

Joan Brunet

List of Publications by Year in descending order

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Version: 2024-02-01

186
papers

7,950
citations

57758

44
h-index

69250

77
g-index

199
all docs

199
docs citations

199
times ranked

13259
citing authors

#	ARTICLE	IF	CITATIONS
1	Quality of Colonoscopy Is Associated With Adenoma Detection and Postcolonoscopy Colorectal Cancer Prevention in Lynch Syndrome. <i>Clinical Gastroenterology and Hepatology</i> , 2022, 20, 611-621.e9.	4.4	17
2	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.	6.3	19
3	Time-Dependent COVID-19 Mortality in Patients With Cancer. <i>JAMA Oncology</i> , 2022, 8, 114.	7.1	50
4	DGCR8 and the six hit, three-step model of schwannomatosis. <i>Acta Neuropathologica</i> , 2022, 143, 115-117.	7.7	10
5	Predicting Ovarian-Cancer Burden in Catalonia by 2030: An Age-Period Cohort Modelling. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 1404.	2.6	4
6	Potential Involvement of NSD1, KRT24 and ACACA in the Genetic Predisposition to Colorectal Cancer. <i>Cancers</i> , 2022, 14, 699.	3.7	0
7	Mosaicism in PTEN—new case and comment on the literature. <i>European Journal of Human Genetics</i> , 2022, 30, 641-644.	2.8	6
8	Night work, chronotype and risk of endometrial cancer in the Screenwide case-control study. <i>Occupational and Environmental Medicine</i> , 2022, , oemed-2021-108080.	2.8	6
9	Modification of BRCA1-associated breast cancer risk by HMMR overexpression. <i>Nature Communications</i> , 2022, 13, 1895.	12.8	19
10	Clinical Management of COVID-19 in Cancer Patients with the STAT3 Inhibitor Silibinin. <i>Pharmaceuticals</i> , 2022, 15, 19.	3.8	2
11	COVID-19 Sequelae and the Host Proinflammatory Response: An Analysis From the OnCovid Registry. <i>Journal of the National Cancer Institute</i> , 2022, 114, 979-987.	6.3	14
12	Persistence of long-term COVID-19 sequelae in patients with cancer: An analysis from the OnCovid registry. <i>European Journal of Cancer</i> , 2022, 170, 10-16.	2.8	11
13	Vaccination against SARS-CoV-2 protects from morbidity, mortality and sequelae from COVID19 in patients with cancer. <i>European Journal of Cancer</i> , 2022, 171, 64-74.	2.8	19
14	Outcomes of the SARS-CoV-2 omicron (B.1.1.529) variant outbreak among vaccinated and unvaccinated patients with cancer in Europe: results from the retrospective, multicentre, OnCovid registry study. <i>Lancet Oncology</i> , The, 2022, 23, 865-875.	10.7	50
15	An Integrated Approach for the Early Detection of Endometrial and Ovarian Cancers (Screenwide) Tj ETQq1 1 0.784314 rgBT /Overlo	2.5	6
16	<i>TP53</i> , a gene for colorectal cancer predisposition in the absence of Li-Fraumeni-associated phenotypes. <i>Gut</i> , 2021, 70, 1139-1146.	12.1	10
17	Predicting the rising incidence and mortality of endometrial cancers among women aged 65-74 years in Catalonia. <i>Maturitas</i> , 2021, 144, 11-15.	2.4	2
18	Characteristics of Adrenocortical Carcinoma Associated With Lynch Syndrome. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021, 106, 318-325.	3.6	20

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19	Specialist palliative and end-of-life care for patients with cancer and SARS-CoV-2 infection: a European perspective. <i>Therapeutic Advances in Medical Oncology</i> , 2021, 13, 175883592110422.	3.2	4
20	BARD1 Pathogenic Variants Are Associated with Triple-Negative Breast Cancer in a Spanish Hereditary Breast and Ovarian Cancer Cohort. <i>Genes</i> , 2021, 12, 150.	2.4	11
21	Systemic pro-inflammatory response identifies patients with cancer with adverse outcomes from SARS-CoV-2 infection: the OnCovid Inflammatory Score. , 2021, 9, e002277.		30
22	Patientsâ€™ and professionalsâ€™ perspective of non-in-person visits in hereditary cancer: predictors and impact of the COVID-19 pandemic. <i>Genetics in Medicine</i> , 2021, 23, 1450-1457.	2.4	1
23	Response to letter entitled: Re: ERCC3 a new ovarian cancer susceptibility gene?. <i>European Journal of Cancer</i> , 2021, 150, 281-282.	2.8	0
24	Determinants of enhanced vulnerability to coronavirus disease 2019 in UK patients with cancer: a European study. <i>European Journal of Cancer</i> , 2021, 150, 190-202.	2.8	37
25	No Difference in Penetrance between Truncating and Missense/Aberrant Splicing Pathogenic Variants in MLH1 and MSH2: A Prospective Lynch Syndrome Database Study. <i>Journal of Clinical Medicine</i> , 2021, 10, 2856.	2.4	11
26	Non-Lynch Familial and Early-Onset Colorectal Cancer Explained by Accumulation of Low-Risk Genetic Variants. <i>Cancers</i> , 2021, 13, 3857.	3.7	8
27	Altered regulation of <i>BRCA1</i> exon 11 splicing is associated with breast cancer risk in carriers of <i>BRCA1</i> pathogenic variants. <i>Human Mutation</i> , 2021, 42, 1488-1502.	2.5	7
28	Paired Somatic-Germline Testing of 15 Polyposis and Colorectal Cancerâ€“Predisposing Genes Highlights the Role of APC Mosaicism in de Novo Familial Adenomatous Polyposis. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 1452-1459.	2.8	10
29	Sensitivity of cervical cytology in endometrial cancer detection in a tertiary hospital in Spain. <i>Cancer Medicine</i> , 2021, 10, 6762-6766.	2.8	6
30	Prevalence and impact of COVID-19 sequelae on treatment and survival of patients with cancer who recovered from SARS-CoV-2 infection: evidence from the OnCovid retrospective, multicentre registry study. <i>Lancet Oncology</i> , The, 2021, 22, 1669-1680.	10.7	73
31	RNA assay identifies a previous misclassification of BARD1 c.1977A>G variant. <i>Scientific Reports</i> , 2021, 11, 22948.	3.3	0
32	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.	0.9	39
33	The â€œPsychosocial Aspects in Hereditary Cancerâ€–questionnaire in women attending breast cancer genetic clinics: Psychometric validation across Frenchâ€, Germanâ€–and Spanishâ€–language versions. <i>European Journal of Cancer Care</i> , 2020, 29, e13173.	1.5	2
34	Complete Loss of EPCAM Immunoexpression Identifies EPCAM Deletion Carriers in MSH2-Negative Colorectal Neoplasia. <i>Cancers</i> , 2020, 12, 2803.	3.7	4
35	Presenting Features and Early Mortality from SARS-CoV-2 Infection in Cancer Patients during the Initial Stage of the COVID-19 Pandemic in Europe. <i>Cancers</i> , 2020, 12, 1841.	3.7	58
36	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	2.4	82

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37	Comprehensive Constitutional Genetic and Epigenetic Characterization of Lynch-Like Individuals. <i>Cancers</i> , 2020, 12, 1799.	3.7	15
38	Primary breast cancer and health related quality of life in Spanish women: The EpiGEICAM case-control study. <i>Scientific Reports</i> , 2020, 10, 7741.	3.3	9
39	Screening of CNVs using NGS data improves mutation detection yield and decreases costs in genetic testing for hereditary cancer. <i>Journal of Medical Genetics</i> , 2020, , jmedgenet-2020-107366.	3.2	3
40	Assessing Effectiveness of Colonic and Gynecological Risk Reducing Surgery in Lynch Syndrome Individuals. <i>Cancers</i> , 2020, 12, 3419.	3.7	11
41	Clinical Portrait of the SARS-CoV-2 Epidemic in European Patients with Cancer. <i>Cancer Discovery</i> , 2020, 10, 1465-1474.	9.4	151
42	Immune Cell Associations with Cancer Risk. <i>IScience</i> , 2020, 23, 101296.	4.1	6
43	Role of POLE and POLD1 in familial cancer. <i>Genetics in Medicine</i> , 2020, 22, 2089-2100.	2.4	76
44	ERCC3, a new ovarian cancer susceptibility gene?. <i>European Journal of Cancer</i> , 2020, 141, 1-8.	2.8	8
45	Comprehensive analysis and ACMG-based classification of <i>CHEK2</i> variants in hereditary cancer patients. <i>Human Mutation</i> , 2020, 41, 2128-2142.	2.5	10
46	High-sensitivity microsatellite instability assessment for the detection of mismatch repair defects in normal tissue of biallelic germline mismatch repair mutation carriers. <i>Journal of Medical Genetics</i> , 2020, 57, 269-273.	3.2	20
47	Evaluation of CNV detection tools for NGS panel data in genetic diagnostics. <i>European Journal of Human Genetics</i> , 2020, 28, 1645-1655.	2.8	67
48	Silibinin and SARS-CoV-2: Dual Targeting of Host Cytokine Storm and Virus Replication Machinery for Clinical Management of COVID-19 Patients. <i>Journal of Clinical Medicine</i> , 2020, 9, 1770.	2.4	42
49	Sensitivity of cervico-vaginal cytology in endometrial carcinoma: A systematic review and meta-analysis. <i>Cancer Cytopathology</i> , 2020, 128, 792-802.	2.4	23
50	Tumors defective in homologous recombination rely on oxidative metabolism: relevance to treatments with PARP inhibitors. <i>EMBO Molecular Medicine</i> , 2020, 12, e11217.	6.9	37
51	High Prevalence of Somatic Oncogenic Driver Alterations in Patients With NSCLC and Li-Fraumeni Syndrome. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1232-1239.	1.1	29
52	The Spectrum of FANCM Protein Truncating Variants in European Breast Cancer Cases. <i>Cancers</i> , 2020, 12, 292.	3.7	11
53	Association of premenopausal risk-reducing salpingo-oophorectomy with breast cancer risk in BRCA1/2 mutation carriers: Maximising bias-reduction. <i>European Journal of Cancer</i> , 2020, 132, 53-60.	2.8	16
54	Exploring the Role of Mutations in Fanconi Anemia Genes in Hereditary Cancer Patients. <i>Cancers</i> , 2020, 12, 829.	3.7	41

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55	Resveratrol targets PD-L1 glycosylation and dimerization to enhance antitumor T-cell immunity. <i>Aging</i> , 2020, 12, 8-34.	3.1	99
56	The LSD1 inhibitor iadademstat (ORY-1001) targets SOX2-driven breast cancer stem cells: a potential epigenetic therapy in luminal-B and HER2-positive breast cancer subtypes. <i>Aging</i> , 2020, 12, 4794-4814.	3.1	38
57	Improving Genetic Testing in Hereditary Cancer by RNA Analysis. <i>Journal of Molecular Diagnostics</i> , 2020, 22, 1453-1468.	2.8	9
58	Psychosocial problems in women attending French, German and Spanish genetics clinics before and after targeted or multigene testing results: an observational prospective study. <i>BMJ Open</i> , 2019, 9, e029926.	1.9	9
59	Germline variation in O6-methylguanine-DNA methyltransferase (MGMT) as cause of hereditary colorectal cancer. <i>Cancer Letters</i> , 2019, 447, 86-92.	7.2	12
60	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
61	Large scale multifactorial likelihood quantitative analysis of <i>BRCA1</i> and <i>BRCA2</i> variants: An ENIGMA resource to support clinical variant classification. <i>Human Mutation</i> , 2019, 40, 1557-1578.	2.5	102
62	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	6.4	19
63	<i>NTHL1</i> biallelic mutations seldom cause colorectal cancer, serrated polyposis or a multi-tumor phenotype, in absence of colorectal adenomas. <i>Scientific Reports</i> , 2019, 9, 9020.	3.3	23
64	Contribution to colonic polyposis of recently proposed predisposing genes and assessment of the prevalence of <i>NTHL1</i> and <i>MSH3</i> associated polyposes. <i>Human Mutation</i> , 2019, 40, 1910-1923. ^{2,5}		24
65	Defining a mutational signature for endometrial cancer screening and early detection. <i>Cancer Epidemiology</i> , 2019, 61, 129-132.	1.9	7
66	Computational de-orphanization of the olive oil biophenol oleacein: Discovery of new metabolic and epigenetic targets. <i>Food and Chemical Toxicology</i> , 2019, 131, 110529.	3.6	15
67	New perspectives on screening and early detection of endometrial cancer. <i>International Journal of Cancer</i> , 2019, 145, 3194-3206.	5.1	58
68	The C Allele of <i>ATM</i> rs11212617 Associates With Higher Pathological Complete Remission Rate in Breast Cancer Patients Treated With Neoadjuvant Metformin. <i>Frontiers in Oncology</i> , 2019, 9, 193.	2.8	17
69	Opportunistic testing of <i>BRCA1</i> , <i>BRCA2</i> and mismatch repair genes improves the yield of phenotype driven hereditary cancer gene panels. <i>International Journal of Cancer</i> , 2019, 145, 2682-2691.	5.1	30
70	The extra virgin olive oil phenolic oleacein is a dual substrate-inhibitor of catechol-O-methyltransferase. <i>Food and Chemical Toxicology</i> , 2019, 128, 35-45.	3.6	27
71	Neoadjuvant Metformin Added to Systemic Therapy Decreases the Proliferative Capacity of Residual Breast Cancer. <i>Journal of Clinical Medicine</i> , 2019, 8, 2180.	2.4	12
72	Highly sensitive <i>MLH1</i> methylation analysis in blood identifies a cancer patient with low-level mosaic <i>MLH1</i> epimutation. <i>Clinical Epigenetics</i> , 2019, 11, 171.	4.1	7

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73	Does multilocus inherited neoplasia alleles syndrome have severe clinical expression?. Journal of Medical Genetics, 2019, 56, 521-525.	3.2	11
74	Novel <i>POLE</i> pathogenic germline variant in a family with multiple primary tumors results in distinct mutational signatures. Human Mutation, 2019, 40, 36-41.	2.5	21
75	Height and Body Mass Index as Modifiers of Breast Cancer Risk in <i>BRCA1</i> / <i>BRCA2</i> Mutation Carriers: A Mendelian Randomization Study. Journal of the National Cancer Institute, 2019, 111, 350-364.	6.3	30
76	Metformin induces a fasting- and antifolate-mimicking modification of systemic host metabolism in breast cancer patients. Aging, 2019, 11, 2874-2888.	3.1	25
77	Decapping protein EDC4 regulates DNA repair and phenocopies <i>BRCA1</i> . Nature Communications, 2018, 9, 967.	12.8	33
78	Evidence suggests that germline <i>RNF43</i> mutations are a rare cause of serrated polyposis. Gut, 2018, 67, 2230-2232.	12.1	48
79	Association Between Germline Mutations in <i>BRF1</i> , a Subunit of the RNA Polymerase III Transcription Complex, and Hereditary Colorectal Cancer. Gastroenterology, 2018, 154, 181-194.e20.	1.3	32
80	Metformin Is a Direct <i>SIRT1</i> -Activating Compound: Computational Modeling and Experimental Validation. Frontiers in Endocrinology, 2018, 9, 657.	3.5	85
81	Surveillance Guidelines for Hereditary Colorectal Cancer Syndromes. , 2018, , 305-326.		0
82	Germline mutations in the spindle assembly checkpoint genes <i>BUB1</i> and <i>BUB3</i> are infrequent in familial colorectal cancer and polyposis. Molecular Cancer, 2018, 17, 23.	19.2	19
83	Elucidating the molecular basis of <i>MSH2</i> -deficient tumors by combined germline and somatic analysis. International Journal of Cancer, 2017, 141, 1365-1380.	5.1	26
84	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	21.4	356
85	Benchmarking of Whole Exome Sequencing and Ad Hoc Designed Panels for Genetic Testing of Hereditary Cancer. Scientific Reports, 2017, 7, 37984.	3.3	35
86	A comprehensive custom panel design for routine hereditary cancer testing: preserving control, improving diagnostics and revealing a complex variation landscape. Scientific Reports, 2017, 7, 39348.	3.3	45
87	Stem cell-like transcriptional reprogramming mediates metastatic resistance to <i>mTOR</i> inhibition. Oncogene, 2017, 36, 2737-2749.	5.9	34
88	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	21.4	289
89	Attenuation of RNA polymerase II pausing mitigates <i>BRCA1</i> -associated R-loop accumulation and tumorigenesis. Nature Communications, 2017, 8, 15908.	12.8	118
90	Prediction of Breast and Prostate Cancer Risks in Male <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers Using Polygenic Risk Scores. Journal of Clinical Oncology, 2017, 35, 2240-2250.	1.6	152

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91	<i>BRCA1</i> haploinsufficiency cell-autonomously activates RANKL expression and generates denosumab-responsive breast cancer-initiating cells. <i>Oncotarget</i> , 2017, 8, 35019-35032.	1.8	12
92	Fatty acid synthase expression and its association with clinico-histopathological features in triple-negative breast cancer. <i>Oncotarget</i> , 2017, 8, 74391-74405.	1.8	40
93	Ethics competences in the undergraduate medical education curriculum: the Spanish experience. <i>Croatian Medical Journal</i> , 2016, 57, 493-503.	0.7	11
94	Response of brain metastasis from lung cancer patients to an oral nutraceutical product containing silibinin. <i>Oncotarget</i> , 2016, 7, 32006-32014.	1.8	47
95	Somatic <i>DICER1</i> mutations in adult-onset pulmonary blastoma. <i>European Respiratory Journal</i> , 2016, 47, 1879-1882.	6.7	22
96	An international survey of surveillance schemes for unaffected BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2016, 157, 319-327.	2.5	26
97	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	12.8	78
98	Scarce evidence of the causal role of germline mutations in UNC5C in hereditary colorectal cancer and polyposis. <i>Scientific Reports</i> , 2016, 6, 20697.	3.3	9
99	Investigating the effect of 28 BRCA1 and BRCA2 mutations on their related transcribed mRNA. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 253-260.	2.5	6
100	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	21.4	125
101	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.	6.3	77
102	POLE and POLD1 mutations in 529 kindred with familial colorectal cancer and/or polyposis: review of reported cases and recommendations for genetic testing and surveillance. <i>Genetics in Medicine</i> , 2016, 18, 325-332.	2.4	209
103	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	1.4	18
104	Synthetic lethal interaction of cetuximab with MEK1/2 inhibition in <i>NRAS</i>-mutant metastatic colorectal cancer. <i>Oncotarget</i> , 2016, 7, 82185-82199.	1.8	16
105	Mammographic density and breast cancer in women from high risk families. <i>Breast Cancer Research</i> , 2015, 17, 93.	5.0	22
106	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.	2.5	34
107	SEOM clinical guidelines in Hereditary Breast and ovarian cancer. <i>Clinical and Translational Oncology</i> , 2015, 17, 956-961.	2.4	82
108	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	21.4	221

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109	Germline Mutations in FAN1 Cause Hereditary Colorectal Cancer by Impairing DNA Repair. <i>Gastroenterology</i> , 2015, 149, 563-566.	1.3	94
110	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.	2.5	22
111	Integrating gene expression and epidemiological data for the discovery of genetic interactions associated with cancer risk. <i>Carcinogenesis</i> , 2014, 35, 578-585.	2.8	1
112	VAV3 mediates resistance to breast cancer endocrine therapy. <i>Breast Cancer Research</i> , 2014, 16, R53.	5.0	28
113	GALNT12 is Not a Major Contributor of Familial Colorectal Cancer Type X. <i>Human Mutation</i> , 2014, 35, 50-52.	2.5	22
114	New insights into POLE and POLD1 germline mutations in familial colorectal cancer and polyposis. <i>Human Molecular Genetics</i> , 2014, 23, 3506-3512.	2.9	135
115	Successful empirical erlotinib treatment of a mechanically ventilated patient newly diagnosed with metastatic lung adenocarcinoma. <i>Lung Cancer</i> , 2014, 86, 102-104.	2.0	27
116	Prevalence of germline MUTYH mutations among Lynch-like syndrome patients. <i>European Journal of Cancer</i> , 2014, 50, 2241-2250.	2.8	66
117	Silibinin administration improves hepatic failure due to extensive liver infiltration in a breast cancer patient. <i>Anticancer Research</i> , 2014, 34, 4323-7.	1.1	21
118	Refining the role of <i>pms2</i> in Lynch syndrome: germline mutational analysis improved by comprehensive assessment of variants. <i>Journal of Medical Genetics</i> , 2013, 50, 552-563.	3.2	47
119	Next-generation sequencing meets genetic diagnostics: development of a comprehensive workflow for the analysis of <i>BRCA1</i> and <i>BRCA2</i> genes. <i>European Journal of Human Genetics</i> , 2013, 21, 864-870.	2.8	94
120	Genome-Wide Association Study in <i>BRCA1</i> Mutation Carriers Identifies Novel Loci Associated with Breast and Ovarian Cancer Risk. <i>PLoS Genetics</i> , 2013, 9, e1003212.	3.5	244
121	Analysis of <i>PALB2</i> Gene in <i>BRCA1/BRCA2</i> Negative Spanish Hereditary Breast/Ovarian Cancer Families with Pancreatic Cancer Cases. <i>PLoS ONE</i> , 2013, 8, e67538.	2.5	44
122	Functional and Structural Analysis of C-Terminal <i>BRCA1</i> Missense Variants. <i>PLoS ONE</i> , 2013, 8, e61302.	2.5	16
123	Cross-suppression of EGFR ligands amphiregulin and epiregulin and de-repression of FGFR3 signalling contribute to cetuximab resistance in wild-type <i>KRAS</i> tumour cells. <i>British Journal of Cancer</i> , 2012, 106, 1406-1414.	6.4	42
124	Common Variants at the 19p13.1 and <i>ZNF365</i> Loci Are Associated with ER Subtypes of Breast Cancer and Ovarian Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 645-657.	2.5	47
125	<i>MLH1</i> promoter hypermethylation in the analytical algorithm of Lynch syndrome: a cost-effectiveness study. <i>European Journal of Human Genetics</i> , 2012, 20, 762-768.	2.8	76
126	Pathology of Breast and Ovarian Cancers among <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Results from the Consortium of Investigators of Modifiers of <i>BRCA1/2</i> (CIMBA). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 134-147.	2.5	513

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127	Transcriptional upregulation of HER2 expression in the absence of HER2 gene amplification results in cetuximab resistance that is reversed by trastuzumab treatment. <i>Oncology Reports</i> , 2012, 27, 1887-92.	2.6	5
128	MLH1 methylation screening is effective in identifying epimutation carriers. <i>European Journal of Human Genetics</i> , 2012, 20, 1256-1264.	2.8	36
129	Cancer develops, progresses and responds to therapies through restricted perturbation of the protein-protein interaction network. <i>Integrative Biology (United Kingdom)</i> , 2012, 4, 1038.	1.3	10
130	What factors may influence psychological well being at three months and one year post BRCA genetic result disclosure?. <i>Breast</i> , 2012, 21, 755-760.	2.2	36
131	Comprehensive functional assessment of <i>MLH1</i> variants of unknown significance. <i>Human Mutation</i> , 2012, 33, 1576-1588.	2.5	30
132	Assessing the RNA effect of 26 DNA variants in the BRCA1 and BRCA2 genes. <i>Breast Cancer Research and Treatment</i> , 2012, 132, 979-992.	2.5	20
133	Analysis of SLX4/FANCP in non-BRCA1/2-mutated breast cancer families. <i>BMC Cancer</i> , 2012, 12, 84.	2.6	14
134	Detection of a large rearrangement in PALB2 in Spanish breast cancer families with male breast cancer. <i>Breast Cancer Research and Treatment</i> , 2012, 132, 307-315.	2.5	50
135	Evolution of the predictive markers amphiregulin and epiregulin mRNAs during long-term cetuximab treatment of KRAS wild-type tumor cells. <i>Investigational New Drugs</i> , 2012, 30, 846-852.	2.6	11
136	Neuroendocrine tumors: A population-based study of incidence and survival in Girona province, 1994-2004. <i>Cancer Epidemiology</i> , 2011, 35, e49-e54.	1.9	20
137	Inhibitor of Apoptosis (IAP) survivin is indispensable for survival of HER2 gene-amplified breast cancer cells with primary resistance to HER1/2-targeted therapies. <i>Biochemical and Biophysical Research Communications</i> , 2011, 407, 412-419.	2.1	44
138	Exploring the link between MORF4L1 and risk of breast cancer. <i>Breast Cancer Research</i> , 2011, 13, R40.	5.0	23
139	Interferon/STAT1 and neuregulin signaling pathways are exploratory biomarkers of cetuximab (Erbix) efficacy in KRAS wild-type squamous carcinomas: A pathway-based analysis of whole human-genome microarray data from cetuximab-adapted tumor cell-line models. <i>International Journal of Oncology</i> , 2011, 39, 1455-79.	3.3	15
140	Costs and Ethical Issues Related to First-Line Treatment of Metastatic Non-Small-Cell Lung Cancer: Considerations From a Public Healthcare System Perspective. <i>Clinical Lung Cancer</i> , 2011, 12, 335-340.	2.6	2
141	eXIT*CBR: A framework for case-based medical diagnosis development and experimentation. <i>Artificial Intelligence in Medicine</i> , 2011, 51, 81-91.	6.5	31
142	Evidence for a link between TNFRSF11A and risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011, 129, 947-954.	2.5	12
143	Identification of a new complex rearrangement affecting exon 20 of BRCA1. <i>Breast Cancer Research and Treatment</i> , 2011, 130, 341-344.	2.5	3
144	SEOM clinical guidelines for hereditary cancer. <i>Clinical and Translational Oncology</i> , 2011, 13, 580-586.	2.4	13

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