

# Julie Livingstone

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6426641/publications.pdf>

Version: 2024-02-01

20  
papers

2,677  
citations

623574

14  
h-index

752573

20  
g-index

22  
all docs

22  
docs citations

22  
times ranked

5324  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular landmarks of tumor hypoxia across cancer types. <i>Nature Genetics</i> , 2019, 51, 308-318.	9.4	480
2	Genomic hallmarks of localized, non-indolent prostate cancer. <i>Nature</i> , 2017, 541, 359-364.	13.7	462
3	Spatial genomic heterogeneity within localized, multifocal prostate cancer. <i>Nature Genetics</i> , 2015, 47, 736-745.	9.4	395
4	Widespread and Functional RNA Circularization in Localized Prostate Cancer. <i>Cell</i> , 2019, 176, 831-843.e22.	13.5	317
5	Germline BRCA2 mutations drive prostate cancers with distinct evolutionary trajectories. <i>Nature Communications</i> , 2017, 8, 13671.	5.8	182
6	The Evolutionary Landscape of Localized Prostate Cancers Drives Clinical Aggression. <i>Cell</i> , 2018, 173, 1003-1013.e15.	13.5	176
7	The Proteogenomic Landscape of Curable Prostate Cancer. <i>Cancer Cell</i> , 2019, 35, 414-427.e6.	7.7	168
8	A Prostate Cancer "Nimbus" of Genomic Instability and SCHLAP1 Dysregulation Underpin Aggression of Intraductal and Cribriform Subpathologies. <i>European Urology</i> , 2017, 72, 665-674.	0.9	142
9	The genomic landscape of metastatic castration-resistant prostate cancers reveals multiple distinct genotypes with potential clinical impact. <i>Nature Communications</i> , 2019, 10, 5251.	5.8	130
10	Noncoding mutations target cis-regulatory elements of the FOXA1 plexus in prostate cancer. <i>Nature Communications</i> , 2020, 11, 441.	5.8	51
11	Genome-wide germline correlates of the epigenetic landscape of prostate cancer. <i>Nature Medicine</i> , 2019, 25, 1615-1626.	15.2	45
12	Mammary molecular portraits reveal lineage-specific features and progenitor cell vulnerabilities. <i>Journal of Cell Biology</i> , 2018, 217, 2951-2974.	2.3	35
13	Estimation of tumor cell total mRNA expression in 15 cancer types predicts disease progression. <i>Nature Biotechnology</i> , 2022, 40, 1624-1633.	9.4	31
14	Transcriptomic Heterogeneity of Gleason Grade Group 5 Prostate Cancer. <i>European Urology</i> , 2020, 78, 327-332.	0.9	18
15	Somatic driver mutation prevalence in 1844 prostate cancers identifies ZNRF3 loss as a predictor of metastatic relapse. <i>Nature Communications</i> , 2021, 12, 6248.	5.8	15
16	Immune-focused multi-omics analysis of prostate cancer: leukocyte Ig-Like receptors are associated with disease progression. <i>Oncotmmunology</i> , 2020, 9, 1851950.	2.1	8
17	The telomere length landscape of prostate cancer. <i>Nature Communications</i> , 2021, 12, 6893.	5.8	7
18	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.	0.2	5

#	ARTICLE	IF	CITATIONS
19	Methylation Subtypes of Primary Prostate Cancer Predict Poor Prognosis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1473-1482.	1.1	4
20	Copy Number Profiles of Prostate Cancer in Men of Middle Eastern Ancestry. <i>Cancers</i> , 2021, 13, 2363.	1.7	1