Mark A Rubin

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.

67,812 126 496 252 h-index g-index papers citations 10.8 77,903 7.11 531 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
496	What Experts Think About Prostate Cancer Management During the COVID-19 Pandemic: Report from the Advanced Prostate Cancer Consensus Conference 2021 <i>European Urology</i> , 2022 ,	10.2	2
495	Comparative genomics of primary prostate cancer and paired metastases: insights from 12 molecular case studies <i>Journal of Pathology</i> , 2022 ,	9.4	1
494	Alterations in homologous recombination repair genes in prostate cancer brain metastases <i>Nature Communications</i> , 2022 , 13, 2400	17.4	3
493	The evolving landscape of prostate cancer somatic mutations. <i>Prostate</i> , 2022 , 82,	4.2	0
492	PARP Inhibition in Prostate Cancer With Homologous Recombination Repair Alterations. <i>JCO Precision Oncology</i> , 2021 , 5,	3.6	3
491	G3BP1 inhibits Cul3 to amplify AR signaling and promote prostate cancer. <i>Nature Communications</i> , 2021 , 12, 6662	17.4	3
490	Dynamic prostate cancer transcriptome analysis delineates the trajectory to disease progression. <i>Nature Communications</i> , 2021 , 12, 7033	17.4	4
489	Comparative pathology of dog and human prostate cancer. Veterinary Medicine and Science, 2021,	2.1	5
488	Extracellular Matrix in Synthetic Hydrogel-Based Prostate Cancer Organoids Regulate Therapeutic Response to EZH2 and DRD2 Inhibitors. <i>Advanced Materials</i> , 2021 , e2100096	24	3
487	NKX3.1 Localization to Mitochondria Suppresses Prostate Cancer Initiation. <i>Cancer Discovery</i> , 2021 , 11, 2316-2333	24.4	8
486	Targeting the epichaperome as an effective precision medicine approach in a novel PML-SYK fusion acute myeloid leukemia. <i>Npj Precision Oncology</i> , 2021 , 5, 44	9.8	6
485	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
484	Histological and immunohistochemical investigation of canine prostate carcinoma with identification of common intraductal carcinoma component. <i>Veterinary and Comparative Oncology</i> , 2021 ,	2.5	3
483	PI5P4Ks drive metabolic homeostasis through peroxisome-mitochondria interplay. <i>Developmental Cell</i> , 2021 , 56, 1661-1676.e10	10.2	7
482	CD38 in Advanced Prostate Cancers. European Urology, 2021 , 79, 736-746	10.2	0
481	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
480	The 2019 Genitourinary Pathology Society (GUPS) White Paper on Contemporary Grading of Prostate Cancer. <i>Archives of Pathology and Laboratory Medicine</i> , 2021 , 145, 461-493	5	41

(2020-2021)

479	Practice patterns related to prostate cancer grading: results of a 2019 Genitourinary Pathology Society clinician survey. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2021 , 39, 295.e1-295.e8	2.8	1
478	Loss and revival of androgen receptor signaling in advanced prostate cancer. <i>Oncogene</i> , 2021 , 40, 1205-1	9216	13
477	Prostate cancer hijacks the microenvironment. <i>Nature Cell Biology</i> , 2021 , 23, 3-5	23.4	6
476	Dual functions of SPOP and ERG dictate androgen therapy responses in prostate cancer. <i>Nature Communications</i> , 2021 , 12, 734	17.4	10
475	Molecular medicine tumor board: whole-genome sequencing to inform on personalized medicine for a man with advanced prostate cancer. <i>Prostate Cancer and Prostatic Diseases</i> , 2021 , 24, 786-793	6.2	3
474	Patient-derived xenografts and organoids model therapy response in prostate cancer. <i>Nature Communications</i> , 2021 , 12, 1117	17.4	18
473	Co-occurrence and mutual exclusivity: what cross-cancer mutation patterns can tell us. <i>Trends in Cancer</i> , 2021 , 7, 823-836	12.5	3
472	The long noncoding RNA H19 regulates tumor plasticity in neuroendocrine prostate cancer <i>Nature Communications</i> , 2021 , 12, 7349	17.4	10
471	Role of specialized composition of SWI/SNF complexes in prostate cancer lineage plasticity. <i>Nature Communications</i> , 2020 , 11, 5549	17.4	31
470	Report From the International Society of Urological Pathology (ISUP) Consultation Conference on Molecular Pathology of Urogenital Cancers. I. Molecular Biomarkers in Prostate Cancer. <i>American Journal of Surgical Pathology</i> , 2020 , 44, e15-e29	6.7	21
469	Fusions involving BCOR and CREBBP are rare events in infiltrating glioma. <i>Acta Neuropathologica Communications</i> , 2020 , 8, 80	7.3	3
468	Small Cell Carcinoma of the Ovary, Hypercalcemic Type (SCCOHT) beyond Mutations: A Comprehensive Genomic Analysis. <i>Cells</i> , 2020 , 9,	7.9	14
467	Management of Patients with Advanced Prostate Cancer: Report of the Advanced Prostate Cancer Consensus Conference 2019. <i>European Urology</i> , 2020 , 77, 508-547	10.2	155
466	Pathway and network analysis of more than 2500 whole cancer genomes. <i>Nature Communications</i> , 2020 , 11, 729	17.4	38
465	Analyses of non-coding somatic drivers in 2,658 cancer whole genomes. <i>Nature</i> , 2020 , 578, 102-111	50.4	220
464	Integrative multiplatform molecular profiling of benign prostatic hyperplasia identifies distinct subtypes. <i>Nature Communications</i> , 2020 , 11, 1987	17.4	14
463	Performance Characteristics of a Targeted Sequencing Platform for Simultaneous Detection of Single Nucleotide Variants, Insertions/Deletions, Copy Number Alterations, and Gene Fusions in Cancer Genome. <i>Archives of Pathology and Laboratory Medicine</i> , 2020 , 144, 1535-1546	5	4
462	A MYC and RAS co-activation signature in localized prostate cancer drives bone metastasis and castration resistance. <i>Nature Cancer</i> , 2020 , 1, 1082-1096	15.4	18

461	Clinical deployment of AI for prostate cancer diagnosis. The Lancet Digital Health, 2020, 2, e383-e384	14.4	3
460	Common germline-somatic variant interactions in advanced urothelial cancer. <i>Nature Communications</i> , 2020 , 11, 6195	17.4	6
459	Impact of Lineage Plasticity to and from a Neuroendocrine Phenotype on Progression and Response in Prostate and Lung Cancers. <i>Molecular Cell</i> , 2020 , 80, 562-577	17.6	12
458	Sex differences in oncogenic mutational processes. <i>Nature Communications</i> , 2020 , 11, 4330	17.4	23
457	SETting Up for Epigenetic Regulation of Advanced Prostate Cancer. Cancer Cell, 2020, 38, 309-311	24.3	1
456	GRAM: A GeneRAlized Model to predict the molecular effect of a non-coding variant in a cell-type specific manner. <i>PLoS Genetics</i> , 2019 , 15, e1007860	6	1
455	Cancer-Specific Thresholds Adjust for Whole Exome Sequencing-based Tumor Mutational Burden Distribution. <i>JCO Precision Oncology</i> , 2019 , 3,	3.6	8
454	Integrative Molecular Analysis of Patients With Advanced and Metastatic Cancer. <i>JCO Precision Oncology</i> , 2019 , 3,	3.6	15
453	Genomic correlates of clinical outcome in advanced prostate cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 11428-11436	11.5	383
452	Characterization of the ERG-regulated Kinome in Prostate Cancer Identifies TNIK as a Potential Therapeutic Target. <i>Neoplasia</i> , 2019 , 21, 389-400	6.4	10
451	CHD1 Loss Alters AR Binding at Lineage-Specific Enhancers and Modulates Distinct Transcriptional Programs to Drive Prostate Tumorigenesis. <i>Cancer Cell</i> , 2019 , 35, 603-617.e8	24.3	29
450	DNA Hypermethylation Encroachment at CpG Island Borders in Cancer Is Predisposed by H3K4 Monomethylation Patterns. <i>Cancer Cell</i> , 2019 , 35, 297-314.e8	24.3	34
449	Proteomic and genomic signatures of repeat instability in cancer and adjacent normal tissues. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 16987-16996	6 ^{11.5}	6
448	The Role of Lineage Plasticity in Prostate Cancer Therapy Resistance. <i>Clinical Cancer Research</i> , 2019 , 25, 6916-6924	12.9	94
447	Upper tract urothelial carcinoma has a luminal-papillary T-cell depleted contexture and activated FGFR3 signaling. <i>Nature Communications</i> , 2019 , 10, 2977	17.4	71
446	A Phase II Trial of the Aurora Kinase A Inhibitor Alisertib for Patients with Castration-resistant and Neuroendocrine Prostate Cancer: Efficacy and Biomarkers. <i>Clinical Cancer Research</i> , 2019 , 25, 43-51	12.9	110
445	Prostate Cancer Research at the Crossroads. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2019 , 9,	5.4	2
444	The Genomics of Prostate Cancer: A Historic Perspective. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2019 , 9,	5.4	10

(2018-2018)

443	Rapid autopsy of a patient with recurrent anaplastic ependymoma. <i>Palliative and Supportive Care</i> , 2018 , 16, 238-242	2.5	2
442	Loss of an Androgen-Inactivating and Isoform-Specific HSD17B4 Splice Form Enables Emergence of Castration-Resistant Prostate Cancer. <i>Cell Reports</i> , 2018 , 22, 809-819	10.6	23
441	Clinical Outcome of Prostate Cancer Patients with Germline DNA Repair Mutations: Retrospective Analysis from an International Study. <i>European Urology</i> , 2018 , 73, 687-693	10.2	70
440	Bone biopsy protocol for advanced prostate cancer in the era of precision medicine. <i>Cancer</i> , 2018 , 124, 1008-1015	6.4	24
439	The Genomics of Prostate Cancer: emerging understanding with technologic advances. <i>Modern Pathology</i> , 2018 , 31, S1-11	9.8	31
438	Phosphatidylinositol-5-Phosphate 4-Kinases Regulate Cellular Lipid Metabolism By Facilitating Autophagy. <i>Molecular Cell</i> , 2018 , 70, 531-544.e9	17.6	35
437	The long tail of oncogenic drivers in prostate cancer. <i>Nature Genetics</i> , 2018 , 50, 645-651	36.3	380
436	Characterization of CD34-deficient myofibroblastomas of the breast. <i>Breast Journal</i> , 2018 , 24, 55-61	1.2	4
435	Management of Patients with Advanced Prostate Cancer: The Report of the Advanced Prostate Cancer Consensus Conference APCCC 2017. <i>European Urology</i> , 2018 , 73, 178-211	10.2	313
434	SPOP-Mutated/CHD1-Deleted Lethal Prostate Cancer and Abiraterone Sensitivity. <i>Clinical Cancer Research</i> , 2018 , 24, 5585-5593	12.9	74
433	Sequence of events in prostate cancer. <i>Nature</i> , 2018 , 560, 557-559	50.4	3
432	Suppression of insulin feedback enhances the efficacy of PI3K inhibitors. <i>Nature</i> , 2018 , 560, 499-503	50.4	277
431	Patient derived organoids to model rare prostate cancer phenotypes. <i>Nature Communications</i> , 2018 , 9, 2404	17.4	149
430	Transcriptomic heterogeneity in multifocal prostate cancer. JCI Insight, 2018, 3,	9.9	41
429	Differential impact of RB status on E2F1 reprogramming in human cancer. <i>Journal of Clinical Investigation</i> , 2018 , 128, 341-358	15.9	58
428	SPOP mutation drives prostate neoplasia without stabilizing oncogenic transcription factor ERG. <i>Journal of Clinical Investigation</i> , 2018 , 128, 381-386	15.9	23
427	Upper tract urothelial carcinoma is non-basal and T-cell depleted <i>Journal of Clinical Oncology</i> , 2018 , 36, 4525-4525	2.2	1
426	Molecular and clinical implications of CHD1 loss and SPOP mutations in advanced prostate cancer Journal of Clinical Oncology, 2018 , 36, 5064-5064	2.2	2

425	Integrative molecular profiling challenges robustness of prognostic signature scores in multifocal prostate cancer <i>Journal of Clinical Oncology</i> , 2018 , 36, 96-96	2.2	1
424	Clinical outcome of patients with germline DNA repair mutations: Results from a retrospective international study <i>Journal of Clinical Oncology</i> , 2018 , 36, 218-218	2.2	
423	Targeting the Epichaperome As an Effective Precision Medicine Approach in a Novel PML-SYK Fusion Acute Myeloid Leukemia. <i>Blood</i> , 2018 , 132, 1435-1435	2.2	1
422	Clinical and Genomic Characterization of Treatment-Emergent Small-Cell Neuroendocrine Prostate Cancer: A Multi-institutional Prospective Study. <i>Journal of Clinical Oncology</i> , 2018 , 36, 2492-2503	2.2	271
421	Impact of the SPOP Mutant Subtype on the Interpretation of Clinical Parameters in Prostate Cancer. <i>JCO Precision Oncology</i> , 2018 , 2018,	3.6	17
420	Immunogenomic analyses associate immunological alterations with mismatch repair defects in prostate cancer. <i>Journal of Clinical Investigation</i> , 2018 , 128, 4441-4453	15.9	84
419	NSD2 is a conserved driver of metastatic prostate cancer progression. <i>Nature Communications</i> , 2018 , 9, 5201	17.4	44
418	TET2 Deficiency Causes Germinal Center Hyperplasia, Impairs Plasma Cell Differentiation, and Promotes B-cell Lymphomagenesis. <i>Cancer Discovery</i> , 2018 , 8, 1632-1653	24.4	77
417	Linking prostate cancer cell AR heterogeneity to distinct castration and enzalutamide responses. <i>Nature Communications</i> , 2018 , 9, 3600	17.4	60
416	Prostate Power Play: Does Accelerate -Deficient Cancer Progression?. Cancer Discovery, 2018 , 8, 682-6	8524.4	4
415	The long noncoding RNA landscape of neuroendocrine prostate cancer and its clinical implications. <i>GigaScience</i> , 2018 , 7,	7.6	35
414	SOX2 promotes lineage plasticity and antiandrogen resistance in TP53- and RB1-deficient prostate cancer. <i>Science</i> , 2017 , 355, 84-88	33.3	491
413	Racial Variation in the Utility of Urinary Biomarkers PCA3 and T2ERG in a Large Multicenter Study. Journal of Urology, 2017 , 198, 42-49	2.5	10
412	Non-coding genetic variation in cancer. Current Opinion in Systems Biology, 2017, 1, 9-15	3.2	23
411	Transdifferentiation as a Mechanism of Treatment Resistance in a Mouse Model of Castration-Resistant Prostate Cancer. <i>Cancer Discovery</i> , 2017 , 7, 736-749	24.4	182
410	Transplantation of engineered organoids enables rapid generation of metastatic mouse models of colorectal cancer. <i>Nature Biotechnology</i> , 2017 , 35, 577-582	44.5	137
409	Exome Sequencing of African-American Prostate Cancer Reveals Loss-of-Function Mutations. <i>Cancer Discovery</i> , 2017 , 7, 973-983	24.4	65

407	Prostate cancer: Clinical hallmarks in whole cancer genomes. <i>Nature Reviews Clinical Oncology</i> , 2017 , 14, 265-266	19.4	1
406	Personalized and Cancer Models to Guide Precision Medicine. <i>Cancer Discovery</i> , 2017 , 7, 462-477	24.4	477
405	SPOP Mutation Drives Prostate Tumorigenesis In Vivo through Coordinate Regulation of PI3K/mTOR and AR Signaling. <i>Cancer Cell</i> , 2017 , 31, 436-451	24.3	116
404	Quantification of mutant SPOP proteins in prostate cancer using mass spectrometry-based targeted proteomics. <i>Journal of Translational Medicine</i> , 2017 , 15, 175	8.5	5
403	Identification of novel prostate cancer drivers using RegNetDriver: a framework for integration of genetic and epigenetic alterations with tissue-specific regulatory network. <i>Genome Biology</i> , 2017 , 18, 141	18.3	20
402	Next-Generation Rapid Autopsies Enable Tumor Evolution Tracking and Generation of Preclinical Models. <i>JCO Precision Oncology</i> , 2017 , 2017,	3.6	23
401	Prostate cancer-associated SPOP mutations confer resistance to BET inhibitors through stabilization of BRD4. <i>Nature Medicine</i> , 2017 , 23, 1063-1071	50.5	169
400	Aberrant Activation of a Gastrointestinal Transcriptional Circuit in Prostate Cancer Mediates Castration Resistance. <i>Cancer Cell</i> , 2017 , 32, 792-806.e7	24.3	39
399	A germline FANCA alteration that is associated with increased sensitivity to DNA damaging agents. Journal of Physical Education and Sports Management, 2017, 3,	2.8	15
398	Inherited determinants of early recurrent somatic mutations in prostate cancer. <i>Nature Communications</i> , 2017 , 8, 48	17.4	16
397	The Master Neural Transcription Factor BRN2 Is an Androgen Receptor-Suppressed Driver of Neuroendocrine Differentiation in Prostate Cancer. <i>Cancer Discovery</i> , 2017 , 7, 54-71	24.4	173
396	The cancer precision medicine knowledge base for structured clinical-grade mutations and interpretations. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2017 , 24, 513-519	8.6	66
395	The Emergence of Precision Urologic Oncology: A Collaborative Review on Biomarker-driven Therapeutics. <i>European Urology</i> , 2017 , 71, 237-246	10.2	41
394	DNA Repair in Prostate Cancer: Biology and Clinical Implications. <i>European Urology</i> , 2017 , 71, 417-425	10.2	132
393	Re: Prognostic Significance of Percentage and Architectural Types of Contemporary Gleason Pattern 4 Prostate Cancer in Radical Prostatectomy. <i>European Urology</i> , 2017 , 71, 301	10.2	1
392	Biology and evolution of poorly differentiated neuroendocrine tumors. <i>Nature Medicine</i> , 2017 , 23, 1-10	50.5	109
391	On-site Cytology for Development of Patient-Derived Three-dimensional Organoid Cultures - A Pilot Study. <i>Anticancer Research</i> , 2017 , 37, 1569-1573	2.3	5
390	Punctuated evolution of copy-number alterations to define two molecular subtypes of muscle-invasive urothelial carcinoma <i>Journal of Clinical Oncology</i> , 2017 , 35, 299-299	2.2	

389	The long tail of significantly mutated genes in prostate cancer <i>Journal of Clinical Oncology</i> , 2017 , 35, 131-131	2.2	
388	Clonal evolution of chemotherapy-resistant urothelial carcinoma. <i>Nature Genetics</i> , 2016 , 48, 1490-1499	36.3	161
387	Development and validation of a whole-exome sequencing test for simultaneous detection of point mutations, indels and copy-number alterations for precision cancer care. <i>Npj Genomic Medicine</i> , 2016 , 1,	6.2	51
386	Prostate cancer risk regions at 8q24 and 17q24 are differentially associated with somatic TMPRSS2:ERG fusion status. <i>Human Molecular Genetics</i> , 2016 , 25, 5490-5499	5.6	6
385	A Computational Drug Repositioning Approach for Targeting Oncogenic Transcription Factors. <i>Cell Reports</i> , 2016 , 15, 2348-56	10.6	25
384	Patient-Level DNA Damage and Repair Pathway Profiles and Prognosis After Prostatectomy for High-Risk Prostate Cancer. <i>JAMA Oncology</i> , 2016 , 2, 471-80	13.4	38
383	Role of non-coding sequence variants in cancer. <i>Nature Reviews Genetics</i> , 2016 , 17, 93-108	30.1	301
382	Genomic Correlates to the Newly Proposed Grading Prognostic Groups for Prostate Cancer. <i>European Urology</i> , 2016 , 69, 557-560	10.2	50
381	Divergent clonal evolution of castration-resistant neuroendocrine prostate cancer. <i>Nature Medicine</i> , 2016 , 22, 298-305	50.5	775
380	Clonal evaluation of prostate cancer foci in biopsies with discontinuous tumor involvement by dual ERG/SPINK1 immunohistochemistry. <i>Modern Pathology</i> , 2016 , 29, 157-65	9.8	25
379	Inherited mutations in DNA repair genes in men with metastatic castration-resistant prostate cancer <i>Journal of Clinical Oncology</i> , 2016 , 34, 5009-5009	2.2	1
378	Clinical and genomic characterization of metastatic small cell/neuroendocrine prostate cancer (SCNC) and intermediate atypical prostate cancer (IAC): Results from the SU2C/PCF/AACRWest Coast Prostate Cancer Dream Team (WCDT) <i>Journal of Clinical Oncology</i> , 2016 , 34, 5019-5019	2.2	13
377	Generating a neoantigen map of advanced urothelial carcinoma by whole exome sequencing Journal of Clinical Oncology, 2016 , 34, 354-354	2.2	3
376	Serum Autoantibodies in Chronic Prostate Inflammation in Prostate Cancer Patients. <i>PLoS ONE</i> , 2016 , 11, e0147739	3.7	12
375	Integrated whole exome and RNA sequencing to reveal distinct genomic and transcriptomic landscape of upper tract urothelial carcinoma <i>Journal of Clinical Oncology</i> , 2016 , 34, 379-379	2.2	
374	Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. <i>New England Journal of Medicine</i> , 2016 , 375, 443-53	59.2	791
373	An emerging role for cytopathology in precision oncology. Cancer Cytopathology, 2016, 124, 167-73	3.9	14
372	The Molecular Evolution of Castration-resistant Prostate Cancer. European Urology Focus, 2016 , 2, 506-5	5 †3 :	23

(2015-2016)

Image-based computational quantification and visualization of genetic alterations and tumour heterogeneity. <i>Scientific Reports</i> , 2016 , 6, 24146	4.9	21
N-Myc Induces an EZH2-Mediated Transcriptional Program Driving Neuroendocrine Prostate Cancer. <i>Cancer Cell</i> , 2016 , 30, 563-577	24.3	256
Characterization of the leiomyomatous variant of myofibroblastoma: a rare subset distinct from other smooth muscle tumors of the breast. <i>Human Pathology</i> , 2016 , 58, 54-61	3.7	9
Chromatin to Clinic: The Molecular Rationale for PARP1 Inhibitor Function. <i>Molecular Cell</i> , 2015 , 58, 925	5- 3/1 .6	102
Multicenter Evaluation of the Prostate Health Index to Detect Aggressive Prostate Cancer in Biopsy NaWe Men. <i>Journal of Urology</i> , 2015 , 194, 65-72	2.5	110
MAGI3-AKT3 fusion in breast cancer amended. <i>Nature</i> , 2015 , 520, E11-2	50.4	20
Genomics and Epigenomics of Prostate Cancer 2015 , 149-170		
Genomic rearrangements in prostate cancer. Current Opinion in Urology, 2015, 25, 71-6	2.8	23
DNA-Repair Defects and Olaparib in Metastatic Prostate Cancer. <i>New England Journal of Medicine</i> , 2015 , 373, 1697-708	59.2	1345
The Placental Gene PEG10 Promotes Progression of Neuroendocrine Prostate Cancer. <i>Cell Reports</i> , 2015 , 12, 922-36	10.6	155
Toward a prostate cancer precision medicine. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2015 , 33, 73-4	2.8	7
Health: Make precision medicine work for cancer care. <i>Nature</i> , 2015 , 520, 290-1	50.4	45
SPOP mutation leads to genomic instability in prostate cancer. ELife, 2015, 4,	8.9	110
Integrative clinical genomics of advanced prostate cancer. <i>Cell</i> , 2015 , 161, 1215-1228	56.2	1765
Functional characterization of BC039389-GATM and KLK4-KRSP1 chimeric read-through transcripts which are up-regulated in renal cell cancer. <i>BMC Genomics</i> , 2015 , 16, 247	4.5	11
Whole-Exome Sequencing of Metastatic Cancer and Biomarkers of Treatment Response. <i>JAMA Oncology</i> , 2015 , 1, 466-74	13.4	207
Whole exome sequencing to reveal chemotherapy-driven evolution of platinum-resistant metastatic urothelial cancer <i>Journal of Clinical Oncology</i> , 2015 , 33, 4513-4513	2.2	1
Defining a molecular subclass of treatment resistant prostate cancer <i>Journal of Clinical Oncology</i> , 2015 , 33, 5004-5004	2.2	3
	heterogeneity. Scientific Reports, 2016, 6, 24146 N-Myc Induces an EZH2-Mediated Transcriptional Program Driving Neuroendocrine Prostate Cancer. Cancer Cell, 2016, 30, 563-577 Characterization of the leiomyomatous variant of myofibroblastoma: a rare subset distinct from other smooth muscle tumors of the breast. Human Pathology, 2016, 58, 54-61 Chromatin to Clinic: The Molecular Rationale for PARP1 Inhibitor Function. Molecular Cell, 2015, 58, 92. Multicenter Evaluation of the Prostate Health Index to Detect Aggressive Prostate Cancer in Biopsy Naße Men. Journal of Urology, 2015, 194, 65-72 MAGI3-AKT3 fusion in breast cancer amended. Nature, 2015, 520, E11-2 Genomics and Epigenomics of Prostate Cancer 2015, 149-170 Genomic rearrangements in prostate cancer. Current Opinion in Urology, 2015, 25, 71-6 DNA-Repair Defects and Olaparib in Metastatic Prostate Cancer. New England Journal of Medicine, 2015, 373, 1697-708 The Placental Gene PEG10 Promotes Progression of Neuroendocrine Prostate Cancer. Cell Reports, 2015, 12, 922-36 Toward a prostate cancer precision medicine. Urologic Oncology: Seminars and Original Investigations, 2015, 33, 73-4 Health: Make precision medicine work for cancer care. Nature, 2015, 520, 290-1 SPOP mutation leads to genomic instability in prostate cancer. ELIfe, 2015, 4, Integrative clinical genomics of advanced prostate cancer. Cell, 2015, 161, 1215-1228 Functional characterization of BC039389-GATM and KLK4-KRSP1 chimeric read-through transcripts which are up-regulated in renal cell cancer. BMC Genomics, 2015, 16, 247 Whole-Exome Sequencing of Metastatic Cancer and Biomarkers of Treatment Response. JAMA Oncology, 2015, 1, 466-74 Whole exome sequencing to reveal chemotherapy-driven evolution of platinum-resistant metastatic urothelial cancer Journal of Clinical Oncology, 2015, 33, 4513-4513	heterogeneity. Scientific Reports, 2016, 6, 24146 N-Myc Induces an EZH2-Mediated Transcriptional Program Driving Neuroendocrine Prostate Cancer. Cancer Cell, 2016, 30, 563-577 Characterization of the leiomyomatous variant of myofibroblastoma: a rare subset distinct from other smooth muscle tumors of the breast. Human Pathology, 2016, 58, 54-61 37 Chromatin to Clinic: The Molecular Rationale for PARP1 Inhibitor Function. Molecular Cell, 2015, 58, 925-346 Multicenter Evaluation of the Prostate Health Index to Detect Aggressive Prostate Cancer in Biopsy NaWe Men. Journal of Urology, 2015, 194, 65-72 MAGI3-AKT3 fusion in breast cancer amended. Nature, 2015, 520, E11-2 Genomics and Epigenomics of Prostate Cancer 2015, 149-170 Genomic rearrangements in prostate cancer. Current Opinion in Urology, 2015, 25, 71-6 2.8 DNA-Repair Defects and Olaparib in Metastatic Prostate Cancer. New England Journal of Medicine, 2015, 373, 1697-708 The Placental Gene PEG10 Promotes Progression of Neuroendocrine Prostate Cancer. Cell Reports, 10.6 Toward a prostate cancer precision medicine. Urologic Oncology: Seminars and Original Investigations, 2015, 33, 73-4 Health: Make precision medicine work for cancer care. Nature, 2015, 520, 290-1 SPOP mutation leads to genomic instability in prostate cancer. ELife, 2015, 4, 89 Integrative clinical genomics of advanced prostate cancer. Cell, 2015, 161, 1215-1228 Functional characterization of BC039389-GATM and KLK4-KRSP1 chimeric read-through transcripts which are up-regulated in renal cell cancer. BMC Genomics, 2015, 16, 247 Whole-Exome Sequencing of Metastatic Cancer and Biomarkers of Treatment Response. JAMA Oncology, 2015, 1, 466-74 Whole exome sequencing to reveal chemotherapy-driven evolution of platinum-resistant metastatic urothelial cancer Journal of Clinical Oncology, 2015, 33, 4513-4513

353	Phenotypic characterization of circulating tumor cells (CTCs) from neuroendocrine prostate cancer (NEPC) and metastatic castration-resistant prostate cancer (mCRPC) patients to identify a novel diagnostic algorithm for the presence of NEPC <i>Journal of Clinical Oncology</i> , 2015 , 33, 197-197	2.2	2
352	Precision medicine program for whole-exome sequencing (WES) provides new insight on platinum sensitivity in advanced prostate cancer (PCa) <i>Journal of Clinical Oncology</i> , 2015 , 33, 158-158	2.2	1
351	Clonal heterogeneity in platinum-resistant metastatic urothelial cancer <i>Journal of Clinical Oncology</i> , 2015 , 33, 290-290	2.2	
350	SPOP mutations in prostate cancer across demographically diverse patient cohorts. <i>Neoplasia</i> , 2014 , 16, 14-20	6.4	113
349	PCAT-1, a long noncoding RNA, regulates BRCA2 and controls homologous recombination in cancer. <i>Cancer Research</i> , 2014 , 74, 1651-60	10.1	204
348	TMPRSS2:ERG gene fusion predicts subsequent detection of prostate cancer in patients with high-grade prostatic intraepithelial neoplasia. <i>Journal of Clinical Oncology</i> , 2014 , 32, 206-11	2.2	79
347	Insights into the mechanism of organ-specific cancer metastasis. Cancer Discovery, 2014, 4, 1262-4	24.4	6
346	MYB-NFIB gene fusion in adenoid cystic carcinoma of the breast with special focus paid to the solid variant with basaloid features. <i>Human Pathology</i> , 2014 , 45, 2270-80	3.7	56
345	Prostate cancer. Ubiquitylome analysis identifies dysregulation of effector substrates in SPOP-mutant prostate cancer. <i>Science</i> , 2014 , 346, 85-89	33.3	157
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(2013-2014)

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(2006-2006)

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10	H&E-stained Whole Slide Image Deep Learning Predicts SPOP Mutation State in Prostate Cancer		27	
9	Discovery and reporting of clinically-relevant germline variants in advanced cancer patients assessed using whole-exome sequencing		2	
8	Role of Specialized Composition of SWI/SNF Complexes in Prostate Cancer Lineage Plasticity		1	
7	Patient-derived xenografts and organoids model therapy response in prostate cancer		1	
6	The Genomic Landscape of Prostate Cancer Brain Metastases		2	
5	Extracellular Microenvironment in Patient-derived Hydrogel Organoids of Prostate Cancer Regulates Therapeutic Response		2	
4	Dual Functions of SPOP and ERG Dictate Androgen Therapy Responses in Prostate Cancer		1	
3	Discovery and characterization of coding and non-coding driver mutations in more than 2,500 whole cancer genomes		12	
2	Pathway and network analysis of more than 2,500 whole cancer genomes		4	
1	The WHO 2022 landscape of papillary and chromophobe renal cell carcinoma. <i>Histopathology</i> ,	7.3	О	