Matti Annala

List of Publications by Year in descending order

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315616 304602 6,140 37 22 38 h-index citations g-index papers 39 39 39 10303 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The Molecular Taxonomy of Primary Prostate Cancer. Cell, 2015, 163, 1011-1025.	13.5	2,435
2	The evolutionary history of lethal metastatic prostate cancer. Nature, 2015, 520, 353-357.	13.7	1,185
3	Circulating Tumor DNA Genomics Correlate with Resistance to Abiraterone and Enzalutamide in Prostate Cancer. Cancer Discovery, 2018, 8, 444-457.	7.7	376
4	Genomic Alterations in Cell-Free DNA and Enzalutamide Resistance in Castration-Resistant Prostate Cancer. JAMA Oncology, 2016, 2, 1598.	3.4	290
5	Concordance of Circulating Tumor DNA and Matched Metastatic Tissue Biopsy in Prostate Cancer. Journal of the National Cancer Institute, 2017, 109, .	3.0	288
6	Optimal sequencing of enzalutamide and abiraterone acetate plus prednisone in metastatic castration-resistant prostate cancer: a multicentre, randomised, open-label, phase 2, crossover trial. Lancet Oncology, The, 2019, 20, 1730-1739.	5.1	227
7	Treatment Outcomes and Tumor Loss of Heterozygosity in Germline DNA Repair–deficient Prostate Cancer. European Urology, 2017, 72, 34-42.	0.9	179
8	Circulating Tumor DNA Abundance and Potential Utility in De Novo Metastatic Prostate Cancer. European Urology, 2019, 75, 667-675.	0.9	131
9	Circulating Tumor DNA Reveals Clinically Actionable Somatic Genome of Metastatic Bladder Cancer. Clinical Cancer Research, 2017, 23, 6487-6497.	3.2	121
10	Integrative proteomics in prostate cancer uncovers robustness against genomic and transcriptomic aberrations during disease progression. Nature Communications, 2018, 9, 1176.	5.8	117
11	Plasma ctDNA is a tumor tissue surrogate and enables clinical-genomic stratification of metastatic bladder cancer. Nature Communications, 2021, 12, 184.	5.8	85
12	Impact of Therapy on Genomics and Transcriptomics in High-Risk Prostate Cancer Treated with Neoadjuvant Docetaxel and Androgen Deprivation Therapy. Clinical Cancer Research, 2017, 23, 6802-6811.	3.2	69
13	Transcriptome Sequencing Reveals <i>PCAT5</i> as a Novel ERG-Regulated Long Noncoding RNA in Prostate Cancer. Cancer Research, 2015, 75, 4026-4031.	0.4	68
14	Constitutively active androgen receptor splice variants AR-V3, AR-V7 and AR-V9 are co-expressed in castration-resistant prostate cancer metastases. British Journal of Cancer, 2018, 119, 347-356.	2.9	63
15	<i>BRCA2</i> , <i>ATM</i> , and <i>CDK12</i> Defects Differentially Shape Prostate Tumor Driver Genomics and Clinical Aggression. Clinical Cancer Research, 2021, 27, 1650-1662.	3.2	52
16	Activating AKT1 and PIK3CA Mutations in Metastatic Castration-Resistant Prostate Cancer. European Urology, 2020, 78, 834-844.	0.9	47
17	Evolution of Castration-Resistant Prostate Cancer in ctDNA during Sequential Androgen Receptor Pathway Inhibition. Clinical Cancer Research, 2021, 27, 4610-4623.	3.2	41
18	Epigenetically altered miRâ€193b targets cyclin D1 in prostate cancer. Cancer Medicine, 2015, 4, 1417-1425.	1.3	39

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19	The expression of AURKA is androgen regulated in castration-resistant prostate cancer. Scientific Reports, 2017, 7, 17978.	1.6	38
20	Morphologic and genomic characterization of urothelial to sarcomatoid transition in muscle-invasive bladder cancer. Urologic Oncology: Seminars and Original Investigations, 2019, 37, 826-836.	0.8	33
21	Plasma Circulating Tumor DNA and Clonal Hematopoiesis in Metastatic Renal Cell Carcinoma. Clinical Genitourinary Cancer, 2020, 18, 322-331.e2.	0.9	30
22	Microseminoprotein-Beta Expression in Different Stages of Prostate Cancer. PLoS ONE, 2016, 11, e0150241.	1.1	28
23	Evaluation of Commercial Circulating Tumor DNA Test in Metastatic Prostate Cancer. JCO Precision Oncology, 2019, 3, 1-9.	1.5	26
24	Integrated clinical, whole-genome, and transcriptome analysis of multisampled lethal metastatic prostate cancer. Journal of Physical Education and Sports Management, 2016, 2, a000752.	0.5	24
25	Recurrent SKIL-activating rearrangements in ETS-negative prostate cancer. Oncotarget, 2015, 6, 6235-6250.	0.8	23
26	Frequent mutation of the FOXA1 untranslated region in prostate cancer. Communications Biology, 2018, 1, 122.	2.0	21
27	DOT1Lâ€HES6 fusion drives androgen independent growth in prostate cancer. EMBO Molecular Medicine, 2014, 6, 1121-1123.	3.3	17
28	OUP accepted manuscript. Neuro-Oncology, 2017, 19, 1206-1216.	0.6	17
29	Moving Toward Personalized Care: Liquid Biopsy Predicts Response to Cisplatin in an Unusual Case of BRCA2-Null Neuroendocrine Prostate Cancer. Clinical Genitourinary Cancer, 2016, 14, e233-e236.	0.9	15
30	Morphologic and genomic characterization of urothelial to sarcomatoid transition in muscle-invasive bladder cancer. Urologic Oncology: Seminars and Original Investigations, 2019, 37, 573.e19-573.e29.	0.8	13
31	Segmentum: a tool for copy number analysis of cancer genomes. BMC Bioinformatics, 2017, 18, 215.	1.2	7
32	Genomic Features of Lung-Recurrent Hormone-Sensitive Prostate Cancer. JCO Precision Oncology, 2022, 6, e2100543.	1.5	7
33	Somatic Features of Response and Relapse in Non–muscle-invasive Bladder Cancer Treated with Bacillus Calmette-Guérin Immunotherapy. European Urology Oncology, 2022, 5, 677-686.	2.6	6
34	Whole-exome sequencing identifies germline mutation in <i>TP53</i> and <i>ATRX</i> in a child with genomically aberrant AT/RT and her mother with anaplastic astrocytoma. Journal of Physical Education and Sports Management, 2018, 4, a002246.	0.5	5
35	AR and ERG drive the expression of prostate cancer specific long noncoding RNAs. Oncogene, 2020, 39, 5241-5251.	2.6	4
36	Moderate-to-strong expression of FGFR3 and TP53 alterations in a subpopulation of choroid plexus tumors. Histology and Histopathology, 2020, 35, 673-680.	0.5	3

#	Article	IF	CITATIONS
37	Molecular Dissection of Complete Response to Receptor Tyrosine Kinase Inhibition in Type II Papillary Renal Cell Carcinoma. Clinical Genitourinary Cancer, 2017, 15, e145-e150.	0.9	1