Lisa Cannon-Albright

List of Publications by Year
in descending order

255 papers	21,954 citations	31902 53 h-index	9839 141 g-index
268 all docs	268 docs citations	268 times ranked	23970 citing authors

#	Article	IF	Citations
1	Evidence for an Inherited Contribution to Sepsis Susceptibility Among a Cohort of U.S. Veterans. , 2022, 4, e0603.		0
2	A whole-exome case-control association study to characterize the contribution of rare coding variation to pancreatic cancer risk. Human Genetics and Genomics Advances, 2022, 3, 100078.	1.0	0
3	Evidence for excess familial clustering of Post Traumatic Stress Disorder in the US Veterans Genealogy resource. Journal of Psychiatric Research, 2022, 150, 332-337.	1.5	1
4	The Effect of Sex Hormone Deficiency on the Incidence of Rotator Cuff Repair. Journal of Bone and Joint Surgery - Series A, 2022, 104, 774-779.	1.4	9
5	Genome-wide analysis of high-risk primary brain cancer pedigrees identifies PDXDC1 as a candidate brain cancer predisposition gene. Neuro-Oncology, 2021, 23, 277-283.	0.6	3
6	A role for the MEGF6 gene in predisposition to osteoporosis. Annals of Human Genetics, 2021, 85, 58-72.	0.3	15
7	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nature Genetics, 2021, 53, 65-75.	9.4	264
8	Polygenic hazard score is associated with prostate cancer in multi-ethnic populations. Nature Communications, 2021, 12, 1236.	5.8	40
9	KLK3 SNPâ€“SNP interactions for prediction of prostate cancer aggressiveness. Scientific Reports, 2021, 11, 9264.	1.6	5
10	A Rare Variant in ERF (rs144812092) Predisposes to Prostate and Bladder Cancers in an Extended Pedigree. Cancers, 2021, 13, 2399.	1.7	4
11	Differential methylation of G-protein coupled receptor signaling genes in gastrointestinal neuroendocrine tumors. Scientific Reports, 2021, 11, 12303.	1.6	7
12	An intronic variant in the CELF4 gene is associated with risk for colorectal cancer. Cancer Epidemiology, 2021, 72, 101941.	0.8	7
13	Early-onset colorectal cancer risk extends to second- and third-degree relatives. Cancer Epidemiology, 2021, 73, 101973.	0.8	4
14	Rare Germline Variants in ATM Predispose to Prostate Cancer: A PRACTICAL Consortium Study. European Urology Oncology, 2021, 4, 570-579.	2.6	38
15	Targeted germline sequencing of patients with three or more primary melanomas reveals high rate of pathogenic variants. Melanoma Research, 2020, 30, 247-251.	0.6	5
16	Colorectal cancer risk based on extended family history and body mass index. Genetic Epidemiology, 2020, 44, 778-784.	0.6	2
17	Legal terms of use and public genealogy websites. Journal of Law and the Biosciences, 2020, 7, lsaa063.	0.8	1
18	Risk of pelvic organ prolapse treatment based onÂextended family history. American Journal of Obstetrics and Gynecology, 2020, 223, 105.e1-105.e8.	0.7	13

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19	The HOXB13 p.Gly84Glu variant observed in an extended five generation high-risk prostate cancer pedigree supports risk association for multiple cancer sites. Cancer Epidemiology, 2020, 69, 101834.	0.8	1
20	FANCM c5791C>T stopgain mutation (rs144567652) is a familial colorectal cancer risk factor. Molecular Genetics & Genomic Medicine, 2020, 8, e1532.	0.6	5
21	A novel ribosomal protein <scp>S20</scp> variant in a family with unexplained colorectal cancer and polyposis. Clinical Genetics, 2020, 97, 943-944.	1.0	14
22	The effect of sample size on polygenic hazard models for prostate cancer. European Journal of Human Genetics, 2020, 28, 1467-1475.	1.4	14
23	Pathogenic Germline DNA Repair Gene and HOXB13 Mutations in Men With Metastatic Prostate Cancer. JCO Precision Oncology, 2020, 4, 139-151.	1.5	7
24	A populationâ€based study of testicular cancer risk among children and young adults from Norway and Utah, USA. International Journal of Cancer, 2020, 147, 1604-1611.	2.3	1
25	A Genetic Risk Score to Personalize Prostate Cancer Screening, Applied to Population Data. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1731-1738.	1.1	27
26	Identification and genomic analysis of pedigrees with exceptional longevity identifies candidate rare variants. Neurobiology of Disease, 2020, 143, 104972.	2.1	7
27	Early life exposures associated with risk of small intestinal neuroendocrine tumors. PLoS ONE, 2020, 15, e0231991.	1.1	6
28	Childhood central nervous system tumors and leukemia: Incidence and familial risk. A comparative populationâ€based study in Utah and Norway. Pediatric Blood and Cancer, 2020, 67, e28408.	0.8	1
29	Early life exposures associated with risk of small intestinal neuroendocrine tumors. , 2020, 15, e0231991.		0
30	Early life exposures associated with risk of small intestinal neuroendocrine tumors. , 2020, 15, e0231991.		0
31	Early life exposures associated with risk of small intestinal neuroendocrine tumors. , 2020, 15, e0231991.		0
32	Early life exposures associated with risk of small intestinal neuroendocrine tumors. , 2020, 15, e0231991.		0
33	Population-Based Relative Risks for Lung Cancer Based on Complete Family History of Lung Cancer. Journal of Thoracic Oncology, 2019, 14, 1184-1191.	0.5	27
34	Increased risk for other cancers in individuals with Ewing sarcoma and their relatives. Cancer Medicine, 2019, 8, 7924-7930.	1.3	3
35	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
36	Predictors of Response Outcomes for Research Recruitment Through a Central Cancer Registry: Evidence From 17 Recruitment Efforts for Population-Based Studies. American Journal of Epidemiology, 2019, 188, 928-939.	1.6	9

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37	Breast cancer histologic subtypes show excess familial clustering. Cancer, 2019, 125, 3131-3138.	2.0	10
38	Population-based relative risks for specific family history constellations of breast cancer. Cancer Causes and Control, 2019, 30, 581-590.	0.8	7
39	Evidence for a Heritable Contribution toÂAtrial Fibrillation Associated WithÂFibrosis. JACC: Clinical Electrophysiology, 2019, 5, 493-500.	1.3	8
40	Relative risk for Alzheimer disease based on complete family history. Neurology, 2019, 92, e1745-e1753.	1.5	45
41	Family history of cancer and risk of paediatric and young adultâ€™s testicular cancer: A Norwegian cohort study. British Journal of Cancer, 2019, 120, 1007-1014.	2.9	12
42	Association of Rare Coding Mutations With Alzheimer Disease and Other Dementias Among Adults of European Ancestry. JAMA Network Open, 2019, 2, e191350.	2.8	58
43	Chronic Thromboembolic Pulmonary Hypertension Cases Cluster in Families. Chest, 2019, 155, 384-390.	0.4	10
44	Associations of Tobacco and Alcohol Use with Risk of Neuroendocrine Tumors of the Small Intestine in Utah. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 1998-2004.	1.1	6
45	O3â€13â€01: HIGHLY PENETRANT LATEâ€ONSET ALZHEIMER DISEASE VARIANTS IN NOTCH3 IN ASHKENAZI JEWS. Alzheimer's and Dementia, 2019, 15, P918.	0.4	0
46	Germline Variant in SLCO2B1 and Response to Abiraterone Acetate Plus Prednisone (AA) in New-onset Metastatic Castration-resistant Prostate Cancer (mCRPC). Molecular Cancer Therapeutics, 2019, 18, 726-729.	1.9	9
47	Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 208-216.	1.1	21
48	Family history of cancer and the risk of childhood solid tumours: a Norwegian nationwide register-based cohort study. British Journal of Cancer, 2018, 118, 905-912.	2.9	13
49	A Nonsynonymous Variant in the GOLM1 Gene in Cutaneous Malignant Melanoma. Journal of the National Cancer Institute, 2018, 110, 1380-1385.	3.0	23
50	Germline Variant in HSD3B1 (1245 A > C) and Response to Abiraterone Acetate Plus Prednisone in Men With New-Onset Metastatic Castration-Resistant Prostate Cancer. Clinical Genitourinary Cancer, 2018, 16, 288-292.	0.9	23
51	Genetic Contribution to Nonsquamous, Nonâ€“Small Cell Lung Cancer in Nonsmokers. Journal of Thoracic Oncology, 2018, 13, 938-945.	0.5	11
52	Polygenic hazard score to guide screening for aggressive prostate cancer: development and validation in large scale cohorts. BMJ: British Medical Journal, 2018, 360, j5757.	2.4	153
53	Similar survival of patients with multiple versus single primary melanomas basedÂon Utah Surveillance, Epidemiology, and End Results data (1973-2011). Journal of the American Academy of Dermatology, 2018, 79, 238-244.	0.6	11
54	Familial clustering of oropharyngeal squamous cell carcinoma in the Utah population. Head and Neck, 2018, 40, 384-393.	0.9	3

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55	A population-based description of familial clustering of Hirschsprung disease. Journal of Pediatric Surgery, 2018, 53, 1355-1359.	0.8	5
56	P2â€600: RELATIVE RISK FOR ALZHEIMER'Sâ€RELATED DEATH BASED ON COMPLETE FAMILY HISTORY OF ALZHEIMER'Sâ€RELATED DEATH. Alzheimer's and Dementia, 2018, 14, P967.	0.4	0
57	P1â€592: A U.S. POPULATION GENEALOGY RESOURCE SHOWS EVIDENCE OF FAMILIAL CLUSTERING FOR ALZHEIMER'S DISEASE IN VETERANS. Alzheimer's and Dementia, 2018, 14, P563.	0.4	0
58	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. Nature Communications, 2018, 9, 4616.	5.8	43
59	Consensus report of the 8 and 9th Weinman Symposia on Gene x Environment Interaction in carcinogenesis: novel opportunities for precision medicine. Cell Death and Differentiation, 2018, 25, 1885-1904.	5.0	31
60	A genealogical assessment of familial clustering of anorectal malformations. Journal of Human Genetics, 2018, 63, 1029-1034.	1.1	8
61	Pancreatic cancer as a sentinel for hereditary cancer predisposition. BMC Cancer, 2018, 18, 697.	1.1	29
62	Population genealogy resource shows evidence of familial clustering for Alzheimer disease. Neurology: Genetics, 2018, 4, e249.	0.9	6
63	AA9int: SNP interaction pattern search using non-hierarchical additive model set. Bioinformatics, 2018, 34, 4141-4150.	1.8	3
64	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. Nature Genetics, 2018, 50, 928-936.	9.4	652
65	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. Nature Communications, 2018, 9, 2256.	5.8	88
66	SNP interaction pattern identifier (SIPI): an intensive search for SNPâ€“SNP interaction patterns. Bioinformatics, 2017, 33, 822-833.	1.8	11
67	Independent Validation of Effect of HSD3B1 Genotype on Response to Androgen-Deprivation Therapy in Prostate Cancer. JAMA Oncology, 2017, 3, 856.	3.4	53
68	The Influential Role of BCL2 Family Members in Synovial Sarcomagenesis. Molecular Cancer Research, 2017, 15, 1733-1740.	1.5	10
69	Germline Variation at CDKN2A and Associations with Nevus Phenotypes amongÂMembers of Melanoma Families. Journal of Investigative Dermatology, 2017, 137, 2606-2612.	0.3	18
70	Cancer Risk in Families Fulfilling the Amsterdam Criteria for Lynch Syndrome. JAMA Oncology, 2017, 3, 1697.	3.4	32
71	Investigating the possible causal role of coffee consumption with prostate cancer risk and progression using Mendelian randomization analysis. International Journal of Cancer, 2017, 140, 322-328.	2.3	17
72	Relative Risks for Lethal Prostate Cancer Based on Complete Family History of Prostate Cancer Death. Prostate, 2017, 77, 41-48.	1.2	18

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73	Alcohol consumption and prostate cancer incidence and progression: A Mendelian randomisation study. International Journal of Cancer, 2017, 140, 75-85.	2.3	28
74	A description of familial clustering of meningiomas in the Utah population. Neuro-Oncology, 2017, 19, 1683-1687.	0.6	4
75	Linkage, whole genome sequence, and biological data implicate variants in RAB10 in Alzheimerâ€™s disease resilience. Genome Medicine, 2017, 9, 100.	3.6	67
76	Independent validation of effect of HSD3B1 genotype on response to androgen deprivation therapy (ADT) in hormone-sensitive prostate cancer (HSPC).. Journal of Clinical Oncology, 2017, 35, 172-172.	0.8	0
77	Discordant Haplotype Sequencing Identifies Functional Variants at the 2q33 Breast Cancer Risk Locus. Cancer Research, 2016, 76, 1916-1925.	0.4	7
78	A population-based survey of risk for cancer in individuals diagnosed with myotonic dystrophy. Muscle and Nerve, 2016, 54, 783-785.	1.0	25
79	PALB2,CHEK2andATMrare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811.	1.5	174
80	P2â€158: LINKAGE AND WHOLE GENOME SEQUENCE ANALYSIS OF ALZHEIMER'S DISEASE RESILIENCE AND RISK. Alzheimer's and Dementia, 2016, 12, P675.	0.4	0
81	Familiality analysis of provoked vestibulodynia treated by vestibulectomy supports genetic predisposition. American Journal of Obstetrics and Gynecology, 2016, 214, 609.e1-609.e7.	0.7	18
82	Population-based risks for cancer in patients with ALS. Neurology, 2016, 87, 289-294.	1.5	40
83	Identification of a genetic variant associated with rotator cuff repair healing. Journal of Shoulder and Elbow Surgery, 2016, 25, 865-872.	1.2	37
84	Inherited Variants in SULT1E1 and Response to Abiraterone Acetate by Men with Metastatic Castration Refractory Prostate Cancer. Journal of Urology, 2016, 196, 1112-1116.	0.2	12
85	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1609-1618.	1.1	18
86	REVEL: An Ensemble Method for Predicting the Pathogenicity of Rare Missense Variants. American Journal of Human Genetics, 2016, 99, 877-885.	2.6	1,555
87	Polyunsaturated fatty acids and prostate cancer risk: a Mendelian randomisation analysis from the PRACTICAL consortium. British Journal of Cancer, 2016, 115, 624-631.	2.9	23
88	Assessing the role of insulinâ€like growth factors and binding proteins in prostate cancer using Mendelian randomization: Genetic variants as instruments for circulating levels. International Journal of Cancer, 2016, 139, 1520-1533.	2.3	26
89	Blood lipids and prostate cancer: a Mendelian randomization analysis. Cancer Medicine, 2016, 5, 1125-1136.	1.3	68
90	Prostate cancer risk regions at 8q24 and 17q24 are differentially associated with somaticTMPRSS2:ERGfusion status. Human Molecular Genetics, 2016, 25, ddw349.	1.4	8

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91	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. Nature Communications, 2016, 7, 10933.	5.8	94
92	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	5.8	50
93	Familial aggregation of Parkinson disease in Utah. Neurology: Genetics, 2016, 2, e65.	0.9	13
94	Genome-wide association of familial prostate cancer cases identifies evidence for a rare segregating haplotype at 8q24.21. Human Genetics, 2016, 135, 923-938.	1.8	37
95	Pubertal development and prostate cancer risk: Mendelian randomization study in a population-based cohort. BMC Medicine, 2016, 14, 66.	2.3	42
96	Evidence for an Environmental and Inherited Predisposition Contributing to the Risk for Global Tendinopathies or Compression Neuropathies in Patients With Rotator Cuff Tears. Orthopaedic Journal of Sports Medicine, 2016, 4, 232596711664217.	0.8	12
97	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. Human Molecular Genetics, 2016, 25, 1663-1676.	1.4	52
98	Phenotypic and Histopathological Tumor Characteristics According to CDKN2A Mutation Status among Affected Members ofÂMelanoma Families. Journal of Investigative Dermatology, 2016, 136, 1066-1069.	0.3	13
99	Gene and pathway level analyses of germline DNA-repair gene variants and prostate cancer susceptibility using the iCOGS-genotyping array. British Journal of Cancer, 2016, 114, 945-952.	2.9	17
100	Evidence for a heritable contribution to neuroendocrine tumors of the small intestine. Endocrine-Related Cancer, 2016, 23, 93-100.	1.6	22
101	Genome-wide association study for rotator cuffÂtears identifies two significant single-nucleotide polymorphisms. Journal of Shoulder and Elbow Surgery, 2016, 25, 174-179.	1.2	46
102	Association of single nucleotide polymorphisms (SNPs) in SULT1E1 with response to treatment with abiraterone acetate (AA) in men with metastatic castration refractory prostate cancer (mCRPC).. Journal of Clinical Oncology, 2016, 34, 222-222.	0.8	0
103	Association of single nucleotide polymorphism in four genes and response to androgen deprivation therapy (ADT) in men with advanced hormone sensitive prostate cancer (aHSPC).. Journal of Clinical Oncology, 2016, 34, 224-224.	0.8	0
104	Association of single nucleotide polymorphisms (SNPs) in TPD52 gene with response to treatment with enzalutamide (ENZA) in men with metastatic castration refractory prostate cancer (mCRPC).. Journal of Clinical Oncology, 2016, 34, 5066-5066.	0.8	0
105	Prostate cancer risk prediction based on complete prostate cancer family history. Prostate, 2015, 75, 390-398.	1.2	91
106	Relative risks for comorbidities associated with myotonic dystrophy: A populationâ€based analysis. Muscle and Nerve, 2015, 52, 659-661.	1.0	23
107	Prediction of individual genetic risk to prostate cancer using a polygenic score. Prostate, 2015, 75, 1467-1474.	1.2	54
108	Risk of Associated Conditions in Relatives of Subjects With Interstitial Cystitis. Obstetrical and Gynecological Survey, 2015, 70, 439-440.	0.2	0

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109	Rarity of the Alzheimer Diseaseâ€“ProtectiveAPPA673T Variant in the United States. JAMA Neurology, 2015, 72, 209.	4.5	41
110	Significant association of full-thickness rotator cuff tears and estrogen-related receptor-Î² (ESRRB). Journal of Shoulder and Elbow Surgery, 2015, 24, e31-e35.	1.2	48
111	A Large-Scale Analysis of Genetic Variants within Putative miRNA Binding Sites in Prostate Cancer. Cancer Discovery, 2015, 5, 368-379.	7.7	56
112	Associations of prostate cancer risk variants with disease aggressiveness: results of the NCI-SPORE Genetics Working Group analysis of 18,343 cases. Human Genetics, 2015, 134, 439-450.	1.8	45
113	A Heritable Predisposition to Osteoarthritis of the Hip. Journal of Arthroplasty, 2015, 30, 125-129.	1.5	6
114	Evidence for pelvic organ prolapse predispositionÂgenesÂon chromosomes 10 and 17. American Journal of Obstetrics and Gynecology, 2015, 212, 771.e1-771.e7.	0.7	24
115	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. Nature Communications, 2015, 6, 7539.	5.8	38
116	Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1121-1129.	1.1	56
117	Risk of Associated Conditions in Relatives of Subjects With Interstitial Cystitis. Female Pelvic Medicine and Reconstructive Surgery, 2015, 21, 93-98.	0.6	14
118	The effects of height and BMI on prostate cancer incidence and mortality: a Mendelian randomization study in 20,848 cases and 20,214 controls from the PRACTICAL consortium. Cancer Causes and Control, 2015, 26, 1603-1616.	0.8	77
119	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. Human Molecular Genetics, 2015, 24, 5589-5602.	1.4	67
120	Evidence for a genetical contribution to non-smoking-related lung cancer. Thorax, 2015, 70, 1033-1039.	2.7	21
121	Genome-Wide Association Study of Prostate Cancerâ€“Specific Survival. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1796-1800.	1.1	27
122	BRCA1, BRCA2, PALB2, and CDKN2A mutations in familial pancreatic cancer: a PACGENE study. Genetics in Medicine, 2015, 17, 569-577.	1.1	231
123	Association of single nucleotide polymorphisms (SNPs) in ESR1 and PRMT8 and response to treatment with abiraterone acetate (AA) in men with metastatic castration refractory prostate cancer (mCRPC).. Journal of Clinical Oncology, 2015, 33, 5048-5048.	0.8	2
124	A genome-wide linkage study of lethal prostate cancer predisposition gene in a set of high-risk pedigrees.. Journal of Clinical Oncology, 2015, 33, 159-159.	0.8	0
125	Germ line predictors of response to androgen deprivation therapy in men with advanced prostate cancer.. Journal of Clinical Oncology, 2015, 33, 162-162.	0.8	0
126	Risk for death from prostate cancer predicted from complete family history of lethal prostate cancer (LPC).. Journal of Clinical Oncology, 2015, 33, 175-175.	0.8	1

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127	Significant Evidence of Linkage for a Gene Predisposing to Colorectal Cancer and Multiple Primary Cancers on 22q11. Clinical and Translational Gastroenterology, 2014, 5, e50.	1.3	5
128	Fine-Mapping the HOXB Region Detects Common Variants Tagging a Rare Coding Allele: Evidence for Synthetic Association in Prostate Cancer. PLoS Genetics, 2014, 10, e1004129.	1.5	34
129	Co-prevalence of other tumors in patients harboring pituitary tumors. Journal of Neurosurgery, 2014, 121, 1474-1477.	0.9	12
130	Familial clustering of ALS in a population-based resource. Neurology, 2014, 82, 17-22.	1.5	33
131	Identification of specific Y chromosomes associated with increased prostate cancer risk. Prostate, 2014, 74, 991-998.	1.2	11
132	Association analysis of 9,560 prostate cancer cases from the International Consortium of Prostate Cancer Genetics confirms the role of reported prostate cancer associated SNPs for familial disease. Human Genetics, 2014, 133, 347-356.	1.8	24
133	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	9.4	408
134	The familiality of pelvic organ prolapse in the Utah Population Database. International Urogynecology Journal, 2013, 24, 413-418.	0.7	21
135	Use of a Genealogical Database Demonstrates Heritability of Pulmonary Fibrosis. Lung, 2013, 191, 475-481.	1.4	16
136	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. Human Molecular Genetics, 2013, 22, 4239-4239.	1.4	2
137	HOXB13 is a susceptibility gene for prostate cancer: results from the International Consortium for Prostate Cancer Genetics (ICPCG). Human Genetics, 2013, 132, 5-14.	1.8	166
138	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. Nature Genetics, 2013, 45, 385-391.	9.4	492
139	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. Nature Genetics, 2013, 45, 868-876.	9.4	179
140	Empiric evidence for a genetic contribution to predisposition to surgical site infection. Wound Repair and Regeneration, 2013, 21, 211-215.	1.5	8
141	Report of a novel OCA2 gene mutation and an investigation of OCA2 variants on melanoma risk in a familial melanoma pedigree. Journal of Dermatological Science, 2013, 69, 30-37.	1.0	32
142	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. Human Molecular Genetics, 2013, 22, 408-415.	1.4	118
143	Creation of a national resource with linked genealogy and phenotypic data: the Veterans Genealogy Project. Genetics in Medicine, 2013, 15, 541-547.	1.1	11
144	Linkage Analysis of Extended High-Risk Pedigrees Replicates a Cutaneous Malignant Melanoma Predisposition Locus on Chromosome 9q21. Journal of Investigative Dermatology, 2013, 133, 128-134.	0.3	9

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145	Lack of GNAQ and GNA11 Germ-Line Mutations in Familial Melanoma Pedigrees with Uveal Melanoma or Blue Nevi. Frontiers in Oncology, 2013, 3, 160.	1.3	15
146	Genetic Analysis of Low BMI Phenotype in the Utah Population Database. PLoS ONE, 2013, 8, e80287.	1.1	8
147	A population-based analysis of clustering identifies a strong genetic contribution to lethal prostate cancer. Frontiers in Genetics, 2013, 4, 152.	1.1	10
148	Associations of ATR and CHEK1 Single Nucleotide Polymorphisms with Breast Cancer. PLoS ONE, 2013, 8, e68578.	1.1	13
149	Strong Evidence for a Genetic Contribution to Late-Onset Alzheimerâ€™s Disease Mortality: A Population-Based Study. PLoS ONE, 2013, 8, e77087.	1.1	14
150	At-Risk Populations for Osteosarcoma: The Syndromes and Beyond. Sarcoma, 2012, 2012, 1-9.	0.7	66
151	Fine-Mapping CASP8 Risk Variants in Breast Cancer. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 176-181.	1.1	21
152	A comprehensive survey of cancer risks in extended families. Genetics in Medicine, 2012, 14, 107-114.	1.1	56
153	Evidence of an Inherited Predisposition for Cervical Spondylotic Myelopathy. Spine, 2012, 37, 26-29.	1.0	30
154	The Impact of Family History on the Risk of Colorectal Neoplasia: Donâ€™t Change the Guidelines Just Yet!. Digestive Diseases and Sciences, 2012, 57, 3047-3049.	1.1	3
155	Analysis of Xq27-28 linkage in the international consortium for prostate cancer genetics (ICPCG) families. BMC Medical Genetics, 2012, 13, 46.	2.1	5
156	Significant evidence for a heritable contribution to cancer predisposition: a review of cancer familiality by site. BMC Cancer, 2012, 12, 138.	1.1	52
157	Shared Predispositions of Parkinsonism and Cancer. Archives of Neurology, 2012, 69, 1572.	4.9	71
158	Pairwise shared genomic segment analysis in three Utah high-risk breast cancer pedigrees. BMC Genomics, 2012, 13, 676.	1.2	0
159	Chromosomes 4 and 8 implicated in a genome wide SNP linkage scan of 762 prostate cancer families collected by the ICPCG. Prostate, 2012, 72, 410-426.	1.2	14
160	Validation of prostate cancer risk-related loci identified from genome-wide association studies using family-based association analysis: evidence from the International Consortium for Prostate Cancer Genetics (ICPCG). Human Genetics, 2012, 131, 1095-1103.	1.8	21
161	A unique genome-wide association analysis in extended Utah high-risk pedigrees identifies a novel melanoma risk variant on chromosome arm 10q. Human Genetics, 2012, 131, 77-85.	1.8	24
162	Genome-wide association study identifies three new melanoma susceptibility loci. Nature Genetics, 2011, 43, 1108-1113.	9.4	230

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163	Identification of Six Loci Associated With Pelvic Organ Prolapse Using Genome-Wide Association Analysis. Obstetrics and Gynecology, 2011, 118, 1345-1353.	1.2	47
164	Comparison of compliance for colorectal cancer screening and surveillance by colonoscopy based on risk. Genetics in Medicine, 2011, 13, 737-743.	1.1	57
165	Identification of regions of positive selection using Shared Genomic Segment analysis. European Journal of Human Genetics, 2011, 19, 667-671.	1.4	17
166	Familial clustering of endometrial cancer in a well-defined population. Gynecologic Oncology, 2011, 122, 75-78.	0.6	15
167	A role for XRCC2 gene polymorphisms in breast cancer risk and survival. Journal of Medical Genetics, 2011, 48, 477-484.	1.5	47
168	Evidence for a heritable predisposition to Chronic Fatigue Syndrome. BMC Neurology, 2011, 11, 62.	0.8	68
169	Strategies for selection of subjects for sequencing after detection of a linkage peak. BMC Proceedings, 2011, 5, S77.	1.8	4
170	High quality and quantity Genome-wide germline genotypes from FFPE normal tissue. BMC Research Notes, 2011, 4, 159.	0.6	12
171	How well does family history predict who will get colorectal cancer? Implications for cancer screening and counseling. Genetics in Medicine, 2011, 13, 385-391.	1.1	25
172	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. Nature Genetics, 2011, 43, 785-791.	9.4	265
173	Complex Genotype Sarcomas Display Familial Inheritance Independent of Known Cancer Predisposition Syndromes. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 751-757.	1.1	6
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