

Salvatore Piscuoglio

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

138
papers

5,273
citations

38
h-index

70
g-index

211
ext. papers

6,742
ext. citations

8.3
avg, IF

5.2
L-index

#	Paper	IF	Citations
138	Circulating Cell-Free DNA Captures the Intratumor Heterogeneity in Multinodular Hepatocellular Carcinoma.. <i>JCO Precision Oncology</i> , 2022 , 6, e2100335	3.6	1
137	GATA3 and MDM2 are synthetic lethal in estrogen receptor-positive breast cancers.. <i>Communications Biology</i> , 2022 , 5, 373	6.7	0
136	Alterations in homologous recombination repair genes in prostate cancer brain metastases.. <i>Nature Communications</i> , 2022 , 13, 2400	17.4	3
135	Integrative proteogenomic characterization of hepatocellular carcinoma across etiologies and stages.. <i>Nature Communications</i> , 2022 , 13, 2436	17.4	1
134	Standardizing Patient-Derived Organoid Generation Workflow to Avoid Microbial Contamination From Colorectal Cancer Tissues.. <i>Frontiers in Oncology</i> , 2021 , 11, 781833	5.3	1
133	YAP/TAZ and ATF4 drive resistance to Sorafenib in hepatocellular carcinoma by preventing ferroptosis. <i>EMBO Molecular Medicine</i> , 2021 , 13, e14351	12	11
132	The Genomic Landscape of Serrated Lesion of the Colorectum: Similarities and Differences With Tubular and Tubulovillous Adenomas. <i>Frontiers in Oncology</i> , 2021 , 11, 668466	5.3	0
131	Nestin and CD34 expression in colorectal cancer predicts improved overall survival. <i>Acta Oncologica</i> , 2021 , 60, 727-734	3.2	2
130	The Tumor Profiler Study: integrated, multi-omic, functional tumor profiling for clinical decision support. <i>Cancer Cell</i> , 2021 , 39, 288-293	24.3	21
129	Genomic evolutionary trajectory of metastatic squamous cell carcinoma of the lung. <i>Translational Lung Cancer Research</i> , 2021 , 10, 1792-1803	4.4	0
128	Genomic characterization of small cell carcinomas of the uterine cervix. <i>Molecular Oncology</i> , 2021 ,	7.9	1
127	Prostate cancer patient-derived organoids: detailed outcome from a prospective cohort of 81 clinical specimens. <i>Journal of Pathology</i> , 2021 , 254, 543-555	9.4	7
126	Non-viral gene delivery of the oncotoxic protein NS1 for treatment of hepatocellular carcinoma. <i>Journal of Controlled Release</i> , 2021 , 334, 138-152	11.7	1
125	ZNRF3 and RNF43 cooperate to safeguard metabolic liver zonation and hepatocyte proliferation. <i>Cell Stem Cell</i> , 2021 , 28, 1822-1837.e10	18	10
124	Mapping of mA and Its Regulatory Targets in Prostate Cancer Reveals a METTL3-Low Induction of Therapy Resistance. <i>Molecular Cancer Research</i> , 2021 , 19, 1398-1411	6.6	2
123	Systematic identification of novel cancer genes through analysis of deep shRNA perturbation screens. <i>Nucleic Acids Research</i> , 2021 , 49, 8488-8504	20.1	0
122	The genetic landscape of metaplastic breast cancers and uterine carcinosarcomas. <i>Molecular Oncology</i> , 2021 , 15, 1024-1039	7.9	6

121	Prognostic significance of CD8+ T-cells density in stage III colorectal cancer depends on SDF-1 expression. <i>Scientific Reports</i> , 2021 , 11, 775	4.9	5
120	Adenylosuccinate lyase is oncogenic in colorectal cancer by causing mitochondrial dysfunction and independent activation of NRF2 and mTOR-MYC-axis. <i>Theranostics</i> , 2021 , 11, 4011-4029	12.1	7
119	Transcriptional Enhancer Factor Domain Family member 4 Exerts an Oncogenic Role in Hepatocellular Carcinoma by Hippo-Independent Regulation of Heat Shock Protein 70 Family Members. <i>Hepatology Communications</i> , 2021 , 5, 661-674	6	3
118	Patient-derived xenografts and organoids model therapy response in prostate cancer. <i>Nature Communications</i> , 2021 , 12, 1117	17.4	18
117	USP29-mediated HIF1 β stabilization is associated with Sorafenib resistance of hepatocellular carcinoma cells by upregulating glycolysis. <i>Oncogenesis</i> , 2021 , 10, 52	6.6	8
116	Multi-omics data integration reveals novel drug targets in hepatocellular carcinoma. <i>BMC Genomics</i> , 2021 , 22, 592	4.5	1
115	mTOR signaling mediates ILC3-driven immunopathology. <i>Mucosal Immunology</i> , 2021 , 14, 1323-1334	9.2	2
114	Establishing standardized immune phenotyping of metastatic melanoma by digital pathology. <i>Laboratory Investigation</i> , 2021 , 101, 1561-1570	5.9	0
113	Utility of Serial cfDNA NGS for Prospective Genomic Analysis of Patients on a Phase I Basket Study. <i>JCO Precision Oncology</i> , 2021 , 5,	3.6	2
112	Role of specialized composition of SWI/SNF complexes in prostate cancer lineage plasticity. <i>Nature Communications</i> , 2020 , 11, 5549	17.4	31
111	Discovery of heterozygous KRT10 alterations in MAUIE cases underlines the importance of regular skin cancer screening in ichthyosis with confetti. <i>British Journal of Dermatology</i> , 2020 , 183, 954-955	4	
110	High Expression of FAP in Colorectal Cancer Is Associated With Angiogenesis and Immunoregulation Processes. <i>Frontiers in Oncology</i> , 2020 , 10, 979	5.3	17
109	Identification of Somatic Mutations in Thirty-year-old Serum Cell-free DNA From Patients With Breast Cancer: A Feasibility Study. <i>Clinical Breast Cancer</i> , 2020 , 20, 413-421.e1	3	1
108	A Pygopus 2-Histone Interaction Is Critical for Cancer Cell Dedifferentiation and Progression in Malignant Breast Cancer. <i>Cancer Research</i> , 2020 , 80, 3631-3648	10.1	4
107	Reliability of liquid biopsy analysis: an inter-laboratory comparison of circulating tumor DNA extraction and sequencing with different platforms. <i>Laboratory Investigation</i> , 2020 , 100, 1475-1484	5.9	6
106	Prevention of dsRNA-induced interferon signaling by AGO1x is linked to breast cancer cell proliferation. <i>EMBO Journal</i> , 2020 , 39, e103922	13	10
105	SCIM: universal single-cell matching with unpaired feature sets. <i>Bioinformatics</i> , 2020 , 36, i919-i927	7.2	8
104	Hormone receptor and HER2 assessment in breast carcinoma metastatic to bone: A comparison between FNA cell blocks and decalcified core needle biopsies. <i>Cancer Cytopathology</i> , 2020 , 128, 133-145 ^{3,9}		1

103	Cell-free DNA in hepatocellular carcinoma 2020 , 199-209		0
102	Deciphering the clonal relationship between glandular and squamous components in adenosquamous carcinoma of the lung using whole exome sequencing. <i>Lung Cancer</i> , 2020 , 150, 132-138 ^{5,9}		4
101	Infiltration by IL22-Producing T Cells Promotes Neutrophil Recruitment and Predicts Favorable Clinical Outcome in Human Colorectal Cancer. <i>Cancer Immunology Research</i> , 2020 , 8, 1452-1462	12.5	3
100	Expression of RET is associated with Oestrogen receptor expression but lacks prognostic significance in breast cancer. <i>BMC Cancer</i> , 2019 , 19, 41	4.8	8
99	Genomic characterization of metastatic breast cancers. <i>Nature</i> , 2019 , 569, 560-564	50.4	256
98	Convergent Evolution of Copy Number Alterations in Multi-Centric Hepatocellular Carcinoma. <i>Scientific Reports</i> , 2019 , 9, 4611	4.9	2
97	Homologous recombination DNA repair defects in associated breast cancers. <i>Npj Breast Cancer</i> , 2019 , 5, 23	7.8	20
96	Genetic Alterations in Benign Breast Biopsies of Subsequent Breast Cancer Patients. <i>Frontiers in Medicine</i> , 2019 , 6, 166	4.9	5
95	Hepatocellular Carcinoma Xenografts Established From Needle Biopsies Preserve the Characteristics of the Originating Tumors. <i>Hepatology Communications</i> , 2019 , 3, 971-986	6	15
94	Infiltration by myeloperoxidase-positive neutrophils is an independent prognostic factor in breast cancer. <i>Breast Cancer Research and Treatment</i> , 2019 , 177, 581-589	4.4	22
93	PipeIT: A Singularity Container for Molecular Diagnostic Somatic Variant Calling on the Ion Torrent Next-Generation Sequencing Platform. <i>Journal of Molecular Diagnostics</i> , 2019 , 21, 884-894	5.1	5
92	Nectin-4 Expression Is an Independent Prognostic Biomarker and Associated With Better Survival in Triple-Negative Breast Cancer. <i>Frontiers in Medicine</i> , 2019 , 6, 200	4.9	7
91	Therapeutic Targeting of CD146/MCAM Reduces Bone Metastasis in Prostate Cancer. <i>Molecular Cancer Research</i> , 2019 , 17, 1049-1062	6.6	12
90	Preoperative plasma fatty acid metabolites inform risk of prostate cancer progression and may be used for personalized patient stratification. <i>BMC Cancer</i> , 2019 , 19, 1216	4.8	8
89	LATS1 but not LATS2 represses autophagy by a kinase-independent scaffold function. <i>Nature Communications</i> , 2019 , 10, 5755	17.4	22
88	Lobular Carcinomas Display Intralesion Genetic Heterogeneity and Clonal Evolution in the Progression to Invasive Lobular Carcinoma. <i>Clinical Cancer Research</i> , 2019 , 25, 674-686	12.9	31
87	Fibroepithelial Breast Lesion: When Sequencing Can Help to Make a Clinical Decision. A Case Report. <i>Clinical Breast Cancer</i> , 2019 , 19, e1-e6	3	1
86	The Genomic Landscape of Mucinous Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 737-741	9.7	41

85	Genetic profiling using plasma-derived cell-free DNA in therapy-naïve hepatocellular carcinoma patients: a pilot study. <i>Annals of Oncology</i> , 2018 , 29, 1286-1291	10.3	49
84	Mutation Profiling of Key Cancer Genes in Primary Breast Cancers and Their Distant Metastases. <i>Cancer Research</i> , 2018 , 78, 3112-3121	10.1	37
83	NGS-pipe: a flexible, easily extendable and highly configurable framework for NGS analysis. <i>Bioinformatics</i> , 2018 , 34, 107-108	7.2	11
82	The protein histidine phosphatase LHPP is a tumour suppressor. <i>Nature</i> , 2018 , 555, 678-682	50.4	96
81	High expression of HOXA13 correlates with poorly differentiated hepatocellular carcinomas and modulates sorafenib response in in vitro models. <i>Laboratory Investigation</i> , 2018 , 98, 95-105	5.9	30
80	Contralateral breast cancers: Independent cancers or metastases?. <i>International Journal of Cancer</i> , 2018 , 142, 347-356	7.5	23
79	Organoid Models of Human Liver Cancers Derived from Tumor Needle Biopsies. <i>Cell Reports</i> , 2018 , 24, 1363-1376	10.6	166
78	Genomic Analysis Revealed New Oncogenic Signatures in -Mutant Hepatocellular Carcinoma. <i>Frontiers in Genetics</i> , 2018 , 9, 2	4.5	21
77	Circulating Cell-Free DNA in Hepatocellular Carcinoma: Current Insights and Outlook. <i>Frontiers in Medicine</i> , 2018 , 5, 78	4.9	31
76	Cancer Diagnosis Using a Liquid Biopsy: Challenges and Expectations. <i>Diagnostics</i> , 2018 , 8,	3.8	69
75	The Role of Long Non-Coding RNAs in Hepatocarcinogenesis. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	49
74	Recurrent hotspot mutations in HRAS Q61 and PI3K-AKT pathway genes as drivers of breast adenomyoepitheliomas. <i>Nature Communications</i> , 2018 , 9, 1816	17.4	82
73	Diagnostic Targeted Sequencing Panel for Hepatocellular Carcinoma Genomic Screening. <i>Journal of Molecular Diagnostics</i> , 2018 , 20, 836-848	5.1	9
72	Hepatocellular Carcinoma: Pathology and Genetics 2018 , 198-198		0
71	Reliability of Whole-Exome Sequencing for Assessing Intratumor Genetic Heterogeneity. <i>Cell Reports</i> , 2018 , 25, 1446-1457	10.6	55
70	Genetic heterogeneity and actionable mutations in HER2-positive primary breast cancers and their brain metastases. <i>Oncotarget</i> , 2018 , 9, 20617-20630	3.3	26
69	Genetic analysis of uterine adenosarcomas and phyllodes tumors of the breast. <i>Molecular Oncology</i> , 2017 , 11, 913-926	7.9	6
68	Whole-genome single-cell copy number profiling from formalin-fixed paraffin-embedded samples. <i>Nature Medicine</i> , 2017 , 23, 376-385	50.5	82

67	The Landscape of Somatic Genetic Alterations in Metaplastic Breast Carcinomas. <i>Clinical Cancer Research</i> , 2017 , 23, 3859-3870	12.9	92
66	Genetic analysis of a morphologically heterogeneous ovarian endometrioid carcinoma. <i>Histopathology</i> , 2017 , 71, 480-487	7.3	2
65	High mobility group A1 enhances tumorigenicity of human cholangiocarcinoma and confers resistance to therapy. <i>Molecular Carcinogenesis</i> , 2017 , 56, 2146-2157	5	15
64	Genetic Heterogeneity in Therapy-Naïve Synchronous Primary Breast Cancers and Their Metastases. <i>Clinical Cancer Research</i> , 2017 , 23, 4402-4415	12.9	69
63	Bi-allelic alterations in DNA repair genes underpin homologous recombination DNA repair defects in breast cancer. <i>Journal of Pathology</i> , 2017 , 242, 165-177	9.4	35
62	The genetic landscape of breast carcinomas with neuroendocrine differentiation. <i>Journal of Pathology</i> , 2017 , 241, 405-419	9.4	35
61	Phyllodes tumors with and without fibroadenoma-like areas display distinct genomic features and may evolve through distinct pathways. <i>Npj Breast Cancer</i> , 2017 , 3, 40	7.8	33
60	Vascular endothelial growth factor A amplification in colorectal cancer is associated with reduced M1 and M2 macrophages and diminished PD-1-expressing lymphocytes. <i>PLoS ONE</i> , 2017 , 12, e0175563	3.7	12
59	Phosphoprotein enriched in diabetes (PED/PEA15) promotes migration in hepatocellular carcinoma and confers resistance to sorafenib. <i>Cell Death and Disease</i> , 2017 , 8, e3138	9.8	17
58	Pregnancy at early age is associated with a reduction of progesterone-responsive cells and epithelial Wnt signaling in human breast tissue. <i>Oncotarget</i> , 2017 , 8, 22353-22360	3.3	6
57	Diverse and Reversion Mutations in Circulating Cell-Free DNA of Therapy-Resistant Breast or Ovarian Cancer. <i>Clinical Cancer Research</i> , 2017 , 23, 6708-6720	12.9	132
56	Genomic and transcriptomic heterogeneity in metaplastic carcinomas of the breast. <i>Npj Breast Cancer</i> , 2017 , 3, 48	7.8	38
55	Genetic analysis of microglandular adenosis and acinic cell carcinomas of the breast provides evidence for the existence of a low-grade triple-negative breast neoplasia family. <i>Modern Pathology</i> , 2017 , 30, 69-84	9.8	34
54	Preclinical evaluation of the PARP inhibitor BMN-673 for the treatment of ovarian clear cell cancer. <i>Oncotarget</i> , 2017 , 8, 6057-6066	3.3	11
53	Cancer-associated fibroblasts release exosomal microRNAs that dictate an aggressive phenotype in breast cancer. <i>Oncotarget</i> , 2017 , 8, 19592-19608	3.3	208
52	HMGA1 overexpression is associated with a particular subset of human breast carcinomas. <i>Journal of Clinical Pathology</i> , 2016 , 69, 117-21	3.9	12
51	RNASeq analysis reveals biological processes governing the clinical behaviour of endometrioid and serous endometrial cancers. <i>European Journal of Cancer</i> , 2016 , 64, 149-58	7.5	4
50	HMGA1 Expression in Human Hepatocellular Carcinoma Correlates with Poor Prognosis and Promotes Tumor Growth and Migration in in vitro Models. <i>Neoplasia</i> , 2016 , 18, 724-731	6.4	25

49	Uterine adenosarcomas are mesenchymal neoplasms. <i>Journal of Pathology</i> , 2016 , 238, 381-8	9.4	70
48	TP53 Mutational Spectrum in Endometrioid and Serous Endometrial Cancers. <i>International Journal of Gynecological Pathology</i> , 2016 , 35, 289-300	3.2	51
47	Targeted capture massively parallel sequencing analysis of LCIS and invasive lobular cancer: Repertoire of somatic genetic alterations and clonal relationships. <i>Molecular Oncology</i> , 2016 , 10, 360-70	7.9	36
46	The Genomic Landscape of Male Breast Cancers. <i>Clinical Cancer Research</i> , 2016 , 22, 4045-56	12.9	85
45	Massively Parallel Sequencing-Based Clonality Analysis of Synchronous Endometrioid Endometrial and Ovarian Carcinomas. <i>Journal of the National Cancer Institute</i> , 2016 , 108, djv427	9.7	111
44	Abstract S4-03: A functional assay for homologous recombination (HR) DNA repair and whole exome sequencing reveal that HR-defective sporadic breast cancers are enriched for genetic alterations in DNA repair genes 2016 ,		3
43	Pilot trial of an implantable microdevice for In Vivo drug sensitivity testing in patients with early stage, triple negative breast cancer receiving neoadjuvant therapy.. <i>Journal of Clinical Oncology</i> , 2016 , 34, TPS1101-TPS1101	2.2	0
42	Serial next generation sequencing (NGS) of cell free DNA (cfDNA) and clonal evolution of AKT1 E17K mutant tumors: Analyses from patients (pts) enrolled on a phase I basket study of an AKT inhibitor (AZD5363).. <i>Journal of Clinical Oncology</i> , 2016 , 34, 11500-11500	2.2	
41	Lack of PRKD2 and PRKD3 kinase domain somatic mutations in PRKD1 wild-type classic polymorphous low-grade adenocarcinomas of the salivary gland. <i>Histopathology</i> , 2016 , 68, 1055-62	7.3	21
40	Infiltrating epitheliosis of the breast: characterization of histological features, immunophenotype and genomic profile. <i>Histopathology</i> , 2016 , 68, 1030-9	7.3	21
39	Massively parallel sequencing of phyllodes tumours of the breast reveals actionable mutations, and TERT promoter hotspot mutations and TERT gene amplification as likely drivers of progression. <i>Journal of Pathology</i> , 2016 , 238, 508-18	9.4	80
38	Microglandular adenosis associated with triple-negative breast cancer is a neoplastic lesion of triple-negative phenotype harbouring TP53 somatic mutations. <i>Journal of Pathology</i> , 2016 , 238, 677-88	9.4	42
37	Resolving quandaries: basaloid adenoid cystic carcinoma or breast cylindroma? The role of massively parallel sequencing. <i>Histopathology</i> , 2016 , 68, 262-71	7.3	16
36	Massively parallel sequencing analysis of synchronous fibroepithelial lesions supports the concept of progression from fibroadenoma to phyllodes tumor. <i>Npj Breast Cancer</i> , 2016 , 2, 16035	7.8	23
35	IDH2 Mutations Define a Unique Subtype of Breast Cancer with Altered Nuclear Polarity. <i>Cancer Research</i> , 2016 , 76, 7118-7129	10.1	70
34	Genetic alterations of triple negative breast cancer by targeted next-generation sequencing and correlation with tumor morphology. <i>Modern Pathology</i> , 2016 , 29, 476-88	9.8	67
33	Genetic events in the progression of adenoid cystic carcinoma of the breast to high-grade triple-negative breast cancer. <i>Modern Pathology</i> , 2016 , 29, 1292-1305	9.8	52
32	PIKING the type and pattern of PI3K pathway mutations in endometrioid endometrial carcinomas. <i>Gynecologic Oncology</i> , 2015 , 137, 321-8	4.9	14

31	Response to dual HER2 blockade in a patient with HER3-mutant metastatic breast cancer. <i>Annals of Oncology</i> , 2015 , 26, 1704-9	10.3	16
30	Genomic landscape of adenoid cystic carcinoma of the breast. <i>Journal of Pathology</i> , 2015 , 237, 179-89	9.4	101
29	Intra-tumor genetic heterogeneity and alternative driver genetic alterations in breast cancers with heterogeneous HER2 gene amplification. <i>Genome Biology</i> , 2015 , 16, 107	18.3	83
28	A survey of DICER1 hotspot mutations in ovarian and testicular sex cord-stromal tumors. <i>Modern Pathology</i> , 2015 , 28, 1603-12	9.8	71
27	High-resolution breakpoint analysis provides evidence for the sequence-directed nature of genome rearrangements in hereditary disorders. <i>Human Mutation</i> , 2015 , 36, 250-9	4.7	1
26	The repertoire of somatic genetic alterations of acinic cell carcinomas of the breast: an exploratory, hypothesis-generating study. <i>Journal of Pathology</i> , 2015 , 237, 166-78	9.4	42
25	Are acinic cell carcinomas of the breast and salivary glands distinct diseases?. <i>Histopathology</i> , 2015 , 67, 529-37	7.3	25
24	MED12 somatic mutations in fibroadenomas and phyllodes tumours of the breast. <i>Histopathology</i> , 2015 , 67, 719-29	7.3	68
23	Cerebrospinal fluid-derived circulating tumour DNA better represents the genomic alterations of brain tumours than plasma. <i>Nature Communications</i> , 2015 , 6, 8839	17.4	416
22	HIPK2 deficiency causes chromosomal instability by cytokinesis failure and increases tumorigenicity. <i>Oncotarget</i> , 2015 , 6, 10320-34	3.3	20
21	Characterization of the genomic features and expressed fusion genes in micropapillary carcinomas of the breast. <i>Journal of Pathology</i> , 2014 , 232, 553-65	9.4	75
20	Breast cancer intra-tumor heterogeneity. <i>Breast Cancer Research</i> , 2014 , 16, 210	8.3	188
19	Hotspot activating PRKD1 somatic mutations in polymorphous low-grade adenocarcinomas of the salivary glands. <i>Nature Genetics</i> , 2014 , 46, 1166-9	36.3	150
18	Capturing intra-tumor genetic heterogeneity by de novo mutation profiling of circulating cell-free tumor DNA: a proof-of-principle. <i>Annals of Oncology</i> , 2014 , 25, 1729-1735	10.3	258
17	Integrative genomic and transcriptomic characterization of papillary carcinomas of the breast. <i>Molecular Oncology</i> , 2014 , 8, 1588-602	7.9	38
16	Benchmarking mutation effect prediction algorithms using functionally validated cancer-related missense mutations. <i>Genome Biology</i> , 2014 , 15, 484	18.3	95
15	3SUTR poly(T/U) tract deletions and altered expression of EWSR1 are a hallmark of mismatch repair-deficient cancers. <i>Cancer Research</i> , 2014 , 74, 224-34	10.1	8
14	Long noncoding RNA HOTTIP/HOXA13 expression is associated with disease progression and predicts outcome in hepatocellular carcinoma patients. <i>Hepatology</i> , 2014 , 59, 911-23	11.2	333

13	SH2D4A is frequently downregulated in hepatocellular carcinoma and cirrhotic nodules. <i>European Journal of Cancer</i> , 2014 , 50, 731-8	7.5	3
12	90 HOXA13 AND HOTTIP EXPRESSION LEVELS PREDICT PATIENTSSSURVIVAL AND METASTASIS FORMATION IN HEPATOCELLULAR CARCINOMA. <i>Journal of Hepatology</i> , 2013 , 58, S39-S40	13.4	3
11	HMGA1 and HMGA2 protein expression correlates with advanced tumour grade and lymph node metastasis in pancreatic adenocarcinoma. <i>Histopathology</i> , 2012 , 60, 397-404	7.3	66
10	Effect of EpCAM, CD44, CD133 and CD166 expression on patient survival in tumours of the ampulla of Vater. <i>Journal of Clinical Pathology</i> , 2012 , 65, 140-5	3.9	27
9	MAGE-A10 is a nuclear protein frequently expressed in high percentages of tumor cells in lung, skin and urothelial malignancies. <i>International Journal of Cancer</i> , 2011 , 129, 1137-48	7.5	34
8	TIA-1 cytotoxic granule-associated RNA binding protein improves the prognostic performance of CD8 in mismatch repair-proficient colorectal cancer. <i>PLoS ONE</i> , 2010 , 5, e14282	3.7	42
7	Survivin expression in renal cell carcinoma. <i>Cancer Investigation</i> , 2008 , 26, 929-35	2.1	31
6	Reliability of Whole-Exome Sequencing for Assessing Intratumor Genetic Heterogeneity. <i>SSRN Electronic Journal</i> ,	1	2
5	Role of Specialized Composition of SWI/SNF Complexes in Prostate Cancer Lineage Plasticity		1
4	Patient-derived xenografts and organoids model therapy response in prostate cancer		1
3	The Genomic Landscape of Prostate Cancer Brain Metastases		2
2	Discovery of synthetic lethal interactions from large-scale pan-cancer perturbation screens		2
1	Democratisation of Enterprise Risk Management239-251		