

Francesca M Buffa

List of Publications by Year in descending order

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155
papers

10,943
citations

30068

54
h-index

32838

100
g-index

171
all docs

171
docs citations

171
times ranked

21118
citing authors

#	ARTICLE	IF	CITATIONS
1	hsa-miR-210 Is Induced by Hypoxia and Is an Independent Prognostic Factor in Breast Cancer. <i>Clinical Cancer Research</i> , 2008, 14, 1340-1348.	7.0	617
2	Direct targeting of Sec23a by miR-200s influences cancer cell secretome and promotes metastatic colonization. <i>Nature Medicine</i> , 2011, 17, 1101-1108.	30.7	552
3	Fatty Acid Uptake and Lipid Storage Induced by HIF-1 α Contribute to Cell Growth and Survival after Hypoxia-Reoxygenation. <i>Cell Reports</i> , 2014, 9, 349-365.	6.4	498
4	Large meta-analysis of multiple cancers reveals a common, compact and highly prognostic hypoxia metagene. <i>British Journal of Cancer</i> , 2010, 102, 428-435.	6.4	440
5	miR-182-Mediated Downregulation of BRCA1 Impacts DNA Repair and Sensitivity to PARP Inhibitors. <i>Molecular Cell</i> , 2011, 41, 210-220.	9.7	409
6	Relation of a Hypoxia Metagene Derived from Head and Neck Cancer to Prognosis of Multiple Cancers. <i>Cancer Research</i> , 2007, 67, 3441-3449.	0.9	349
7	Regulation of autophagy by ATF4 in response to severe hypoxia. <i>Oncogene</i> , 2010, 29, 4424-4435.	5.9	320
8	Glucose Utilization via Glycogen Phosphorylase Sustains Proliferation and Prevents Premature Senescence in Cancer Cells. <i>Cell Metabolism</i> , 2012, 16, 751-764.	16.2	320
9	microRNA-Associated Progression Pathways and Potential Therapeutic Targets Identified by Integrated mRNA and microRNA Expression Profiling in Breast Cancer. <i>Cancer Research</i> , 2011, 71, 5635-5645.	0.9	285
10	MicroRNA-210 Regulates Mitochondrial Free Radical Response to Hypoxia and Krebs Cycle in Cancer Cells by Targeting Iron Sulfur Cluster Protein ISCU. <i>PLoS ONE</i> , 2010, 5, e10345.	2.5	276
11	Tumor hypoxia induces nuclear paraspeckle formation through HIF-2 α dependent transcriptional activation of NEAT1 leading to cancer cell survival. <i>Oncogene</i> , 2015, 34, 4482-4490.	5.9	245
12	The small-nucleolar RNAs commonly used for microRNA normalisation correlate with tumour pathology and prognosis. <i>British Journal of Cancer</i> , 2011, 104, 1168-1177.	6.4	244
13	A Core Human Primary Tumor Angiogenesis Signature Identifies the Endothelial Orphan Receptor ELTD1 as a Key Regulator of Angiogenesis. <i>Cancer Cell</i> , 2013, 24, 229-241.	16.8	238
14	hsa-miR-210 is a marker of tumor hypoxia and a prognostic factor in head and neck cancer. <i>Cancer</i> , 2010, 116, 2148-2158.	4.1	215
15	A 26-Gene Hypoxia Signature Predicts Benefit from Hypoxia-Modifying Therapy in Laryngeal Cancer but Not Bladder Cancer. <i>Clinical Cancer Research</i> , 2013, 19, 4879-4888.	7.0	214
16	Hypoxia promotes stem cell phenotypes and poor prognosis through epigenetic regulation of DICER. <i>Nature Communications</i> , 2014, 5, 5203.	12.8	195
17	Translation reprogramming is an evolutionarily conserved driver of phenotypic plasticity and therapeutic resistance in melanoma. <i>Genes and Development</i> , 2017, 31, 18-33.	5.9	184
18	The anti-malarial atovaquone increases radiosensitivity by alleviating tumour hypoxia. <i>Nature Communications</i> , 2016, 7, 12308.	12.8	173

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19	The Histone Demethylase JMJD2B Is Regulated by Estrogen Receptor $\hat{\pm}$ and Hypoxia, and Is a Key Mediator of Estrogen Induced Growth. <i>Cancer Research</i> , 2010, 70, 6456-6466.	0.9	167
20	Assessment of tumour hypoxia for prediction of response to therapy and cancer prognosis. <i>Journal of Cellular and Molecular Medicine</i> , 2010, 14, 18-29.	3.6	143
21	Integrated analysis of microRNA and mRNA expression and association with HIF binding reveals the complexity of microRNA expression regulation under hypoxia. <i>Molecular Cancer</i> , 2014, 13, 28.	19.2	135
22	MicroRNA-10b and breast cancer metastasis. <i>Nature</i> , 2008, 455, E8-E9.	27.8	134
23	Development and Validation of a 28-gene Hypoxia-related Prognostic Signature for Localized Prostate Cancer. <i>EBioMedicine</i> , 2018, 31, 182-189.	6.1	132
24	Transcriptional up-regulation of ULK1 by ATF4 contributes to cancer cell survival. <i>Biochemical Journal</i> , 2013, 449, 389-400.	3.7	128
25	Clinical whole-genome sequencing from routine formalin-fixed, paraffin-embedded specimens: pilot study for the 100,000 Genomes Project. <i>Genetics in Medicine</i> , 2018, 20, 1196-1205.	2.4	125
26	Phosphorylated ER $\hat{\pm}$, HIF-1 $\hat{\pm}$, and MAPK Signaling As Predictors of Primary Endocrine Treatment Response and Resistance in Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2009, 27, 227-234.	1.6	116
27	Hypoxia-induced p53 modulates both apoptosis and radiosensitivity via AKT. <i>Journal of Clinical Investigation</i> , 2015, 125, 2385-2398.	8.2	111
28	Estrogen receptor- $\hat{\pm}$ directly regulates the hypoxia-inducible factor 1 pathway associated with antiestrogen response in breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 15172-15177.	7.1	110
29	Pan-cancer characterisation of microRNA across cancer hallmarks reveals microRNA-mediated downregulation of tumour suppressors. <i>Nature Communications</i> , 2018, 9, 5228.	12.8	110
30	A Gene Signature for Selecting Benefit from Hypoxia Modification of Radiotherapy for High-Risk Bladder Cancer Patients. <i>Clinical Cancer Research</i> , 2017, 23, 4761-4768.	7.0	107
31	miR-139-5p Modulates Radiotherapy Resistance in Breast Cancer by Repressing Multiple Gene Networks of DNA Repair and ROS Defense. <i>Cancer Research</i> , 2018, 78, 501-515.	0.9	105
32	Overexpression of <i>POLQ</i> Confers a Poor Prognosis in Early Breast Cancer Patients. <i>Oncotarget</i> , 2010, 1, 175-184.	1.8	100
33	Gene Expression Signatures as Biomarkers of Tumour Hypoxia. <i>Clinical Oncology</i> , 2015, 27, 547-560.	1.4	95
34	Targeting tumour hypoxia to prevent cancer metastasis. From biology, biosensing and technology to drug development: the METOXIA consortium. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2015, 30, 689-721.	5.2	93
35	Integrated Pharmacodynamic Analysis Identifies Two Metabolic Adaption Pathways to Metformin in Breast Cancer. <i>Cell Metabolism</i> , 2018, 28, 679-688.e4.	16.2	92
36	Epigenetic downregulation of human disabled homolog 2 switches TGF- $\hat{1}^2$ from a tumor suppressor to a tumor promoter. <i>Journal of Clinical Investigation</i> , 2010, 120, 2842-2857.	8.2	87

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37	Hypoxia-induced switch in SNAT2/SLC38A2 regulation generates endocrine resistance in breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12452-12461.	7.1	86
38	Molecular Marker Profiles Predict Locoregional Control of Head and Neck Squamous Cell Carcinoma in a Randomized Trial of Continuous Hyperfractionated Accelerated Radiotherapy. <i>Clinical Cancer Research</i> , 2004, 10, 3745-3754.	7.0	83
39	Differential clonal evolution in oesophageal cancers in response to neo-adjuvant chemotherapy. <i>Nature Communications</i> , 2016, 7, 11111.	12.8	83
40	Human CHCHD4 mitochondrial proteins regulate cellular oxygen consumption rate and metabolism and provide a critical role in hypoxia signaling and tumor progression. <i>Journal of Clinical Investigation</i> , 2012, 122, 600-611.	8.2	82
41	Alternate RASSF1 Transcripts Control SRC Activity, E-Cadherin Contacts, and YAP-Mediated Invasion. <i>Current Biology</i> , 2015, 25, 3019-3034.	3.9	74
42	Gene expression and hypoxia in breast cancer. <i>Genome Medicine</i> , 2011, 3, 55.	8.2	73
43	FANCD2 limits replication stress and genome instability in cells lacking BRCA2. <i>Nature Structural and Molecular Biology</i> , 2016, 23, 755-757.	8.2	73
44	Spectral Clustering of Microarray Data Elucidates the Roles of Microenvironment Remodeling and Immune Responses in Survival of Head and Neck Squamous Cell Carcinoma. <i>Journal of Clinical Oncology</i> , 2010, 28, 2881-2888.	1.6	72
45	RASSF1A uncouples Wnt from Hippo signalling and promotes YAP mediated differentiation via p73. <i>Nature Communications</i> , 2018, 9, 424.	12.8	72
46	Hypoxia induces a lipogenic cancer cell phenotype via HIF1 α -dependent and -independent pathways. <i>Oncotarget</i> , 2015, 6, 1920-1941.	1.8	72
47	Genomic alterations underlie a pan-cancer metabolic shift associated with tumour hypoxia. <i>Genome Biology</i> , 2016, 17, 140.	8.8	67
48	Dichloroacetate reverses the hypoxic adaptation to bevacizumab and enhances its antitumor effects in mouse xenografts. <i>Journal of Molecular Medicine</i> , 2013, 91, 749-758.	3.9	64
49	Robust prognostic value of a knowledge-based proliferation signature across large patient microarray studies spanning different cancer types. <i>British Journal of Cancer</i> , 2008, 99, 1884-1890.	6.4	62
50	Dosimetric impact of computed tomography calibration on a commercial treatment planning system for external radiation therapy. <i>Radiotherapy and Oncology</i> , 1998, 48, 335-338.	0.6	60
51	High activity Rhenium-186 HEDP with autologous peripheral blood stem cell rescue: a phase I study in progressive hormone refractory prostate cancer metastatic to bone. <i>British Journal of Cancer</i> , 2002, 86, 1715-1720.	6.4	57
52	Hypoxia regulates FGFR3 expression via HIF-1 α and miR-100 and contributes to cell survival in non-muscle invasive bladder cancer. <i>British Journal of Cancer</i> , 2013, 109, 50-59.	6.4	55
53	Incorporating biologic measurements (SF2, CFE) into a tumor control probability model increases their prognostic significance: a study in cervical carcinoma treated with radiation therapy. <i>International Journal of Radiation Oncology Biology Physics</i> , 2001, 50, 1113-1122.	0.8	54
54	Development of a Tissue Array for Primary Melanoma with Long-Term Follow-Up: Discovering Melanoma Cell Adhesion Molecule as an Important Prognostic Marker. <i>Plastic and Reconstructive Surgery</i> , 2005, 115, 367-375.	1.4	53

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55	Phase I/II Trial of Bevacizumab and Radiotherapy for Locally Advanced Inoperable Colorectal Cancer: Vasculature-Independent Radiosensitizing Effect of Bevacizumab. <i>Clinical Cancer Research</i> , 2009, 15, 7069-7076.	7.0	52
56	<sc>SCF</sc> (Fbxl17) ubiquitylation of Sufu regulates Hedgehog signaling and medulloblastoma development. <i>EMBO Journal</i> , 2016, 35, 1400-1416.	7.8	50
57	Gemcitabine-Induced TIMP1 Attenuates Therapy Response and Promotes Tumor Growth and Liver Metastasis in Pancreatic Cancer. <i>Cancer Research</i> , 2017, 77, 5952-5962.	0.9	50
58	Molecular profiles as predictive marker for the effect of overall treatment time of radiotherapy in supraglottic larynx squamous cell carcinomas. <i>Radiotherapy and Oncology</i> , 2004, 72, 275-282.	0.6	46
59	Nutritional Stress Induced by Tryptophan-Degrading Enzymes Results in ATF4-Dependent Reprogramming of the Amino Acid Transporter Profile in Tumor Cells. <i>Cancer Research</i> , 2016, 76, 6193-6204.	0.9	45
60	Assessing Early Therapeutic Response to Bevacizumab in Primary Breast Cancer Using Magnetic Resonance Imaging and Gene Expression Profiles. <i>Journal of the National Cancer Institute Monographs</i> , 2011, 2011, 71-74.	2.1	42
61	Selective Targeting of Bromodomains of the Bromodomain-PHD Fingers Family Impairs Osteoclast Differentiation. <i>ACS Chemical Biology</i> , 2017, 12, 2619-2630.	3.4	41
62	MITF controls the TCA cycle to modulate the melanoma hypoxia response. <i>Pigment Cell and Melanoma Research</i> , 2019, 32, 792-808.	3.3	41
63	E2F1 proteolysis via <sc>SCF</sc> cyclin F underlies synthetic lethality between cyclin F loss and Chk1 inhibition. <i>EMBO Journal</i> , 2019, 38, e101443.	7.8	40
64	Mitochondrial Inhibitor Atovaquone Increases Tumor Oxygenation and Inhibits Hypoxic Gene Expression in Patients with Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 2459-2469.	7.0	40
65	Carbonic anhydrase IX induction defines a heterogeneous cancer cell response to hypoxia and mediates stem cell-like properties and sensitivity to HDAC inhibition. <i>Oncotarget</i> , 2015, 6, 19413-19427.	1.8	39
66	Intensity-modulated Radiotherapy Allows Escalation of the Radiation Dose to the Pelvic Lymph Nodes in Patients with Locally Advanced Prostate Cancer: Preliminary Results of a Phase I Dose Escalation Study. <i>Clinical Oncology</i> , 2010, 22, 236-244.	1.4	38
67	Radiation response and cure rate of human colon adenocarcinoma spheroids of different size: the significance of hypoxia on tumor control modelling. <i>International Journal of Radiation Oncology Biology Physics</i> , 2001, 49, 1109-1118.	0.8	37
68	Prospective technical validation and assessment of intra-tumour heterogeneity of a low density array hypoxia gene profile in head and neck squamous cell carcinoma. <i>European Journal of Cancer</i> , 2013, 49, 156-165.	2.8	36
69	Hypoxia-driven cell motility reflects the interplay between JMY and HIF-1 β . <i>Oncogene</i> , 2011, 30, 4835-4842.	5.9	35
70	IGF-1R associates with adverse outcomes after radical radiotherapy for prostate cancer. <i>British Journal of Cancer</i> , 2017, 117, 1600-1606.	6.4	35
71	Role of Delta-like 4 in Jagged1-induced tumour angiogenesis and tumour growth. <i>Oncotarget</i> , 2017, 8, 40115-40131.	1.8	35
72	The Role of Hypoxia Regulated microRNAs in Cancer. <i>Current Topics in Microbiology and Immunology</i> , 2010, 345, 47-70.	1.1	34

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73	In vitro downregulated hypoxia transcriptome is associated with poor prognosis in breast cancer. <i>Molecular Cancer</i> , 2017, 16, 105.	19.2	33
74	Identification of vitamin B1 metabolism as a tumor-specific radiosensitizing pathway using a high-throughput colony formation screen. <i>Oncotarget</i> , 2015, 6, 5978-5989.	1.8	33
75	COX-2 expression is predictive for early relapse and aromatase inhibitor resistance in patients with ductal carcinoma in situ of the breast, and is a target for treatment. <i>British Journal of Cancer</i> , 2014, 111, 46-54.	6.4	32
76	Close and Stable Relationship between Proliferation and a Hypoxia Metagene in Aromatase Inhibitor-Treated ER-Positive Breast Cancer. <i>Clinical Cancer Research</i> , 2011, 17, 3005-3012.	7.0	31
77	Role of carbohydrate response element-binding protein (ChREBP) in generating an aerobic metabolic phenotype and in breast cancer progression. <i>British Journal of Cancer</i> , 2014, 110, 715-723.	6.4	30
78	Combining lapatinib and pertuzumab to overcome lapatinib resistance due to NRG1-mediated signalling in HER2-amplified breast cancer. <i>Oncotarget</i> , 2015, 6, 5678-5694.	1.8	30
79	Dosimetric features of linac head and phantom scattered radiation outside the clinical photon beam: experimental measurements and comparison with treatment planning system calculations. <i>Radiotherapy and Oncology</i> , 2001, 58, 193-200.	0.6	29
80	Gene Expression Analysis in Human Breast Cancer Associated Blood Vessels. <i>PLoS ONE</i> , 2012, 7, e44294.	2.5	28
81	Transcriptomic analysis of human primary breast cancer identifies fatty acid oxidation as a target for metformin. <i>British Journal of Cancer</i> , 2020, 122, 258-265.	6.4	28
82	Radiogenomics Monitoring in Breast Cancer Identifies Metabolism and Immune Checkpoints as Early Actionable Mechanisms of Resistance to Anti-angiogenic Treatment. <i>EBioMedicine</i> , 2016, 10, 109-116.	6.1	27
83	Monte Carlo dose calculations and radiobiological modelling: analysis of the effect of the statistical noise of the dose distribution on the probability of tumour control. <i>Physics in Medicine and Biology</i> , 2000, 45, 3009-3023.	3.0	25
84	Multiple biomarker tissue microarrays: bioinformatics and practical approaches. <i>Cancer and Metastasis Reviews</i> , 2008, 27, 481-494.	5.9	25
85	A phase 2 study of high-activity ¹⁸⁶ Re-HEDP with autologous peripheral blood stem cell transplant in progressive hormone-refractory prostate cancer metastatic to bone. <i>European Journal of Nuclear Medicine and Molecular Imaging</i> , 2006, 33, 1055-1061.	6.4	24
86	RIPOSTE: a framework for improving the design and analysis of laboratory-based research. <i>ELife</i> , 2015, 4, .	6.0	24
87	Guidelines for using sigQC for systematic evaluation of gene signatures. <i>Nature Protocols</i> , 2019, 14, 1377-1400.	12.0	23
88	CD44v3 levels in primary cutaneous melanoma are predictive of prognosis: Assessment by the use of tissue microarray. <i>International Journal of Cancer</i> , 2006, 118, 1460-1464.	5.1	22
89	Adaptation to HIF1± Deletion in Hypoxic Cancer Cells by Upregulation of GLUT14 and Creatine Metabolism. <i>Molecular Cancer Research</i> , 2019, 17, 1531-1544.	3.4	22
90	Functional evolution of IGF2:IGF2R domain 11 binding generates novel structural interactions and a specific IGF2 antagonist. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E2766-75.	7.1	21

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91	Igf2 ligand dependency of Pten+/ \hat{a} developmental and tumour phenotypes in the mouse. <i>Oncogene</i> , 2012, 31, 3635-3646.	5.9	20
92	TOPK modulates tumour-specific radiosensitivity and correlates with recurrence after prostate radiotherapy. <i>British Journal of Cancer</i> , 2017, 117, 503-512.	6.4	20
93	The mevalonate precursor enzyme HMGS1 is a novel marker and key mediator of cancer stem cell enrichment in luminal and basal models of breast cancer. <i>PLoS ONE</i> , 2020, 15, e0236187.	2.5	20
94	Hypoxia-induced SETX links replication stress with the unfolded protein response. <i>Nature Communications</i> , 2021, 12, 3686.	12.8	19
95	An analysis of the relationship between radiosensitivity and volume effects in tumor control probability modeling. <i>Medical Physics</i> , 2000, 27, 1258-1265.	3.0	18
96	MicroRNA-Related DNA Repair/Cell-Cycle Genes Independently Associated With Relapse After Radiation Therapy for Early Breast Cancer. <i>International Journal of Radiation Oncology Biology Physics</i> , 2015, 93, 1104-1114.	0.8	18
97	Replication catastrophe induced by cyclic hypoxia leads to increased APOBEC3B activity. <i>Nucleic Acids Research</i> , 2021, 49, 7492-7506.	14.5	18
98	Vitamin D Receptor Expression in Plasmablastic Lymphoma and Myeloma Cells Confers Susceptibility to Vitamin D. <i>Endocrinology</i> , 2017, 158, 503-515.	2.8	17
99	Paracrine effect of GTP cyclohydrolase and angiopoietin-1 interaction in stromal fibroblasts on tumor Tie2 activation and breast cancer growth. <i>Oncotarget</i> , 2016, 7, 9353-9367.	1.8	17
100	A model-based method for the prediction of whole-body absorbed dose and bone marrow toxicity for 186 Re-HEDP treatment of skeletal metastases from prostate cancer. <i>European Journal of Nuclear Medicine and Molecular Imaging</i> , 2003, 30, 1114-1124.	6.4	16
101	Interferon- and STING-independent induction of type I interferon stimulated genes during fractionated irradiation. <i>Journal of Experimental and Clinical Cancer Research</i> , 2021, 40, 161.	8.6	16
102	Pre-treatment proliferation and the outcome of conventional and accelerated radiotherapy. <i>European Journal of Cancer</i> , 2006, 42, 363-371.	2.8	15
103	An in vivo hypoxia metagene identifies the novel hypoxia inducible factor target gene SLCO1B3. <i>European Journal of Cancer</i> , 2013, 49, 1741-1751.	2.8	15
104	Multicellular Dosimetry in Voxel Geometry for Targeted Radionuclide Therapy. <i>Cancer Biotherapy and Radiopharmaceuticals</i> , 2003, 18, 451-461.	1.0	14
105	nm23 as a prognostic marker in primary cutaneous melanoma: evaluation using tissue microarray in a patient group with long-term follow-up. <i>Melanoma Research</i> , 2005, 15, 435-440.	1.2	14
106	A semantic interoperability approach to support integration of gene expression and clinical data in breast cancer. <i>Computers in Biology and Medicine</i> , 2017, 87, 179-186.	7.0	14
107	Modeling genotypes in their microenvironment to predict single- and multi-cellular behavior. <i>GigaScience</i> , 2019, 8, .	6.4	14
108	Quantifying effects of lead shielding in electron beams: a Monte Carlo study. <i>Physics in Medicine and Biology</i> , 2001, 46, 757-769.	3.0	13

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109	Dose- and Time-Dependent Changes in Gene Expression in Human Glioma Cells after Low Radiation Doses. <i>Radiation Research</i> , 2007, 168, 199-208.	1.5	13
110	Informed Consent, Biobank Research, and Locality. <i>Journal of Empirical Research on Human Research Ethics</i> , 2014, 9, 48-55.	1.3	12
111	Nucleoporin 54 contributes to homologous recombination repair and post-replicative DNA integrity. <i>Nucleic Acids Research</i> , 2018, 46, 7731-7746.	14.5	11
112	Disruption of hypoxia-inducible fatty acid binding protein 7 induces beige fat-like differentiation and thermogenesis in breast cancer cells. <i>Cancer & Metabolism</i> , 2020, 8, 13.	5.0	11
113	Role of gene signatures combined with pathology in classification of oropharynx head and neck cancer. <i>Scientific Reports</i> , 2020, 10, 10226.	3.3	10
114	Backscatter and Dose Perturbations for Low- to Medium-Energy Electron Point Sources at the Interface between Materials with Different Atomic Numbers. <i>Radiation Research</i> , 2004, 162, 693-701.	1.5	9
115	Identification of P-cadherin in Primary Melanoma Using a Tissue Microarray. <i>Annals of Plastic Surgery</i> , 2005, 55, 316-320.	0.9	9
116	Heritable genetic variants in key cancer genes link cancer risk with anthropometric traits. <i>Journal of Medical Genetics</i> , 2021, 58, 392-399.	3.2	9
117	A new procedure for determining the genetic basis of a physiological process in a non-model species, illustrated by cold induced angiogenesis in the carp. <i>BMC Genomics</i> , 2009, 10, 490.	2.8	8
118	Neoadjuvant Window Studies of Metformin and Biomarker Development for Drugs Targeting Cancer Metabolism. <i>Journal of the National Cancer Institute Monographs</i> , 2015, 2015, 81-86.	2.1	8
119	Depletion of signal recognition particle 72kDa increases radiosensitivity. <i>Cancer Biology and Therapy</i> , 2017, 18, 425-432.	3.4	8
120	Guest Editorial Data Science in Smart Healthcare: Challenges and Opportunities. <i>IEEE Journal of Biomedical and Health Informatics</i> , 2020, 24, 3041-3043.	6.3	8
121	A Monte-Carlo Method for Interface Dosimetry of Beta Emitters. <i>Cancer Biotherapy and Radiopharmaceuticals</i> , 2003, 18, 463-471.	1.0	7
122	Towards an environment for data mining based analysis processes in bioinformatics and personalized medicine. <i>Network Modeling Analysis in Health Informatics and Bioinformatics</i> , 2013, 2, 29-44.	2.1	7
123	A general framework for quantifying the effects of DNA repair inhibitors on radiation sensitivity as a function of dose. <i>Theoretical Biology and Medical Modelling</i> , 2007, 4, 25.	2.1	6
124	Towards an environment for data mining based analysis processes in bioinformatics & personalized medicine. , 2011, , .		6
125	Disease-associated KBTBD4 mutations in medulloblastoma elicit neomorphic ubiquitylation activity to promote CoREST degradation. <i>Cell Death and Differentiation</i> , 2022, 29, 1955-1969.	11.2	6
126	Liver glycogen phosphorylase is upregulated in glioblastoma and provides a metabolic vulnerability to high dose radiation. <i>Cell Death and Disease</i> , 2022, 13, .	6.3	6

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127	The many faces of mathematical modelling in oncology. British Journal of Radiology, 2019, 92, 20180856.	2.2	5
128	Endogenous miRNA sponges mediate the generation of oscillatory dynamics for a non-coding RNA network. Journal of Theoretical Biology, 2019, 481, 54-60.	1.7	5
129	Defining T Cell Subsets in Human Tonsils Using ChipCytometry. Journal of Immunology, 2021, 206, 3073-3082.	0.8	5
130	Identification of anticancer drugs to radiosensitise BRAF-wild-type and mutant colorectal cancer. Cancer Biology and Medicine, 2019, 16, 234.	3.0	4
131	ADGRL4/ELTD1 Expression in Breast Cancer Cells Induces Vascular Normalization and Immune Suppression. Molecular Cancer Research, 2021, 19, 1957-1969.	3.4	4
132	Considerations for modelling MLCs with Monte Carlo techniques. , 2000, , 458-460.		3
133	A Structural Characterisation of the Mitogen-Activated Protein Kinase Network in Cancer. Symmetry, 2022, 14, 1009.	2.2	2
134	A DCE-MRI analysis workflow. , 2016, , .		1
135	Erratum to "radiation response and cure of human colon adenocarcinoma spheroids of different size: the significance of hypoxia tumor control modelling" International Journal of Radiation Oncology Biology Physics, 2004, 58, 1322.	0.8	0
136	Molecular Markers in the CHART Trial. International Journal of Radiation Oncology Biology Physics, 2005, 63, S72.	0.8	0
137	31 Radiation responsive genes in glioma cells. Radiotherapy and Oncology, 2006, 78, S11-S12.	0.6	0
138	189 The response of head and neck tumours to radiotherapy may be influenced by the interaction between cell proliferation and vascularization. Radiotherapy and Oncology, 2006, 78, S65-S66.	0.6	0
139	332 POLQ (DNA polymerase theta) as a novel therapeutic target: preclinical and clinical data. European Journal of Cancer, Supplement, 2010, 8, 106.	2.2	0
140	Correction: Close And Stable Relationship Between Proliferation And A Hypoxia Metagene In Aromatase Inhibitor-Treated ER-Positive Breast Cancer. Clinical Cancer Research, 2011, 17, 4915-4915.	7.0	0
141	Functional comparison of Notch ligands in tumour angiogenesis. Asian Pacific Journal of Tropical Disease, 2014, 4, 229.	0.5	0
142	D6-03: Proliferation and "invasiveness" gene-expression signatures predict survival of surgically treated non-small-cell lung cancer. Journal of Thoracic Oncology, 2007, 2, S407.	1.1	0
143	Fundamental Radiobiology and its Application to Radiation Oncology. NATO Science for Peace and Security Series B: Physics and Biophysics, 2009, , 3-9.	0.3	0
144	Abstract P2-02-07: Predicting Response to Bevacizumab Therapy in Primary Breast Cancer Using Dynamic Contrast-Enhanced Magnetic Resonance Imaging. , 2010, , .		0

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145	Abstract P2-09-28: Integrated Gene Expression and MRI Analysis To Assess Early Therapeutic Response to Bevacizumab in Primary Breast Cancer. , 2010, , .		0
146	Diagnostic and Prognostic Cancer Biomarkers: From Traditional to Systems Approaches. , 2011, , 329-366.		0
147	Abstract 38: Copy number aberration detected in head and neck carcinomas using OncoSNP and influence on gene expression. , 2011, , .		0
148	Abstract P1-06-01: Upregulation of metabolism as a potential resistance mechanism to bevacizumab in primary breast cancer. , 2012, , .		0
149	Abstract P4-01-01: Integrating dynamic magnetic resonance imaging and gene expression profiling reveals novel therapeutic targets in locally advanced breast cancer.. , 2012, , .		0
150	Abstract BS02-1: Hypoxia metabolism in breast cancer – How to overcome resistance to anti-angiogenic therapy. , 2013, , .		0
151	Abstract P2-05-01: The non-coding transcriptome of hypoxic breast cancer: Novel insights of clinical relevant long non-coding RNA in hypoxia signalling. , 2015, , .		0
152	Abstract 2903: Differential regulation of LC3 A and B, GABARAPL 1 and 2 autophagy genes by micro-environmental stress and role in breast cancer survival. , 2015, , .		0
153	Abstract 3750: Genomic alterations underlie a pan-cancer metabolic transcriptome shift. , 2015, , .		0
154	Abstract B1-56: Distinct roles of copy number and loss-of-heterozygosity in predicting prognosis for breast cancer patients. , 2015, , .		0
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