Incident Myocardial Infarction in the Cardiovascular Health Study. Arteriosclerosis, Thrombosis, and Vascular Biology, 2008, 28, 173-179.	2.4	165

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37	Genome-Wide Association Studies Identify <i>CHRNA5/3</i> and <i>HTR4</i> in the Development of Airflow Obstruction. American Journal of Respiratory and Critical Care Medicine, 2012, 186, 622-632.	5.6	164
38	A Re-Evaluation of Fixed Effect(s) Meta-Analysis. Journal of the Royal Statistical Society Series A: Statistics in Society, 2018, 181, 205-227.	1.1	159
39	Exposure assessment of particulate matter for susceptible populations in Seattle Environmental Health Perspectives, 2003, 111, 909-918.	6.0	158
40	Esterified Estrogens and Conjugated Equine Estrogens and the Risk of Venous Thrombosis. JAMA - Journal of the American Medical Association, 2004, 292, 1581.	7.4	157
41	Evaluating the Incremental Value of New Biomarkers With Integrated Discrimination Improvement. American Journal of Epidemiology, 2011, 174, 364-374.	3.4	153
42	Overlap bias in the case-crossover design, with application to air pollution exposures. Statistics in Medicine, 2005, 24, 285-300.	1.6	143
43	A Case-Crossover Analysis of Particulate Matter Air Pollution and Out-of-Hospital Primary Cardiac Arrest. Epidemiology, 2001, 12, 193-199.	2.7	138
44	Four Novel Loci (19q13, 6q24, 12q24, and 5q14) Influence the Microcirculation In Vivo. PLoS Genetics, 2010, 6, e1001184.	3.5	134
45	Genome-wide association analysis identifies six new loci associated with forced vital capacity. Nature Genetics, 2014, 46, 669-677.	21.4	131
46	Using the Whole Cohort in the Analysis of Case-Cohort Data. American Journal of Epidemiology, 2009, 169, 1398-1405.	3.4	130
47	Genome-Wide Joint Meta-Analysis of SNP and SNP-by-Smoking Interaction Identifies Novel Loci for Pulmonary Function. PLoS Genetics, 2012, 8, e1003098.	3.5	130
48	Multiethnic Meta-Analysis of Genome-Wide Association Studies in >100 000 Subjects Identifies 23 Fibrinogen-Associated Loci but No Strong Evidence of a Causal Association Between Circulating Fibrinogen and Cardiovascular Disease. Circulation, 2013, 128, 1310-1324.	1.6	128
49	Prospective Study of Particulate Air Pollution Exposures, Subclinical Atherosclerosis, and Clinical Cardiovascular Disease: The Multi-Ethnic Study of Atherosclerosis and Air Pollution (MESA Air). American Journal of Epidemiology, 2012, 176, 825-837.	3.4	126
50	Effects of ambient air pollution on symptom severity and medication use in children with asthma. Annals of Allergy, Asthma and Immunology, 2003, 91, 346-353.	1.0	119
51	Ambient air pollution, lung function, and airway responsiveness in asthmatic children. Journal of Allergy and Clinical Immunology, 2016, 137, 390-399.	2.9	119
52	Predicting intraâ€urban variation in air pollution concentrations with complex spatioâ€ŧemporal dependencies. Environmetrics, 2010, 21, 606-631.	1.4	116
53	Improved Horvitz–Thompson Estimation of Model Parameters from Two-phase Stratified Samples: Applications in Epidemiology. Statistics in Biosciences, 2009, 1, 32-49.	1.2	112
54	Insights into the Genetic Architecture of Early Stage Age-Related Macular Degeneration: A Genome-Wide Association Study Meta-Analysis. PLoS ONE, 2013, 8, e53830.	2.5	108

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55	Multiple Loci Are Associated with White Blood Cell Phenotypes. PLoS Genetics, 2011, 7, e1002113.	3.5	106
56	Efficient measurement error correction with spatially misaligned data. Biostatistics, 2011, 12, 610-623.	1.5	105
57	A Genomeâ€Wide Association Study for Venous Thromboembolism: The Extended Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Genetic Epidemiology, 2013, 37, 512-521.	1.3	99
58	A stroke prediction score in the elderly. Journal of Clinical Epidemiology, 2002, 55, 129-136.	5.0	97
59	Genetic variation associated with plasma von Willebrand factor levels and the risk of incident venous thrombosis. Blood, 2011, 117, 6007-6011.	1.4	97
60	Lower Risk of Cardiovascular Events in Postmenopausal Women Taking Oral Estradiol Compared With Oral Conjugated Equine Estrogens. JAMA Internal Medicine, 2014, 174, 25.	5.1	95
61	Behavior of QQ-Plots and Genomic Control in Studies of Gene-Environment Interaction. PLoS ONE, 2011, 6, e19416.	2.5	93
62	Cerivastatin, genetic variants, and the risk of rhabdomyolysis. Pharmacogenetics and Genomics, 2011, 21, 280-288.	1.5	90
63	Exhaled Nitric Oxide in Children with Asthma and Short-Term PM2.5Exposure in Seattle. Environmental Health Perspectives, 2005, 113, 1791-1794.	6.0	87
64	\hat{I}^2 2 -Adrenergic Receptor Polymorphisms and Risk of Incident Cardiovascular Events in the Elderly. Circulation, 2003, 107, 2021-2024.	1.6	86
65	Association of Novel Genetic Loci With Circulating Fibrinogen Levels. Circulation: Cardiovascular Genetics, 2009, 2, 125-133.	5.1	86
66	Fitting Regression Models to Survey Data. Statistical Science, 2017, 32, .	2.8	86
67	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
68	Genome-Wide Association Studies of MRI-Defined Brain Infarcts. Stroke, 2010, 41, 210-217.	2.0	82
69	Genomic Variation Associated With Mortality Among Adults of European and African Ancestry With Heart Failure. Circulation: Cardiovascular Genetics, 2010, 3, 248-255.	5.1	80
70	A network meta-analysis combined direct and indirect comparisons between glaucoma drugs to rank effectiveness in lowering intraocular pressure. Journal of Clinical Epidemiology, 2009, 62, 1279-1283.	5.0	75
71	Connections between Survey Calibration Estimators and Semiparametric Models for Incomplete Data. International Statistical Review, 2011, 79, 200-220.	1.9	67
72	Predicting Stroke Through Genetic Risk Functions. Stroke, 2014, 45, 403-412.	2.0	62

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73	Genetic Predictors of Fibrin D-Dimer Levels in Healthy Adults. Circulation, 2011, 123, 1864-1872.	1.6	60
74	Breast cancer recurrence risk in relation to antidepressant use after diagnosis. Breast Cancer Research and Treatment, 2008, 112, 123-132.	2.5	56
75	Genetic variation associated with circulating monocyte count in the eMERGE Network. Human Molecular Genetics, 2013, 22, 2119-2127.	2.9	56
76	Spatial Characteristics of Fine Particulate Matter: Identifying Representative Monitoring Locations in Seattle, Washington. Journal of the Air and Waste Management Association, 2002, 52, 324-333.	1.9	55
77	Large-Scale Genome-Wide Association Studies and Meta-Analyses of Longitudinal Change in Adult Lung Function. PLoS ONE, 2014, 9, e100776.	2.5	52
78	Exposure and measurement contributions to estimates of acute air pollution effects. Journal of Exposure Science and Environmental Epidemiology, 2005, 15, 366-376.	3.9	51
79	Common variation in cytochrome P450 epoxygenase genes and the risk of incident nonfatal myocardial infarction and ischemic stroke. Pharmacogenetics and Genomics, 2008, 18, 535-543.	1.5	51
80	Cholesterol Ester Transfer Protein, Interleukin-8, Peroxisome Proliferator Activator Receptor Alpha, and Toll-Like Receptor 4 Genetic Variations and Risk of Incident Nonfatal Myocardial Infarction and Ischemic Stroke. American Journal of Cardiology, 2008, 101, 1683-1688.	1.6	50
81	Gene Variants Associated With Ischemic Stroke. Stroke, 2009, 40, 363-368.	2.0	46
82	Antihypertensive Treatment With ACE Inhibitors or Â-Blockers and Risk of Incident Atrial Fibrillation in a General Hypertensive Population. American Journal of Hypertension, 2009, 22, 538-544.	2.0	44
83	Time trends in the use of β-blockers and other pharmacotherapies in older adults with congestive heart failure. American Heart Journal, 2004, 148, 710-717.	2.7	43
84	Simple estimates of haplotype relative risks in case-control data. Genetic Epidemiology, 2006, 30, 485-494.	1.3	43
85	Assessing seasonal confounding and model selection bias in air pollution epidemiology using positive and negative control analyses. Environmetrics, 2000, 11, 705-717.	1.4	41
86	Surrogate End Points and FDA Approval. JAMA - Journal of the American Medical Association, 2008, 299, 1474.	7.4	39
87	Model-robust regression and a Bayesian "sandwich―estimator. Annals of Applied Statistics, 2010, 4, .	1.1	39
88	Generalized estimating equations for genomeâ€wide association studies using longitudinal phenotype data. Statistics in Medicine, 2015, 34, 118-130.	1.6	37
89	Dietary fatty acids modulate associations between genetic variants and circulating fatty acids in plasma and erythrocyte membranes: Metaâ€analysis of nine studies in the CHARGE consortium. Molecular Nutrition and Food Research, 2015, 59, 1373-1383.	3.3	37
90	Potential for Revealing Individual-Level Information in Genome-wide Association Studies. JAMA - Journal of the American Medical Association, 2010, 303, 659.	7.4	32

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91	?-adrenergic receptor polymorphisms and determinants of cardiovascular risk: The Cardiovascular Health Study. American Journal of Hypertension, 2005, 18, 392-397.	2.0	28
92	Meta-analysis of genome-wide association studies of HDL cholesterol response to statins. Journal of Medical Genetics, 2016, 53, 835-845.	3.2	28
93	Genetic Loci for Retinal Arteriolar Microcirculation. PLoS ONE, 2013, 8, e65804.	2.5	27
94	White Matter Lesion Progression. Stroke, 2015, 46, 3048-3057.	2.0	27
95	Renin-Angiotensin System Haplotypes and the Risk of Myocardial Infarction and Stroke in Pharmacologically Treated Hypertensive Patients. American Journal of Epidemiology, 2007, 166, 19-27.	3.4	26
96	Common genetic variation in six lipid-related and statin-related genes, statin use and risk of incident nonfatal myocardial infarction and stroke. Pharmacogenetics and Genomics, 2008, 18, 677-682.	1.5	25
97	Graphical exploration of network meta-analysis data: the use of multidimensional scaling. Clinical Trials, 2008, 5, 301-307.	1.6	22
98	Genome-Wide Association Study of Retinopathy in Individuals without Diabetes. PLoS ONE, 2013, 8, e54232.	2.5	22
99	Conjugated Equine Estrogen, Esterified Estrogen, Prothrombotic Variants, and the Risk of Venous Thrombosis in Postmenopausal Women. Arteriosclerosis, Thrombosis, and Vascular Biology, 2006, 26, 2807-2812.	2.4	21
100	Sequence Analysis of Six Blood Pressure Candidate Regions in 4,178 Individuals: The Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Targeted Sequencing Study. PLoS ONE, 2014, 9, e109155.	2.5	19
101	Strategies to Design and Analyze Targeted Sequencing Data. Circulation: Cardiovascular Genetics, 2014, 7, 335-343.	5.1	18
102	Motivating Factors for Physician Ordering of Factor V Leiden Genetic Tests. Archives of Internal Medicine, 2009, 169, 68.	3.8	16
103	Drug-Gene Interactions of Antihypertensive Medications and Risk of Incident Cardiovascular Disease: A Pharmacogenomics Study from the CHARGE Consortium. PLoS ONE, 2015, 10, e0140496.	2.5	15
104	Raking and regression calibration: Methods to address bias from correlated covariate and timeâ€ŧoâ€event error. Statistics in Medicine, 2021, 40, 631-649.	1.6	14
105	Pathway analysis of a genome-wide gene by air pollution interaction study in asthmatic children. Journal of Exposure Science and Environmental Epidemiology, 2019, 29, 539-547.	3.9	13
106	Sequencing of <i>SCN5A</i> Identifies Rare and Common Variants Associated With Cardiac Conduction: Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Circulation: Cardiovascular Genetics, 2014, 7, 365-373.	5.1	12
107	Optimal multiwave sampling for regression modeling in twoâ€phase designs. Statistics in Medicine, 2020, 39, 4912-4921.	1.6	12
108	Validation sampling can reduce bias in health care database studies: an illustration using influenza vaccination effectiveness. Journal of Clinical Epidemiology, 2013, 66, S110-S121.	5.0	11

#	Article	IF	CITATIONS
109	Associations of NINJ2 Sequence Variants with Incident Ischemic Stroke in the Cohorts for Heart and Aging in Genomic Epidemiology (CHARGE) Consortium. PLoS ONE, 2014, 9, e99798.	2.5	11
110	On the Edge: Statistics & Computing. Chance, 2003, 16, 41-45.	0.2	8
111	Partial likelihood ratio tests for the Cox model under complex sampling. Statistics in Medicine, 2013, 32, 110-123.	1.6	8
112	No Evidence for Genome-Wide Interactions on Plasma Fibrinogen by Smoking, Alcohol Consumption and Body Mass Index: Results from Meta-Analyses of 80,607 Subjects. PLoS ONE, 2014, 9, e111156.	2.5	8
113	<i>ADAM19</i> and <i>HTR4</i> Variants and Pulmonary Function. Circulation: Cardiovascular Genetics, 2014, 7, 350-358.	5.1	8
114	Pseudo- <i>R</i> ² statistics under complex sampling. Australian and New Zealand Journal of Statistics, 2017, 59, 187-194.	0.9	8
115	Two-Phase Sampling Designs for Data Validation in Settings with Covariate Measurement Error and Continuous Outcome. Journal of the Royal Statistical Society Series A: Statistics in Society, 2021, 184, 1368-1389.	1.1	7
116	Common Genetic Variation in the Prothrombin Gene, Hormone Therapy, and Incident Nonfatal Myocardial Infarction in Postmenopausal Women. American Journal of Epidemiology, 2006, 163, 600-607.	3.4	6
117	Combining multiple imputation with raking of weights: AnÂefficient and robust approach in the setting of nearly true models. Statistics in Medicine, 2021, 40, 6777-6791.	1.6	5
118	Improved generalized raking estimators to address dependent covariate and failureâ€ŧime outcome error. Biometrical Journal, 2021, 63, 1006-1027.	1.0	4
119	An Introduction to Survival Analysis Using <i>Stata</i> (Revised Edition). Journal of Statistical Software, 2005, 12, .	3.7	4
120	Multiwave validation sampling for errorâ€prone electronic health records. Biometrics, 2023, 79, 2649-2663.	1.4	4
121	Principal Component Analysis and Factor Analysis. Wiley Series in Probability and Statistics, 2004, , 584-639.	0.0	3
122	Longitudinal Data Analysis. Wiley Series in Probability and Statistics, 2004, , 728-765.	0.0	3
123	Analysis of the Time to an Event: Survival Analysis. Wiley Series in Probability and Statistics, 2004, , 661-708.	0.0	3
124	Counting Data. Wiley Series in Probability and Statistics, 2004, , 151-207.	0.0	2
125	Categorical Data: Contingency Tables. Wiley Series in Probability and Statistics, 2004, , 208-252.	0.0	2
126	Nonparametric, Distribution-Free, and Permutation Models: Robust Procedures. Wiley Series in Probability and Statistics, 2004, , 253-290.	0.0	2

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127	Multiple Comparisons. Wiley Series in Probability and Statistics, 2004, , 520-549.	0.0	2
128	Rates and Proportions. Wiley Series in Probability and Statistics, 2004, , 640-660.	0.0	2
129	Discovering novel risk factors for venous thrombosis: a candidate-gene approach. Thrombosis Research, 2009, 123, S25-S29.	1.7	2
130	Optimal sampling for designâ€based estimators of regression models. Statistics in Medicine, 2022, 41, 1482-1497.	1.6	2
131	On the Edge: Statistics & Computing. Chance, 2002, 15, 39-42.	0.2	1
132	On the Edge: Statistics & amp; Computing. Chance, 2002, 15, 40-42.	0.2	1
133	Sample Sizes for Observational Studies. Wiley Series in Probability and Statistics, 2004, , 709-727.	0.0	1
134	Statistical Inference: Populations and Samples. Wiley Series in Probability and Statistics, 2004, , 61-116.	0.0	1
135	One- and Two-Sample Inference. Wiley Series in Probability and Statistics, 2004, , 117-150.	0.0	1
136	Discrimination and Classification. Wiley Series in Probability and Statistics, 2004, , 550-583.	0.0	1
137	Semiparametric models and two-phase samples: Applications to Cox regression. Institute of Mathematical Statistics Collections, 2013, , 65-77.	0.3	1
138	Cox Models for Ecologic Time-Series Data?. Environmental Health Perspectives, 2006, 114, A690-1; author reply A691.	6.0	1
139	On the Edge: Statistics & Computing. Chance, 2002, 15, 52-55.	0.2	0
140	On the Edge: Statistics & amp; Computing. Chance, 2003, 16, 39-44.	0.2	0
141	On the Edge: Statistics & Computing. Chance, 2003, 16, 44-48.	0.2	0
142	Symbol Index. Wiley Series in Probability and Statistics, 2004, , 867-871.	0.0	0
143	Personal Postscript. Wiley Series in Probability and Statistics, 2004, , 787-816.	0.0	0
144	On the Edge: Statistics & amp; Computing. Chance, 2004, 17, 52-54.	0.2	0

#	Article	IF	CITATIONS
145	Introduction to Biostatistics. Wiley Series in Probability and Statistics, 2004, , 1-9.	0.0	Ο
146	Biostatistical Design of Medical Studies. Wiley Series in Probability and Statistics, 2004, , 10-24.	0.0	0
147	Association and Prediction: Linear Models with One Predictor Variable. Wiley Series in Probability and Statistics, 2004, , 291-356.	0.0	0
148	Randomized Clinical Trials. Wiley Series in Probability and Statistics, 2004, , 766-786.	0.0	0
149	Association and Prediction: Multiple Regression Analysis and Linear Models with Multiple Predictor Variables. Wiley Series in Probability and Statistics, 2004, , 428-519.	0.0	Ο
150	Do Subject Characteristics Modify the Effects of Particulate Air Pollution on Daily Mortality Among the Elderly?. Journal of Occupational and Environmental Medicine, 2005, 47, 543.	1.7	0
151	On the Edge: Statistics & amp; Computing. Chance, 2002, 15, 46-49.	0.2	Ο