Chad J Creighton

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.

81 167 30,302 320 h-index g-index citations papers 10.8 6.82 39,250 342 L-index avg, IF ext. papers ext. citations

#	Paper	IF	Citations
320	UALCAN: An update to the integrated cancer data analysis platform <i>Neoplasia</i> , 2022 , 25, 18-27	6.4	44
319	MAPK4 promotes triple negative breast cancer growth and reduces tumor sensitivity to PI3K blockade <i>Nature Communications</i> , 2022 , 13, 245	17.4	2
318	Abstract P5-05-06: Metformin concentration is a deciding factor of its pro- or anti-tumor function in triple negative breast cancer. <i>Cancer Research</i> , 2022 , 82, P5-05-06-P5-05-06	10.1	
317	A functional genomic approach to actionable gene fusions for precision oncology <i>Science Advances</i> , 2022 , 8, eabm2382	14.3	0
316	Proteogenomic characterization of 2002 human cancers reveals pan-cancer molecular subtypes and associated pathways <i>Nature Communications</i> , 2022 , 13, 2669	17.4	3
315	Transcriptional control of a collagen deposition and adhesion process that promotes lung adenocarcinoma growth and metastasis. <i>JCI Insight</i> , 2021 ,	9.9	2
314	The EMT activator ZEB1 accelerates endosomal trafficking to establish a polarity axis in lung adenocarcinoma cells. <i>Nature Communications</i> , 2021 , 12, 6354	17.4	5
313	Global molecular alterations involving recurrence or progression of pediatric brain tumors. <i>Neoplasia</i> , 2021 , 24, 22-33	6.4	1
312	MAPK6-AKT signaling promotes tumor growth and resistance to mTOR kinase blockade. <i>Science Advances</i> , 2021 , 7, eabi6439	14.3	1
311	Rearrangement-mediated cis-regulatory alterations in advanced patient tumors reveal interactions with therapy. <i>Cell Reports</i> , 2021 , 37, 110023	10.6	0
310	CKB inhibits epithelial-mesenchymal transition and prostate cancer progression by sequestering and inhibiting AKT activation. <i>Neoplasia</i> , 2021 , 23, 1147-1165	6.4	1
309	Identification of diverse tumor endothelial cell populations in malignant glioma. <i>Neuro-Oncology</i> , 2021 , 23, 932-944	1	12
308	SVExpress: identifying gene features altered recurrently in expression with nearby structural variant breakpoints. <i>BMC Bioinformatics</i> , 2021 , 22, 135	3.6	1
307	YAP1 overexpression contributes to the development of enzalutamide resistance by induction of cancer stemness and lipid metabolism in prostate cancer. <i>Oncogene</i> , 2021 , 40, 2407-2421	9.2	8
306	Portals for Exploring Noncoding Variants in Pediatric Cancer. <i>Trends in Genetics</i> , 2021 , 37, 297-298	8.5	
305	Contextual cues from cancer cells govern cancer-associated fibroblast heterogeneity. <i>Cell Reports</i> , 2021 , 35, 109009	10.6	3
304	A collagen glucosyltransferase drives lung adenocarcinoma progression in mice. <i>Communications Biology</i> , 2021 , 4, 482	6.7	3

(2020-2021)

303	Addiction to Golgi-resident PI4P synthesis in chromosome 1q21.3-amplified lung adenocarcinoma cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	1	
302	Endometrial receptivity and implantation require uterine BMP signaling through an ACVR2A-SMAD1/SMAD5 axis. <i>Nature Communications</i> , 2021 , 12, 3386	17.4	3	
301	p53 loss activates prometastatic secretory vesicle biogenesis in the Golgi. <i>Science Advances</i> , 2021 , 7,	14.3	4	
300	A Wnt-Independent LGR4-EGFR Signaling Axis in Cancer Metastasis. <i>Cancer Research</i> , 2021 , 81, 4441-445	5 4 0.1	2	
299	Female Gender Predicts Augmented Immune Infiltration in Lung Adenocarcinoma. <i>Clinical Lung Cancer</i> , 2021 , 22, e415-e424	4.9	5	
298	Neuropeptide Y nerve paracrine regulation of prostate cancer oncogenesis and therapy resistance. <i>Prostate</i> , 2021 , 81, 58-71	4.2	8	
297	Meta-analysis of host transcriptional responses to SARS-CoV-2 infection reveals their manifestation in human tumors. <i>Scientific Reports</i> , 2021 , 11, 2459	4.9	6	
296	Functional Hierarchy and Cooperation of EMT Master Transcription Factors in Breast Cancer Metastasis. <i>Molecular Cancer Research</i> , 2021 , 19, 784-798	6.6	8	
295	A pediatric brain tumor atlas of genes deregulated by somatic genomic rearrangement. <i>Nature Communications</i> , 2021 , 12, 937	17.4	9	
294	Systematic identification of non-coding somatic single nucleotide variants associated with altered transcription and DNA methylation in adult and pediatric cancers. <i>NAR Cancer</i> , 2021 , 3, zcab001	5.2	3	
293	Mass-spectrometry-based proteomic correlates of grade and stage reveal pathways and kinases associated with aggressive human cancers. <i>Oncogene</i> , 2021 , 40, 2081-2095	9.2	8	
292	MAPK4 promotes prostate cancer by concerted activation of androgen receptor and AKT. <i>Journal of Clinical Investigation</i> , 2021 , 131,	15.9	10	
291	A protumorigenic secretory pathway activated by p53 deficiency in lung adenocarcinoma. <i>Journal of Clinical Investigation</i> , 2021 , 131,	15.9	11	
2 90	In vivolmodeling of metastatic human high-grade serous ovarian cancer in mice. <i>PLoS Genetics</i> , 2020 , 16, e1008808	6	15	
289	Accumulation of Molecular Aberrations Distinctive to Hepatocellular Carcinoma Progression. <i>Cancer Research</i> , 2020 , 80, 3810-3819	10.1	9	
288	The Sca-1 and Sca-1 mouse prostatic luminal cell lineages are independently sustained. <i>Stem Cells</i> , 2020 , 38, 1479-1491	5.8	10	
287	PIK3CA variants selectively initiate brain hyperactivity during gliomagenesis. <i>Nature</i> , 2020 , 578, 166-171	50.4	50	
286	PI4KIIIIs a therapeutic target in chromosome 1q-amplified lung adenocarcinoma. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	19	

285	High-coverage whole-genome analysis of 1220 cancers reveals hundreds of genes deregulated by rearrangement-mediated cis-regulatory alterations. <i>Nature Communications</i> , 2020 , 11, 736	17.4	32
284	Genomic basis for RNA alterations in cancer. <i>Nature</i> , 2020 , 578, 129-136	50.4	148
283	Comprehensive Molecular Characterization Identifies Distinct Genomic and Immune Hallmarks of Renal Medullary Carcinoma. <i>Cancer Cell</i> , 2020 , 37, 720-734.e13	24.3	32
282	Comprehensive molecular characterization of mitochondrial genomes in human cancers. <i>Nature Genetics</i> , 2020 , 52, 342-352	36.3	105
281	Proteomic signatures of clear cell renal cell carcinoma. <i>Nature Reviews Nephrology</i> , 2020 , 16, 133-134	14.9	3
280	Constitutive expression of progesterone receptor isoforms promotes the development of hormone-dependent ovarian neoplasms. <i>Science Signaling</i> , 2020 , 13,	8.8	4
279	Renal Carcinoma Is Associated With Increased Risk of Coronavirus Infections. <i>Frontiers in Molecular Biosciences</i> , 2020 , 7, 579422	5.6	8
278	IMPAD1 and KDELR2 drive invasion and metastasis by enhancing Golgi-mediated secretion. <i>Oncogene</i> , 2020 , 39, 5979-5994	9.2	12
277	Pten and Dicer1 loss in the mouse uterus causes poorly differentiated endometrial adenocarcinoma. <i>Oncogene</i> , 2020 , 39, 6286-6299	9.2	4
276	Caveolin-1-mediated sphingolipid oncometabolism underlies a metabolic vulnerability of prostate cancer. <i>Nature Communications</i> , 2020 , 11, 4279	17.4	27
275	Reply to Liu et al.: ALK5-mediated tumor suppressor signaling through SMAD2 and SMAD3 in the uterus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 9166	5-9 1 167	
274	JNK represses Lkb-deficiency-induced lung squamous cell carcinoma progression. <i>Nature Communications</i> , 2019 , 10, 2148	17.4	13
273	Functional Heterogeneity of Mouse Prostate Stromal Cells Revealed by Single-Cell RNA-Seq. <i>IScience</i> , 2019 , 13, 328-338	6.1	17
272	Spatially Restricted Stromal Wnt Signaling Restrains Prostate Epithelial Progenitor Growth through Direct and Indirect Mechanisms. <i>Cell Stem Cell</i> , 2019 , 24, 753-768.e6	18	29
271	Spermidine/spermine N1-acetyltransferase 1 is a gene-specific transcriptional regulator that drives brain tumor aggressiveness. <i>Oncogene</i> , 2019 , 38, 6794-6800	9.2	13
270	Alterations in Wnt- and/or STAT3 signaling pathways and the immune microenvironment during metastatic progression. <i>Oncogene</i> , 2019 , 38, 5942-5958	9.2	7
269	Global impact of somatic structural variation on the DNA methylome of human cancers. <i>Genome Biology</i> , 2019 , 20, 209	18.3	9
268	ZEB1/NuRD complex suppresses TBC1D2b to stimulate E-cadherin internalization and promote metastasis in lung cancer. <i>Nature Communications</i> , 2019 , 10, 5125	17.4	35

(2018-2019)

267	A CTC-Cluster-Specific Signature Derived from OMICS Analysis of Patient-Derived Xenograft Tumors Predicts Outcomes in Basal-Like Breast Cancer. <i>Journal of Clinical Medicine</i> , 2019 , 8,	5.1	19	
266	MAPK4 overexpression promotes tumor progression via noncanonical activation of AKT/mTOR signaling. <i>Journal of Clinical Investigation</i> , 2019 , 129, 1015-1029	15.9	32	
265	MDMA Abuse in Relation to MicroRNA Variation in Human Brain Ventral Tegmental Area and Nucleus Accumbens. <i>Iranian Journal of Pharmaceutical Research</i> , 2019 , 18, 1989-1999	1.1	2	
264	Abnormal oxidative metabolism in a quiet genomic background underlies clear cell papillary renal cell carcinoma. <i>ELife</i> , 2019 , 8,	8.9	17	
263	Pan-cancer molecular subtypes revealed by mass-spectrometry-based proteomic characterization of more than 500 human cancers. <i>Nature Communications</i> , 2019 , 10, 5679	17.4	150	
262	Molecular Correlates of Metastasis by Systematic Pan-Cancer Analysis Across The Cancer Genome Atlas. <i>Molecular Cancer Research</i> , 2019 , 17, 476-487	6.6	15	
261	Mammary Precancerous Stem and Non-Stem Cells Evolve into Cancers of Distinct Subtypes. <i>Cancer Research</i> , 2019 , 79, 61-71	10.1	17	
260	Uterine double-conditional inactivation of and in mice causes endometrial dysregulation, infertility, and uterine cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 3873-3882	11.5	27	
259	MicroRNAs as prognostic markers in prostate cancer. <i>Prostate</i> , 2019 , 79, 265-271	4.2	19	
258	Osteoblast-Secreted Factors Mediate Dormancy of Metastatic Prostate Cancer in the Bone via Activation of the TGFRIII-p38MAPK-pS249/T252RB Pathway. <i>Cancer Research</i> , 2018 , 78, 2911-2924	10.1	73	
257	IL17A Regulates Tumor Latency and Metastasis in Lung Adeno and Squamous SQ.2b and AD.1 Cancer. <i>Cancer Immunology Research</i> , 2018 , 6, 645-657	12.5	19	
256	Comprehensive Characterization of Cancer Driver Genes and Mutations. <i>Cell</i> , 2018 , 173, 371-385.e18	56.2	854	
255	Cell-of-Origin Patterns Dominate the Molecular Classification of 10,000 Tumors from 33 Types of Cancer. <i>Cell</i> , 2018 , 173, 291-304.e6	56.2	888	
254	Perspective on Oncogenic Processes at the End of the Beginning of Cancer Genomics. <i>Cell</i> , 2018 , 173, 305-320.e10	56.2	166	
253	Oncogenic Signaling Pathways in The Cancer Genome Atlas. Cell, 2018, 173, 321-337.e10	56.2	1124	
252	Driver Fusions and Their Implications in the Development and Treatment of Human Cancers. <i>Cell Reports</i> , 2018 , 23, 227-238.e3	10.6	235	
251	Genomic, Pathway Network, and Immunologic Features Distinguishing Squamous Carcinomas. <i>Cell Reports</i> , 2018 , 23, 194-212.e6	10.6	146	
250	The Cancer Genome Atlas Comprehensive Molecular Characterization of Renal Cell Carcinoma. <i>Cell Reports</i> , 2018 , 23, 313-326.e5	10.6	295	

249	The Immune Landscape of Cancer. <i>Immunity</i> , 2018 , 48, 812-830.e14	32.3	1754
248	Genomic and Molecular Landscape of DNA Damage Repair Deficiency across The Cancer Genome Atlas. <i>Cell Reports</i> , 2018 , 23, 239-254.e6	10.6	405
247	A Landscape of Metabolic Variation across Tumor Types. <i>Cell Systems</i> , 2018 , 6, 301-313.e3	10.6	73
246	Pan-Cancer Molecular Classes Transcending Tumor Lineage Across 32 Cancer Types, Multiple Data Platforms, and over 10,000 Cases. <i>Clinical Cancer Research</i> , 2018 , 24, 2182-2193	12.9	49
245	Making Use of Cancer Genomic Databases. Current Protocols in Molecular Biology, 2018, 121, 19.14.1-19	.124913	7
244	Genomic classifications of renal cell carcinoma: a critical step towards the future application of personalized kidney cancer care with pan-omics precision. <i>Journal of Pathology</i> , 2018 , 244, 525-537	9.4	66
243	GPCRs profiling and identification of GPR110 as a potential new target in HER2+ breast cancer. Breast Cancer Research and Treatment, 2018 , 170, 279-292	4.4	14
242	Genomic and Functional Approaches to Understanding Cancer Aneuploidy. <i>Cancer Cell</i> , 2018 , 33, 676-68	3 9.e3	377
241	Atrx inactivation drives disease-defining phenotypes in glioma cells of origin through global epigenomic remodeling. <i>Nature Communications</i> , 2018 , 9, 1057	17.4	39
240	Combinatorial inhibition of PTPN12-regulated receptors leads to a broadly effective therapeutic strategy in triple-negative breast cancer. <i>Nature Medicine</i> , 2018 , 24, 505-511	50.5	28
239	Pan-cancer survey of epithelial-mesenchymal transition markers across the Cancer Genome Atlas. <i>Developmental Dynamics</i> , 2018 , 247, 555-564	2.9	53
238	Functional Annotation of ESR1 Gene Fusions in Estrogen Receptor-Positive Breast Cancer. <i>Cell Reports</i> , 2018 , 24, 1434-1444.e7	10.6	43
237	TMEM106B drives lung cancer metastasis by inducing TFEB-dependent lysosome synthesis and secretion of cathepsins. <i>Nature Communications</i> , 2018 , 9, 2731	17.4	48
236	In vivo screening identifies GATAD2B as a metastasis driver in KRAS-driven lung cancer. <i>Nature Communications</i> , 2018 , 9, 2732	17.4	18
235	A Pan-Cancer Compendium of Genes Deregulated by Somatic Genomic Rearrangement across More Than 1,400 Cases. <i>Cell Reports</i> , 2018 , 24, 515-527	10.6	49
234	Change in Neutrophil-to-lymphocyte ratio (NLR) in response to immune checkpoint blockade for metastatic renal cell carcinoma 2018 , 6, 5		129
233	The clinical applications of The Cancer Genome Atlas project for bladder cancer. <i>Expert Review of Anticancer Therapy</i> , 2018 , 18, 973-980	3.5	9
232	The epithelial-to-mesenchymal transition activator ZEB1 initiates a prometastatic competing endogenous RNA network. <i>Journal of Clinical Investigation</i> , 2018 , 128, 1267-1282	15.9	38

231	Reprogramming of the estrogen responsive transcriptome contributes to tamoxifen-dependent protection against tumorigenesis in the p53 null mammary epithelial cells. <i>PLoS ONE</i> , 2018 , 13, e01949	1 3 .7	5
230	Fibroblast growth factor receptor signaling plays a key role in transformation induced by the TMPRSS2/ERG fusion gene and decreased PTEN. <i>Oncotarget</i> , 2018 , 9, 14456-14471	3.3	5
229	Coactivation of Estrogen Receptor and IKKIInduces a Dormant Metastatic Phenotype in ER-Positive Breast Cancer. <i>Cancer Research</i> , 2018 , 78, 974-984	10.1	21
228	Influence of the neural microenvironment on prostate cancer. <i>Prostate</i> , 2018 , 78, 128-139	4.2	36
227	CBMT-23. MODULATION OF HYPERSYNAPTIC MICROENVIRONMENT DIFFERENTIALLY PROMOTES GLIOMAGENESIS ACROSS PIK3CA VARIANTS. <i>Neuro-Oncology</i> , 2018 , 20, vi37-vi37	1	78
226	miR-205 Regulates Basal Cell Identity and Stem Cell Regenerative Potential During Mammary Reconstitution. <i>Stem Cells</i> , 2018 , 36, 1875-1889	5.8	7
225	Integrated Multi-omic Analysis of Esthesioneuroblastomas Identifies Two Subgroups Linked to Cell Ontogeny. <i>Cell Reports</i> , 2018 , 25, 811-821.e5	10.6	25
224	Comprehensive Molecular Characterization of the Hippo Signaling Pathway in Cancer. <i>Cell Reports</i> , 2018 , 25, 1304-1317.e5	10.6	152
223	Mammary stem cell and macrophage markers are enriched in normal tissue adjacent to inflammatory breast cancer. <i>Breast Cancer Research and Treatment</i> , 2018 , 171, 283-293	4.4	12
222	Ki67 Proliferation Index as a Tool for Chemotherapy Decisions During and After Neoadjuvant Aromatase Inhibitor Treatment of Breast Cancer: Results From the American College of Surgeons Oncology Group Z1031 Trial (Alliance). <i>Journal of Clinical Oncology</i> , 2017 , 35, 1061-1069	2.2	164
221	Identification of diverse astrocyte populations and their malignant analogs. <i>Nature Neuroscience</i> , 2017 , 20, 396-405	25.5	275
220	A Versatile Tumor Gene Deletion System Reveals a Crucial Role for FGFR1 in Breast Cancer Metastasis. <i>Neoplasia</i> , 2017 , 19, 421-428	6.4	8
219	Engineering and Functional Characterization of Fusion Genes Identifies Novel Oncogenic Drivers of Cancer. <i>Cancer Research</i> , 2017 , 77, 3502-3512	10.1	22
218	RGS12 Is a Novel Tumor-Suppressor Gene in African American Prostate Cancer That Represses AKT and MNX1 Expression. <i>Cancer Research</i> , 2017 , 77, 4247-4257	10.1	18
217	A Pan-Cancer Proteogenomic Atlas of PI3K/AKT/mTOR Pathway Alterations. Cancer Cell, 2017, 31, 820-	8 34. 93	3 286
216	A Role for Progesterone-Regulated sFRP4 Expression in Uterine Leiomyomas. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017 , 102, 3316-3326	5.6	3
215	Exosome-Derived miR-25-3p and miR-92a-3p Stimulate Liposarcoma Progression. <i>Cancer Research</i> , 2017 , 77, 3846-3856	10.1	107
214	Decreased epithelial progesterone receptor A at the window of receptivity is required for preparation of the endometrium for embryo attachment. <i>Biology of Reproduction</i> , 2017 , 96, 313-326	3.9	48

213	The SWI/SNF Protein PBRM1 Restrains VHL-Loss-Driven Clear Cell Renal Cell Carcinoma. <i>Cell Reports</i> , 2017 , 18, 2893-2906	10.6	109
212	WNT-Mediated Regulation of FOXO1 Constitutes a Critical Axis Maintaining Pubertal Mammary Stem Cell Homeostasis. <i>Developmental Cell</i> , 2017 , 43, 436-448.e6	10.2	27
211	Glia-specific enhancers and chromatin structure regulate NFIA expression and glioma tumorigenesis. <i>Nature Neuroscience</i> , 2017 , 20, 1520-1528	25.5	24
210	UALCAN: A Portal for Facilitating Tumor Subgroup Gene Expression and Survival Analyses. <i>Neoplasia</i> , 2017 , 19, 649-658	6.4	2229
209	Sarcomatoid Renal Cell Carcinoma Has a Distinct Molecular Pathogenesis, Driver Mutation Profile, and Transcriptional Landscape. <i>Clinical Cancer Research</i> , 2017 , 23, 6686-6696	12.9	48
208	Pan-urologic cancer genomic subtypes that transcend tissue of origin. <i>Nature Communications</i> , 2017 , 8, 199	17.4	35
207	Thy-1 Cancer-associated Fibroblasts Adversely Impact Lung Cancer Prognosis. <i>Scientific Reports</i> , 2017 , 7, 6478	4.9	21
206	Ror2-mediated alternative Wnt signaling regulates cell fate and adhesion during mammary tumor progression. <i>Oncogene</i> , 2017 , 36, 5958-5968	9.2	25
205	Multiplatform-based molecular subtypes of non-small-cell lung cancer. <i>Oncogene</i> , 2017 , 36, 1384-1393	9.2	75
204	ZEB1 induces LOXL2-mediated collagen stabilization and deposition in the extracellular matrix to drive lung cancer invasion and metastasis. <i>Oncogene</i> , 2017 , 36, 1925-1938	9.2	108
203	Daam2 driven degradation of VHL promotes gliomagenesis. <i>ELife</i> , 2017 , 6,	8.9	13
202	Genomic landscape and evolution of metastatic chromophobe renal cell carcinoma. <i>JCI Insight</i> , 2017 , 2,	9.9	72
201	Epithelial-to-mesenchymal transition drives a pro-metastatic Golgi compaction process through scaffolding protein PAQR11. <i>Journal of Clinical Investigation</i> , 2017 , 127, 117-131	15.9	49
200	Angiomotin regulates prostate cancer cell proliferation by signaling through the Hippo-YAP pathway. <i>Oncotarget</i> , 2017 , 8, 10145-10160	3.3	12
199	Macrophages promote the progression of premalignant mammary lesions to invasive cancer. <i>Oncotarget</i> , 2017 , 8, 50731-50746	3.3	47
198	miR-33a is a tumor suppressor microRNA that is decreased in prostate cancer. <i>Oncotarget</i> , 2017 , 8, 6024	43:502	5 6 7
197	SAR405838: A Novel and Potent Inhibitor of the MDM2:p53 Axis for the Treatment of Dedifferentiated Liposarcoma. <i>Clinical Cancer Research</i> , 2016 , 22, 1150-60	12.9	62
196	HDAC Inhibition for the Treatment of Epithelioid Sarcoma: Novel Cross Talk Between Epigenetic Components. <i>Molecular Cancer Research</i> , 2016 , 14, 35-43	6.6	15

(2016-2016)

195	The miR-200 family and the miR-183~96~182 cluster target Foxf2 to inhibit invasion and metastasis in lung cancers. <i>Oncogene</i> , 2016 , 35, 173-86	9.2	91
194	MNX1 Is Oncogenically Upregulated in African-American Prostate Cancer. <i>Cancer Research</i> , 2016 , 76, 6290-6298	10.1	35
193	Temporal Profiling of Astrocyte Precursors Reveals Parallel Roles for Asef during Development and after Injury. <i>Journal of Neuroscience</i> , 2016 , 36, 11904-11917	6.6	24
192	Aurora kinase-A overexpression in mouse mammary epithelium induces mammary adenocarcinomas harboring genetic alterations shared with human breast cancer. <i>Carcinogenesis</i> , 2016 , 37, 1180-1189	4.6	19
191	Neuronal Trans-Differentiation in Prostate Cancer Cells. <i>Prostate</i> , 2016 , 76, 1312-25	4.2	16
190	A genetic cell context-dependent role for ZEB1 in lung cancer. <i>Nature Communications</i> , 2016 , 7, 12231	17.4	44
189	Dysregulation of miRNAs-COUP-TFII-FOXM1-CENPF axis contributes to the metastasis of prostate cancer. <i>Nature Communications</i> , 2016 , 7, 11418	17.4	74
188	Unique Transcriptomic Profile of Collecting Duct Carcinomas Relative to Upper Tract Urothelial Carcinomas and other Kidney Carcinomas. <i>Scientific Reports</i> , 2016 , 6, 30988	4.9	27
187	Functional annotation of rare gene aberration drivers of pancreatic cancer. <i>Nature Communications</i> , 2016 , 7, 10500	17.4	47
186	Multilevel Genomics-Based Taxonomy of Renal Cell Carcinoma. <i>Cell Reports</i> , 2016 , 14, 2476-89	10.6	228
185	DNA Methylation Signature Reveals Cell Ontogeny of Renal Cell Carcinomas. <i>Clinical Cancer Research</i> , 2016 , 22, 6236-6246	12.9	30
184	Identification of miR-139-5p as a saliva biomarker for tongue squamous cell carcinoma: a pilot study. <i>Cellular Oncology (Dordrecht)</i> , 2016 , 39, 187-93	7.2	58
183	Cancer-Associated Fibroblasts Induce a Collagen Cross-link Switch in Tumor Stroma. <i>Molecular Cancer Research</i> , 2016 , 14, 287-95	6.6	114
182	Ampullary Cancers Harbor ELF3 Tumor Suppressor Gene Mutations and Exhibit Frequent WNT Dysregulation. <i>Cell Reports</i> , 2016 , 14, 907-919	10.6	75
181	HER2 Signaling Drives DNA Anabolism and Proliferation through SRC-3 Phosphorylation and E2F1-Regulated Genes. <i>Cancer Research</i> , 2016 , 76, 1463-75	10.1	24
180	Blockade of AP-1 Potentiates Endocrine Therapy and Overcomes Resistance. <i>Molecular Cancer Research</i> , 2016 , 14, 470-81	6.6	27
179	Uterine ALK3 is essential during the window of implantation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E387-95	11.5	31
178	An Integrated Metabolic Atlas of Clear Cell Renal Cell Carcinoma. <i>Cancer Cell</i> , 2016 , 29, 104-116	24.3	335

177	Comprehensive Molecular Characterization of Papillary Renal-Cell Carcinoma. <i>New England Journal of Medicine</i> , 2016 , 374, 135-45	59.2	753
176	The role of miR-145 in stem cell characteristics of human laryngeal squamous cell carcinoma Hep-2 cells. <i>Tumor Biology</i> , 2016 , 37, 4183-92	2.9	26
175	Notch promotes tumor metastasis in a prostate-specific Pten-null mouse model. <i>Journal of Clinical Investigation</i> , 2016 , 126, 2626-41	15.9	38
174	Comprehensive characterization of 412 muscle invasive urothelial carcinomas: Final analysis of The Cancer Genome Atlas (TCGA) project <i>Journal of Clinical Oncology</i> , 2016 , 34, 405-405	2.2	1
173	Ten-Eleven Translocation Genes are Downregulated in Endometriosis. <i>Current Molecular Medicine</i> , 2016 , 16, 288-98	2.5	13
172	Transcriptomic profiling of collecting duct carcinoma to reveal metabolic and immune aberrations <i>Journal of Clinical Oncology</i> , 2016 , 34, 4572-4572	2.2	
171	The microRNA-200/Zeb1 axis regulates ECM-dependent 🛭-integrin/FAK signaling, cancer cell invasion and metastasis through CRKL. <i>Scientific Reports</i> , 2016 , 6, 18652	4.9	50
170	Identification of microRNA profile specific to cancer stem-like cells directly isolated from human larynx cancer specimens. <i>BMC Cancer</i> , 2016 , 16, 853	4.8	12
169	Endometrial Expression of Steroidogenic Factor 1 Promotes Cystic Glandular Morphogenesis. <i>Molecular Endocrinology</i> , 2016 , 30, 518-32		15
168	Fatty Acid Oxidation-Driven Src Links Mitochondrial Energy Reprogramming and Oncogenic Properties in Triple-Negative Breast Cancer. <i>Cell Reports</i> , 2016 , 14, 2154-2165	10.6	159
168167		10.6	159
	Properties in Triple-Negative Breast Cancer. <i>Cell Reports</i> , 2016 , 14, 2154-2165		
167	Properties in Triple-Negative Breast Cancer. <i>Cell Reports</i> , 2016 , 14, 2154-2165 International cancer seminars: a focus on kidney cancer. <i>Annals of Oncology</i> , 2016 , 27, 1382-5 Non-Cell-Autonomous Regulation of Prostate Epithelial Homeostasis by Androgen Receptor.	10.3	11
167 166	Properties in Triple-Negative Breast Cancer. <i>Cell Reports</i> , 2016 , 14, 2154-2165 International cancer seminars: a focus on kidney cancer. <i>Annals of Oncology</i> , 2016 , 27, 1382-5 Non-Cell-Autonomous Regulation of Prostate Epithelial Homeostasis by Androgen Receptor. <i>Molecular Cell</i> , 2016 , 63, 976-89 FOXA1 overexpression mediates endocrine resistance by altering the ER transcriptome and IL-8 expression in ER-positive breast cancer. <i>Proceedings of the National Academy of Sciences of the</i>	10.3	11 52
167 166 165	International cancer seminars: a focus on kidney cancer. <i>Annals of Oncology</i> , 2016 , 27, 1382-5 Non-Cell-Autonomous Regulation of Prostate Epithelial Homeostasis by Androgen Receptor. <i>Molecular Cell</i> , 2016 , 63, 976-89 FOXA1 overexpression mediates endocrine resistance by altering the ER transcriptome and IL-8 expression in ER-positive breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E6600-E6609 Role of neoplastic monocyte-derived fibrocytes in primary myelofibrosis. <i>Journal of Experimental</i>	10.3 17.6 11.5	11 52 91
167166165164	International cancer seminars: a focus on kidney cancer. <i>Annals of Oncology</i> , 2016 , 27, 1382-5 Non-Cell-Autonomous Regulation of Prostate Epithelial Homeostasis by Androgen Receptor. <i>Molecular Cell</i> , 2016 , 63, 976-89 FOXA1 overexpression mediates endocrine resistance by altering the ER transcriptome and IL-8 expression in ER-positive breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E6600-E6609 Role of neoplastic monocyte-derived fibrocytes in primary myelofibrosis. <i>Journal of Experimental Medicine</i> , 2016 , 213, 1723-40 NKD2, a negative regulator of Wnt signaling, suppresses tumor growth and metastasis in	10.3 17.6 11.5	11 52 91
167166165164163	International cancer seminars: a focus on kidney cancer. <i>Annals of Oncology</i> , 2016 , 27, 1382-5 Non-Cell-Autonomous Regulation of Prostate Epithelial Homeostasis by Androgen Receptor. <i>Molecular Cell</i> , 2016 , 63, 976-89 FOXA1 overexpression mediates endocrine resistance by altering the ER transcriptome and IL-8 expression in ER-positive breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E6600-E6609 Role of neoplastic monocyte-derived fibrocytes in primary myelofibrosis. <i>Journal of Experimental Medicine</i> , 2016 , 213, 1723-40 NKD2, a negative regulator of Wnt signaling, suppresses tumor growth and metastasis in osteosarcoma. <i>Oncogene</i> , 2015 , 34, 5069-79 Circulating and disseminated tumor cells from breast cancer patient-derived xenograft-bearing	10.3 17.6 11.5 16.6	11 52 91 93 80

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	, , , , ,	-5.7	
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76 75			
	phosphatase. <i>Cell</i> , 2011 , 144, 703-18 Metabolites of purine nucleoside phosphorylase (NP) in serum have the potential to delineate	56.2	214
75	phosphatase. <i>Cell</i> , 2011 , 144, 703-18 Metabolites of purine nucleoside phosphorylase (NP) in serum have the potential to delineate pancreatic adenocarcinoma. <i>PLoS ONE</i> , 2011 , 6, e17177 Deregulation of MYCN, LIN28B and LET7 in a molecular subtype of aggressive high-grade serous	56.2 3·7	214
75 74	phosphatase. <i>Cell</i> , 2011 , 144, 703-18 Metabolites of purine nucleoside phosphorylase (NP) in serum have the potential to delineate pancreatic adenocarcinoma. <i>PLoS ONE</i> , 2011 , 6, e17177 Deregulation of MYCN, LIN28B and LET7 in a molecular subtype of aggressive high-grade serous ovarian cancers. <i>PLoS ONE</i> , 2011 , 6, e18064 Metabolomic profiling reveals a role for androgen in activating amino acid metabolism and	56.2 3·7 3·7	21414143
75 74 73	Metabolites of purine nucleoside phosphorylase (NP) in serum have the potential to delineate pancreatic adenocarcinoma. <i>PLoS ONE</i> , 2011 , 6, e17177 Deregulation of MYCN, LIN28B and LET7 in a molecular subtype of aggressive high-grade serous ovarian cancers. <i>PLoS ONE</i> , 2011 , 6, e18064 Metabolomic profiling reveals a role for androgen in activating amino acid metabolism and methylation in prostate cancer cells. <i>PLoS ONE</i> , 2011 , 6, e21417 Int6 regulates both proteasomal degradation and translation initiation and is critical for proper	56.2 3·7 3·7	2141414365

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60 59	Functional microRNA involved in endometriosis. <i>Molecular Endocrinology</i> , 2011 , 25, 821-32 The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85	15.9	186
	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a	15.9	
59	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85 Proteomic Profiling of 112 Proteins in Chronic Myeloid Leukemia Patient Samples Using Reverse Phase Proteins Arrays (RPPA) Reveals Distinct Protein Expression Signatures Associated with		154
59 58	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85 Proteomic Profiling of 112 Proteins in Chronic Myeloid Leukemia Patient Samples Using Reverse Phase Proteins Arrays (RPPA) Reveals Distinct Protein Expression Signatures Associated with Advanced Phase Disease and the CD34+ Compartment. <i>Blood</i> , 2011 , 118, 2503-2503 Stem cell antigen-1 (sca-1) regulates mammary tumor development and cell migration. <i>PLoS ONE</i> ,	2.2	154
59 58 57	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85 Proteomic Profiling of 112 Proteins in Chronic Myeloid Leukemia Patient Samples Using Reverse Phase Proteins Arrays (RPPA) Reveals Distinct Protein Expression Signatures Associated with Advanced Phase Disease and the CD34+ Compartment. <i>Blood</i> , 2011 , 118, 2503-2503 Stem cell antigen-1 (sca-1) regulates mammary tumor development and cell migration. <i>PLoS ONE</i> , 2011 , 6, e27841 TAp63 suppresses metastasis through coordinate regulation of Dicer and miRNAs. <i>Nature</i> , 2010 ,	2.2	154 2 25
59 58 57 56	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85 Proteomic Profiling of 112 Proteins in Chronic Myeloid Leukemia Patient Samples Using Reverse Phase Proteins Arrays (RPPA) Reveals Distinct Protein Expression Signatures Associated with Advanced Phase Disease and the CD34+ Compartment. <i>Blood</i> , 2011 , 118, 2503-2503 Stem cell antigen-1 (sca-1) regulates mammary tumor development and cell migration. <i>PLoS ONE</i> , 2011 , 6, e27841 TAp63 suppresses metastasis through coordinate regulation of Dicer and miRNAs. <i>Nature</i> , 2010 , 467, 986-90 SAFB1 mediates repression of immune regulators and apoptotic genes in breast cancer cells.	2.2 3·7 50·4	154 2 25 353
59 58 57 56 55	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85 Proteomic Profiling of 112 Proteins in Chronic Myeloid Leukemia Patient Samples Using Reverse Phase Proteins Arrays (RPPA) Reveals Distinct Protein Expression Signatures Associated with Advanced Phase Disease and the CD34+ Compartment. <i>Blood</i> , 2011 , 118, 2503-2503 Stem cell antigen-1 (sca-1) regulates mammary tumor development and cell migration. <i>PLoS ONE</i> , 2011 , 6, e27841 TAp63 suppresses metastasis through coordinate regulation of Dicer and miRNAs. <i>Nature</i> , 2010 , 467, 986-90 SAFB1 mediates repression of immune regulators and apoptotic genes in breast cancer cells. <i>Journal of Biological Chemistry</i> , 2010 , 285, 3608-3616 Molecular profiling uncovers a p53-associated role for microRNA-31 in inhibiting the proliferation	2.2 3·7 50·4	154 2 25 353 25

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23	Novel microRNA candidates and miRNA-mRNA pairs in embryonic stem (ES) cells. <i>PLoS ONE</i> , 2008 , 3, e2548	3.7	41
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21	A gene transcription signature of the Akt/mTOR pathway in clinical breast tumors. <i>Oncogene</i> , 2007 , 26, 4648-55	9.2	57
20	A gene transcription signature associated with hormone independence in a subset of both breast and prostate cancers. <i>BMC Genomics</i> , 2007 , 8, 199	4.5	10
19	Integrative genomics analysis reveals silencing of beta-adrenergic signaling by polycomb in prostate cancer. <i>Cancer Cell</i> , 2007 , 12, 419-31	24.3	185
18	MDA-MB-435 cells are derived from M14 melanoma cellsa loss for breast cancer, but a boon for melanoma research. <i>Breast Cancer Research and Treatment</i> , 2007 , 104, 13-9	4.4	300
17	Transcriptional profiling of non-small cell lung cancer cells with activating EGFR somatic mutations. <i>PLoS ONE</i> , 2007 , 2, e1226	3.7	26
16	Activation of mitogen-activated protein kinase in estrogen receptor alpha-positive breast cancer cells in vitro induces an in vivo molecular phenotype of estrogen receptor alpha-negative human breast tumors. <i>Cancer Research</i> , 2006 , 66, 3903-11	10.1	190

LIST OF PUBLICATIONS

15	When will tumor gene expression profiling be incorporated into clinical breast cancer decision making?. <i>Breast Cancer Research</i> , 2006 , 8, 302	8.3	1	
14	Genes regulated by estrogen in breast tumor cells in vitro are similarly regulated in vivo in tumor xenografts and human breast tumors. <i>Genome Biology</i> , 2006 , 7, R28	18.3	98	
13	A functional annotation of subproteomes in human plasma. <i>Proteomics</i> , 2005 , 5, 3506-19	4.8	77	
12	Analysis of tumor-host interactions by gene expression profiling of lung adenocarcinoma xenografts identifies genes involved in tumor formation. <i>Molecular Cancer Research</i> , 2005 , 3, 119-29	6.6	52	
11	Gene expression patterns define pathways correlated with loss of differentiation in lung adenocarcinomas. <i>FEBS Letters</i> , 2003 , 540, 167-70	3.8	14	
10	Profiling of pathway-specific changes in gene expression following growth of human cancer cell lines transplanted into mice. <i>Genome Biology</i> , 2003 , 4, R46	18.3	38	
9	Expression of matrix metalloproteinase 9 (MMP-9/gelatinase B) in adenocarcinomas strongly correlated with expression of immune response genes. <i>In Silico Biology</i> , 2003 , 3, 301-11	2	16	
8	Site-specific recombination of temperate Myxococcus xanthus phage Mx8: genetic elements required for integration. <i>Journal of Bacteriology</i> , 1999 , 181, 4050-61	3.5	33	
7	Mutations that confer resistance to 2-deoxyglucose reduce the specific activity of hexokinase from Myxococcus xanthus. <i>Journal of Bacteriology</i> , 1999 , 181, 2225-35	3.5	11	
6	A chaperone in the HSP70 family controls production of extracellular fibrils in Myxococcus xanthus. <i>Journal of Bacteriology</i> , 1998 , 180, 5357-68	3.5	59	
5	The aadA gene of plasmid R100 confers resistance to spectinomycin and streptomycin in Myxococcus xanthus. <i>Journal of Bacteriology</i> , 1998 , 180, 6757-60	3.5	11	
4	Whole genome and RNA sequencing of 1,220 cancers reveals hundreds of genes deregulated by rearrangement of cis-regulatory elements		4	
3	Comprehensive Molecular Characterization of Mitochondrial Genomes in Human Cancers		6	
2	Genomic basis for RNA alterations revealed by whole-genome analyses of 27 cancer types		10	
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