

Julie Livingstone

List of Publications by Citations

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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.

19
papers

1,684
citations

12
h-index

22
g-index

22
ext. papers

2,269
ext. citations

23.5
avg, IF

3.53
L-index

#	Paper	IF	Citations
19	Genomic hallmarks of localized, non-indolent prostate cancer. <i>Nature</i> , 2017 , 541, 359-364	50.4	320
18	Spatial genomic heterogeneity within localized, multifocal prostate cancer. <i>Nature Genetics</i> , 2015 , 47, 736-45	36.3	306
17	Molecular landmarks of tumor hypoxia across cancer types. <i>Nature Genetics</i> , 2019 , 51, 308-318	36.3	255
16	Widespread and Functional RNA Circularization in Localized Prostate Cancer. <i>Cell</i> , 2019 , 176, 831-843.e23	36.2	214
15	Germline BRCA2 mutations drive prostate cancers with distinct evolutionary trajectories. <i>Nature Communications</i> , 2017 , 8, 13671	17.4	128
14	The Evolutionary Landscape of Localized Prostate Cancers Drives Clinical Aggression. <i>Cell</i> , 2018 , 173, 1003-1013.e15	56.2	115
13	A Prostate Cancer "Nimbusus": Genomic Instability and SCHLAP1 Dysregulation Underpin Aggression of Intraductal and Cribriform Subpathologies. <i>European Urology</i> , 2017 , 72, 665-674	10.2	98
12	The Proteogenomic Landscape of Curable Prostate Cancer. <i>Cancer Cell</i> , 2019 , 35, 414-427.e6	24.3	97
11	The genomic landscape of metastatic castration-resistant prostate cancers reveals multiple distinct genotypes with potential clinical impact. <i>Nature Communications</i> , 2019 , 10, 5251	17.4	66
10	Genome-wide germline correlates of the epigenetic landscape of prostate cancer. <i>Nature Medicine</i> , 2019 , 25, 1615-1626	50.5	25
9	Noncoding mutations target cis-regulatory elements of the FOXA1 plexus in prostate cancer. <i>Nature Communications</i> , 2020 , 11, 441	17.4	21
8	Mammary molecular portraits reveal lineage-specific features and progenitor cell vulnerabilities. <i>Journal of Cell Biology</i> , 2018 , 217, 2951-2974	7.3	20
7	Transcriptomic Heterogeneity of Gleason Grade Group 5 Prostate Cancer. <i>European Urology</i> , 2020 , 78, 327-332	10.2	9
6	Somatic driver mutation prevalence in 1844 prostate cancers identifies ZNRF3 loss as a predictor of metastatic relapse. <i>Nature Communications</i> , 2021 , 12, 6248	17.4	3
5	Immune-focused multi-omics analysis of prostate cancer: leukocyte Ig-Like receptors are associated with disease progression. <i>Onc Immunology</i> , 2020 , 9, 1851950	7.2	2
4	Tumor cell total mRNA expression shapes the molecular and clinical phenotype of cancer		2
3	Copy Number Profiles of Prostate Cancer in Men of Middle Eastern Ancestry. <i>Cancers</i> , 2021 , 13,	6.6	1

2	Decreased ATM Protein Expression Is Substantiated with PTEN Loss in Defining Aggressive Phenotype of Prostate Cancer Associated with Lethal Disease. <i>European Urology Open Science</i> , 2021 , 29, 93-101	0.9	1
1	The telomere length landscape of prostate cancer. <i>Nature Communications</i> , 2021 , 12, 6893	17.4	0