

# Manuel R. Teixeira

## List of Publications by Year in descending order

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Version: 2024-02-01

319  
papers

16,465  
citations

18465

62  
h-index

26591

107  
g-index

340  
all docs

340  
docs citations

340  
times ranked

22829  
citing authors

#	ARTICLE	IF	CITATIONS
1	Breast and Prostate Cancer Risks for Male <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variant Carriers Using Polygenic Risk Scores. <i>Journal of the National Cancer Institute</i> , 2022, 114, 109-122.	3.0	19
2	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
3	Expression Profiling in Ovarian Cancer Reveals Coordinated Regulation of <i>BRCA1/2</i> and Homologous Recombination Genes. <i>Biomedicines</i> , 2022, 10, 199.	1.4	3
4	Male gender as a poor prognostic factor in Medullary Thyroid Carcinoma: behaviour or biological difference?. <i>Minerva Endocrinology</i> , 2022, , .	0.6	1
5	Prostate cancer risk stratification improvement across multiple ancestries with new polygenic hazard score. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, 25, 755-761.	2.0	14
6	<i>KRAS</i> and <i>NRAS</i> mutational analysis in plasma ctDNA from patients with metastatic colorectal cancer by real-time PCR and digital PCR. <i>International Journal of Colorectal Disease</i> , 2022, 37, 895-905.	1.0	3
7	The role of <i>TP53</i> pathogenic variants in early-onset <i>HER2</i> -positive breast cancer. <i>Familial Cancer</i> , 2021, 20, 173-180.	0.9	2
8	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , 2021, 53, 65-75.	9.4	264
9	Additional SNPs improve risk stratification of a polygenic hazard score for prostate cancer. <i>Prostate Cancer and Prostatic Diseases</i> , 2021, 24, 532-541.	2.0	16
10	Polygenic hazard score is associated with prostate cancer in multi-ethnic populations. <i>Nature Communications</i> , 2021, 12, 1236.	5.8	40
11	Multi-Gene Panel Testing in Gastroenterology: Are We Ready for the Results?. <i>GE Portuguese Journal of Gastroenterology</i> , 2021, 28, 1-7.	0.3	2
12	Case Report: Pheochromocytoma and Synchronous Neuroblastoma in a Family With Hereditary Pheochromocytoma Associated With a <i>MAX</i> Deleterious Variant. <i>Frontiers in Endocrinology</i> , 2021, 12, 609263.	1.5	4
13	<i>KLK3</i> SNP-SNP interactions for prediction of prostate cancer aggressiveness. <i>Scientific Reports</i> , 2021, 11, 9264.	1.6	5
14	When to Stop TKIs in Patients with Chronic Myeloid Leukemia and How to Follow Them Subsequently. <i>Current Treatment Options in Oncology</i> , 2021, 22, 49.	1.3	1
15	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous <i>BRCA1</i> or <i>BRCA2</i> pathogenic variant. <i>Genetics in Medicine</i> , 2021, 23, 1726-1737.	1.1	16
16	Rare Germline Variants in <i>ATM</i> Predispose to Prostate Cancer: A PRACTICAL Consortium Study. <i>European Urology Oncology</i> , 2021, 4, 570-579.	2.6	38
17	Next Generation Sequencing of Tumor and Matched Plasma Samples: Identification of Somatic Variants in ctDNA From Ovarian Cancer Patients. <i>Frontiers in Oncology</i> , 2021, 11, 754094.	1.3	5
18	Identification of <i>SPRY4</i> as a Novel Candidate Susceptibility Gene for Familial Nonmedullary Thyroid Cancer. <i>Thyroid</i> , 2021, 31, 1366-1375.	2.4	9

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19	A prospective prostate cancer screening programme for men with pathogenic variants in mismatch repair genes (IMPACT): initial results from an international prospective study. <i>Lancet Oncology</i> , The, 2021, 22, 1618-1631.	5.1	48
20	Prognostic Value of Histone Modifying Enzyme EZH2 in RCHOP-Treated Diffuse Large B-Cell Lymphoma and High Grade B-Cell Lymphoma. <i>Journal of Personalized Medicine</i> , 2021, 11, 1384.	1.1	0
21	Association of Genomic Domains in <i>BRCA1</i> and <i>BRCA2</i> with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , 2020, 80, 624-638.	0.4	39
22	Widening the spectrum of Lynch syndrome: first report of testicular seminoma attributable to MSH2 loss. <i>Histopathology</i> , 2020, 76, 486-489.	1.6	5
23	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
24	Pathogenicity reclassification of two <i>BRCA1/BRCA2</i> exonic duplications after identification of genomic breakpoints and tandem orientation. <i>Cancer Genetics</i> , 2020, 248-249, 18-24.	0.2	0
25	Haplotype analysis of the internationally distributed <i>BRCA1</i> c.3331_3334delCAAG founder mutation reveals a common ancestral origin in Iberia. <i>Breast Cancer Research</i> , 2020, 22, 108.	2.2	9
26	Gene Panel Tumor Testing in Ovarian Cancer Patients Significantly Increases the Yield of Clinically Actionable Germline Variants beyond <i>BRCA1/BRCA2</i> . <i>Cancers</i> , 2020, 12, 2834.	1.7	6
27	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	1.1	82
28	Hereditary Predisposition to Prostate Cancer: From Genetics to Clinical Implications. <i>International Journal of Molecular Sciences</i> , 2020, 21, 5036.	1.8	38
29	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. <i>Nature Communications</i> , 2020, 11, 3905.	5.8	28
30	Tumor Testing for Somatic and Germline <i>BRCA1/BRCA2</i> Variants in Ovarian Cancer Patients in the Context of Strong Founder Effects. <i>Frontiers in Oncology</i> , 2020, 10, 1318.	1.3	11
31	Association of germline variation with the survival of women with <i>BRCA1/2</i> pathogenic variants and breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 44.	2.3	5
32	The <i>CHEK2</i> Variant C.349A>G Is Associated with Prostate Cancer Risk and Carriers Share a Common Ancestor. <i>Cancers</i> , 2020, 12, 3254.	1.7	16
33	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581.	9.4	265
34	The effect of sample size on polygenic hazard models for prostate cancer. <i>European Journal of Human Genetics</i> , 2020, 28, 1467-1475.	1.4	14
35	Surveillance of succinate dehydrogenase gene mutation carriers: Insights from a nationwide cohort. <i>Clinical Endocrinology</i> , 2020, 92, 545-553.	1.2	10
36	A Genetic Risk Score to Personalize Prostate Cancer Screening, Applied to Population Data. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1731-1738.	1.1	27

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37	Characterization of the Cancer Spectrum in Men With Germline <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>JAMA Oncology</i> , 2020, 6, 1218.	3.4	48
38	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
39	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
40	The Spectrum of FANCM Protein Truncating Variants in European Breast Cancer Cases. <i>Cancers</i> , 2020, 12, 292.	1.7	11
41	Myeloid Disease with the CSF3R T618I Mutation after CLL. <i>Case Reports in Hematology</i> , 2020, 2020, 1-4.	0.3	1
42	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	2.3	28
43	Interim Results from the IMPACT Study: Evidence for Prostate-specific Antigen Screening in BRCA2 Mutation Carriers. <i>European Urology</i> , 2019, 76, 831-842.	0.9	148
44	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
45	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	2.9	19
46	Negative MR4 chronic myeloid leukaemia and its possible implications for treatment-free remission. <i>British Journal of Haematology</i> , 2019, 186, e181-e184.	1.2	1
47	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
48	The nonsense mutation <i>MSH2</i> c.2152C>T shows a founder effect in Portuguese Lynch syndrome families. <i>Genes Chromosomes and Cancer</i> , 2019, 58, 657-664.	1.5	3
49	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	2.9	52
50	Genetic counselling and testing of susceptibility genes for therapeutic decision-making in breast cancer: an European consensus statement and expert recommendations. <i>European Journal of Cancer</i> , 2019, 106, 54-60.	1.3	25
51	Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 208-216.	1.1	21
52	Height and Body Mass Index as Modifiers of Breast Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019, 111, 350-364.	3.0	30
53	Mutational spectrum in a worldwide study of 29,700 families with <i>BRCA1</i> or <i>BRCA2</i> mutations. <i>Human Mutation</i> , 2018, 39, 593-620.	1.1	224
54	Contribution of <i>MLH1</i> constitutional methylation for Lynch syndrome diagnosis in patients with tumor <i>MLH1</i> downregulation. <i>Cancer Medicine</i> , 2018, 7, 433-444.	1.3	28

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55	Prostate-specific antigen velocity in a prospective prostate cancer screening study of men with genetic predisposition. <i>British Journal of Cancer</i> , 2018, 118, 266-276.	2.9	12
56	Polygenic hazard score to guide screening for aggressive prostate cancer: development and validation in large scale cohorts. <i>BMJ: British Medical Journal</i> , 2018, 360, j5757.	2.4	153
57	Germline pathogenic variants in PALB2 and other cancer-predisposing genes in families with hereditary diffuse gastric cancer without CDH1 mutation: a whole-exome sequencing study. <i>The Lancet Gastroenterology and Hepatology</i> , 2018, 3, 489-498.	3.7	87
58	Genetic Testing and Clinical Management Practices for Variants in Non-BRCA1/2 Breast (and) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 632 for the Interpretation of Germline Mutant Alleles (ENIGMA) Clinical Working Group. <i>JCO Precision Oncology</i> , 2018, 2, 1-42.	1.5	19
59	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. <i>Nature Communications</i> , 2018, 9, 4616.	5.8	43
60	Discontinuation of tyrosine kinase inhibitors in CML patients in real-world clinical practice at a single institution. <i>BMC Cancer</i> , 2018, 18, 1245.	1.1	15
61	Potential clinical applications of circulating cell-free DNA in ovarian cancer patients. <i>Expert Reviews in Molecular Medicine</i> , 2018, 20, e6.	1.6	20
62	Ponatinib induces a sustained deep molecular response in a chronic myeloid leukaemia patient with an early relapse with a T315I mutation following allogeneic hematopoietic stem cell transplantation: a case report. <i>BMC Cancer</i> , 2018, 18, 1229.	1.1	11
63	Screening and characterization of BRCA2 c.156_157insAlu in Brazil: Results from 1380 individuals from the South and Southeast. <i>Cancer Genetics</i> , 2018, 228-229, 93-97.	0.2	6
64	Hybrid oncocytic/chromophobe renal cell tumor: An integrated genetic and epigenetic characterization of a case. <i>Experimental and Molecular Pathology</i> , 2018, 105, 352-356.	0.9	3
65	Targeted next generation sequencing identifies functionally deleterious germline mutations in novel genes in early-onset/familial prostate cancer. <i>PLoS Genetics</i> , 2018, 14, e1007355.	1.5	50
66	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2018, 78, 5419-5430.	0.4	54
67	AA9int: SNP interaction pattern search using non-hierarchical additive model set. <i>Bioinformatics</i> , 2018, 34, 4141-4150.	1.8	3
68	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018, 50, 928-936.	9.4	652
69	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018, 9, 2256.	5.8	88
70	Detection of microsatellite instability (MSI) in colorectal cancer samples with a novel set of highly sensitive markers by means of the Idylla MSI Test prototype.. <i>Journal of Clinical Oncology</i> , 2018, 36, e15639-e15639.	0.8	12
71	Full in-frame exon 3 skipping of <i>BRCA2</i> confers high risk of breast and/or ovarian cancer. <i>Oncotarget</i> , 2018, 9, 17334-17348.	0.8	24
72	SNP interaction pattern identifier (SIPI): an intensive search for SNP-SNP interaction patterns. <i>Bioinformatics</i> , 2017, 33, 822-833.	1.8	11

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73	Identification of somatic <i>TERT</i> promoter mutations in familial nonmedullary thyroid carcinomas. <i>Clinical Endocrinology</i> , 2017, 87, 394-399.	1.2	23
74	Ovarian metastasis from uveal melanoma with MLH1/PMS2 protein loss in a patient with germline MLH1 mutated Lynch syndrome: consequence or coincidence?. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2017, 470, 347-352.	1.4	9
75	Validation of a Next-Generation Sequencing Pipeline for the Molecular Diagnosis of Multiple Inherited Cancer Predisposing Syndromes. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 502-513.	1.2	13
76	Bromodomain protein 4 discriminates tissue-specific super-enhancers containing disease-specific susceptibility loci in prostate and breast cancer. <i>BMC Genomics</i> , 2017, 18, 270.	1.2	26
77	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
78	Germline Mutations in PALB2, BRCA1, and RAD51C, Which Regulate DNA Recombination Repair, in Patients With Gastric Cancer. <i>Gastroenterology</i> , 2017, 152, 983-986.e6.	0.6	98
79	<i>POLE</i> somatic mutations in advanced colorectal cancer. <i>Cancer Medicine</i> , 2017, 6, 2966-2971.	1.3	43
80	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
81	Height, selected genetic markers and prostate cancer risk: results from the PRACTICAL consortium. <i>British Journal of Cancer</i> , 2017, 117, 734-743.	2.9	7
82	Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility. <i>Nature Communications</i> , 2017, 8, 1892.	5.8	40
83	Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 117-134.	1.1	18
84	Investigating the possible causal role of coffee consumption with prostate cancer risk and progression using Mendelian randomization analysis. <i>International Journal of Cancer</i> , 2017, 140, 322-328.	2.3	17
85	Performance of Lynch syndrome predictive models in quantifying the likelihood of germline mutations in patients with abnormal MLH1 immunoreexpression. <i>Familial Cancer</i> , 2017, 16, 73-81.	0.9	2
86	Alcohol consumption and prostate cancer incidence and progression: A Mendelian randomisation study. <i>International Journal of Cancer</i> , 2017, 140, 75-85.	2.3	28
87	Evaluation of Polygenic Risk Scores for Breast and Ovarian Cancer Risk Prediction in BRCA1 and BRCA2 Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	3.0	242
88	Prediction of Breast and Prostate Cancer Risks in Male <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers Using Polygenic Risk Scores. <i>Journal of Clinical Oncology</i> , 2017, 35, 2240-2250.	0.8	152
89	Abstract LB-158: Germline mutations in PALB2, BRCA1 and RAD51C observed in gastric cancer cases. , 2017, , .		0
90	BRCA1 and BRCA2 rearrangements in Brazilian individuals with Hereditary Breast and Ovarian Cancer Syndrome. <i>Genetics and Molecular Biology</i> , 2016, 39, 223-231.	0.6	22

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91	Lapatinib-capecitabine versus capecitabine alone as radiosensitizers in RAS wild-type resectable rectal cancer, an adaptive randomized phase II trial (LaRRC trial): study protocol for a randomized controlled trial. <i>Trials</i> , 2016, 17, 459.	0.7	2
92	Fine-Scale Mapping at 9p22.2 Identifies Candidate Causal Variants That Modify Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>PLoS ONE</i> , 2016, 11, e0158801.	1.1	10
93	Analysis of Founder Mutations in Rare Tumors Associated With Hereditary Breast/Ovarian Cancer Reveals a Novel Association of BRCA2 Mutations with Ampulla of Vater Carcinomas. <i>PLoS ONE</i> , 2016, 11, e0161438.	1.1	15
94	Oncogenic mechanisms of HOXB13 missense mutations in prostate carcinogenesis. <i>Oncoscience</i> , 2016, 3, 288-296.	0.9	11
95	<i>NCOA2</i> is a candidate target gene of 8q gain associated with clinically aggressive prostate cancer. <i>Genes Chromosomes and Cancer</i> , 2016, 55, 365-374.	1.5	14
96	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	1.5	174
97	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	2.2	31
98	Correspondence: SEMA4A variation and risk of colorectal cancer. <i>Nature Communications</i> , 2016, 7, 10611.	5.8	7
99	Prostate Cancer Prognosis Defined by the Combined Analysis of 8q, PTEN and ERG. <i>Translational Oncology</i> , 2016, 9, 575-582.	1.7	12
100	Pathologic Findings in Prophylactic and Nonprophylactic Hysterectomy Specimens of Patients With Lynch Syndrome. <i>American Journal of Surgical Pathology</i> , 2016, 40, 1177-1191.	2.1	23
101	Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , 2016, 18, 15.	2.2	88
102	Genetic basis of PD-L1 overexpression in diffuse large B-cell lymphomas. <i>Blood</i> , 2016, 127, 3026-3034.	0.6	168
103	Polyunsaturated fatty acids and prostate cancer risk: a Mendelian randomisation analysis from the PRACTICAL consortium. <i>British Journal of Cancer</i> , 2016, 115, 624-631.	2.9	23
104	Assessing the role of insulin-like growth factors and binding proteins in prostate cancer using Mendelian randomization: Genetic variants as instruments for circulating levels. <i>International Journal of Cancer</i> , 2016, 139, 1520-1533.	2.3	26
105	Blood lipids and prostate cancer: a Mendelian randomization analysis. <i>Cancer Medicine</i> , 2016, 5, 1125-1136.	1.3	68
106	Implementation of next-generation sequencing for molecular diagnosis of hereditary breast and ovarian cancer highlights its genetic heterogeneity. <i>Breast Cancer Research and Treatment</i> , 2016, 159, 245-256.	1.1	23
107	Truncating and missense <i>PPM1D</i> mutations in early-onset and/or familial/hereditary prostate cancer patients. <i>Genes Chromosomes and Cancer</i> , 2016, 55, 954-961.	1.5	15
108	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	7.7	157

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109	Inheritance of deleterious mutations at both BRCA1 and BRCA2 in an international sample of 32,295 women. <i>Breast Cancer Research</i> , 2016, 18, 112.	2.2	42
110	Prostate cancer risk regions at 8q24 and 17q24 are differentially associated with somatic <i>&gt;</i> TPRSS2:ERG</i>fusion status. <i>Human Molecular Genetics</i> , 2016, 25, ddw349.	1.4	8
111	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
112	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
113	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. <i>Nature Communications</i> , 2016, 7, 10979.	5.8	50
114	Pubertal development and prostate cancer risk: Mendelian randomization study in a population-based cohort. <i>BMC Medicine</i> , 2016, 14, 66.	2.3	42
115	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
116	Gene and pathway level analyses of germline DNA-repair gene variants and prostate cancer susceptibility using the iCOGS-genotyping array. <i>British Journal of Cancer</i> , 2016, 114, 945-952.	2.9	17
117	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv315.	3.0	77
118	Co-occurrence of nonsense mutations in MSH6 and MSH2 in Lynch syndrome families evidencing that not all truncating mutations are equal. <i>Journal of Human Genetics</i> , 2016, 61, 151-156.	1.1	8
119	The role of germline mutations in the BRCA1/2 and mismatch repair genes in men ascertained for early-onset and/or familial prostate cancer. <i>Familial Cancer</i> , 2016, 15, 111-121.	0.9	26
120	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
121	Recurrent Coding Sequence Variation Explains Only A Small Fraction of the Genetic Architecture of Colorectal Cancer. <i>Scientific Reports</i> , 2015, 5, 16286.	1.6	24
122	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369.	1.6	35
123	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2015, 17, 61.	2.2	26
124	Prediction of individual genetic risk to prostate cancer using a polygenic score. <i>Prostate</i> , 2015, 75, 1467-1474.	1.2	54
125	Assessing Associations between the AURKA-HMMR-TPX2-TUBG1 Functional Module and Breast Cancer Risk in BRCA1/2 Mutation Carriers. <i>PLoS ONE</i> , 2015, 10, e0120020.	1.1	34
126	Identification of Two Novel HOXB13 Germline Mutations in Portuguese Prostate Cancer Patients. <i>PLoS ONE</i> , 2015, 10, e0132728.	1.1	34

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127	Frequent alterations in cytoskeleton remodelling genes in primary and metastatic lung adenocarcinomas. <i>Nature Communications</i> , 2015, 6, 10131.	5.8	93
128	A Large-Scale Analysis of Genetic Variants within Putative miRNA Binding Sites in Prostate Cancer. <i>Cancer Discovery</i> , 2015, 5, 368-379.	7.7	56
129	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
130	Association of Type and Location of <i>BRCA1</i> and <i>BRCA2</i> Mutations With Risk of Breast and Ovarian Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1347.	3.8	390
131	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. <i>Cancer Causes and Control</i> , 2015, 26, 1603-1616.	0.8	77
132	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. <i>Human Molecular Genetics</i> , 2015, 24, 5589-5602.	1.4	67
133	Target gene mutational pattern in Lynch syndrome colorectal carcinomas according to tumour location and germline mutation. <i>British Journal of Cancer</i> , 2015, 113, 686-692.	2.9	30
134	Genome-Wide Association Study of Prostate Cancerâ€“Specific Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1796-1800.	1.1	27
135	The role of targeted <i>BRCA1</i> and <i>BRCA2</i> mutation analysis in hereditary breast/ovarian cancer families of Portuguese ancestry. <i>Clinical Genetics</i> , 2015, 88, 41-48.	1.0	24
136	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 1478-1492.	1.4	50
137	Candidate Genetic Modifiers for Breast and Ovarian Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 308-316.	1.1	22
138	Identification of previously unrecognized FAP in children with Gardner fibroma. <i>European Journal of Human Genetics</i> , 2015, 23, 715-718.	1.4	11
139	Uncovering potential downstream targets of oncogenic GRPR overexpression in prostate carcinomas harboring ETS rearrangements. <i>Oncoscience</i> , 2015, 2, 497-507.	0.9	11
140	Specific and redundant activities of <i>ETV1</i> and <i>ETV4</i> in prostate cancer aggressiveness revealed by co-overexpression cellular contexts. <i>Oncotarget</i> , 2015, 6, 5217-5236.	0.8	24
141	Abstract 2739: Transcontinental characterization of the Hispanic <i>BRCA1</i> 3450del4 breast cancer founder mutation. , 2015, , .		0
142	DNA Glycosylases Involved in Base Excision Repair May Be Associated with Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>PLoS Genetics</i> , 2014, 10, e1004256.	1.5	47
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306	Cytogenetic comparison of primary tumors and lymph node metastases in breast cancer patients. <i>Genes Chromosomes and Cancer</i> , 1998, 22, 122-129.	1.5	50

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