

# Deli Liu

## List of Publications by Year in descending order

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29  
papers

1,204  
citations

623574

14  
h-index

642610

23  
g-index

34  
all docs

34  
docs citations

34  
times ranked

2425  
citing authors

#	ARTICLE	IF	CITATIONS
1	Elevated expression of the colony-stimulating factor 1 (CSF1) induces prostatic intraepithelial neoplasia dependent of epithelial-Gp130. <i>Oncogene</i> , 2022, , .	2.6	6
2	Tumor subtype defines distinct pathways of molecular and clinical progression in primary prostate cancer. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	17
3	Tumor size and genomic risk in localized prostate cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2021, 39, 434.e17-434.e22.	0.8	3
4	Reshaping of the androgen-driven chromatin landscape in normal prostate cells by early cancer drivers and effect on therapeutic sensitivity. <i>Cell Reports</i> , 2021, 36, 109625.	2.9	22
5	G3BP1 inhibits Cul3SPOP to amplify AR signaling and promote prostate cancer. <i>Nature Communications</i> , 2021, 12, 6662.	5.8	17
6	Whole genome bisulfite sequencing of human spermatozoa reveals differentially methylated patterns from type 2 diabetic patients. <i>Journal of Diabetes Investigation</i> , 2020, 11, 856-864.	1.1	15
7	Diversity in Androgen Receptor Action Among Treatment-naïve Prostate Cancers Is Reflected in Treatment Response Predictions and Molecular Subtypes. <i>European Urology Open Science</i> , 2020, 22, 34-44.	0.2	7
8	Prognostic value of the SPOP mutant genomic subclass in prostate cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2020, 38, 418-422.	0.8	8
9	Integrative multiplatform molecular profiling of benign prostatic hyperplasia identifies distinct subtypes. <i>Nature Communications</i> , 2020, 11, 1987.	5.8	29
10	MP51-03â€fPROSTATE TUMOR SIZE IS INDEPENDENTLY ASSOCIATED WITH GENOMIC RISK. <i>Journal of Urology</i> , 2020, 203, .	0.2	0
11	FOXA1 mutations alter pioneering activity, differentiation and prostate cancer phenotypes. <i>Nature</i> , 2019, 571, 408-412.	13.7	163
12	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.	7.7	70
13	MP68-12â€fIMPACT OF SPOP MUTATION ON PROSTATE CANCER PROGNOSIS. <i>Journal of Urology</i> , 2019, 201, .	0.2	0
14	Impact of the SPOP Mutant Subtype on the Interpretation of Clinical Parameters in Prostate Cancer. <i>JCO Precision Oncology</i> , 2018, 2018, 1-13.	1.5	29
15	SEG - A Software Program for Finding Somatic Copy Number Alterations in Whole Genome Sequencing Data of Cancer. <i>Computational and Structural Biotechnology Journal</i> , 2018, 16, 335-341.	1.9	7
16	Abstract 2259: Prostate cancer molecular subtype impacts interpretation of clinical parameters. , 2018, , .		0
17	Co-clinical Analysis of a Genetically Engineered Mouse Model and Human Prostate Cancer Reveals Significance of NKX3.1 Expression for Response to 5Î±-reductase Inhibition. <i>European Urology</i> , 2017, 72, 499-506.	0.9	16
18	SPOP Mutation Drives Prostate Tumorigenesis InÂVivo through Coordinate Regulation of PI3K/mTOR and AR Signaling. <i>Cancer Cell</i> , 2017, 31, 436-451.	7.7	152

#	ARTICLE	IF	CITATIONS
19	MP99-08 SPOP MUTANT PROSTATE CANCER IS NOT DRIVEN BY ERG STABILIZATION. Journal of Urology, 2017, 197, .	0.2	0
20	Identification of novel prostate cancer drivers using RegNetDriver: a framework for integration of genetic and epigenetic alterations with tissue-specific regulatory network. Genome Biology, 2017, 18, 141.	3.8	31
21	SPOP mutation drives prostate neoplasia without stabilizing oncogenic transcription factor ERG. Journal of Clinical Investigation, 2017, 128, 381-386.	3.9	29
22	MP04-16 CLINICAL ASSOCIATIONS WITH PROSTATE CANCER MOLECULAR FEATURES IN THE CANCER GENOME ATLAS (TCGA) DATA.. Journal of Urology, 2016, 195, .	0.2	0
23	Abstract A07: Spontaneous canine mammary cancers: A much-needed model for human basal-like breast cancer and an ideal system to understand myoepithelial cells. , 2016, , .		0
24	Abstract 4174: SPOP mutation drives tumorigenesis in mouse prostate - a novel model of ETS negative prostate cancer. , 2016, , .		0
25	Canine Spontaneous Head and Neck Squamous Cell Carcinomas Represent Their Human Counterparts at the Molecular Level. PLoS Genetics, 2015, 11, e1005277.	1.5	37
26	SPOP mutation leads to genomic instability in prostate cancer. ELife, 2015, 4, .	2.8	148
27	Bridger: a new framework for de novo transcriptome assembly using RNA-seq data. Genome Biology, 2015, 16, 30.	3.8	258
28	Abstract 5147: Canine spontaneous head and neck squamous cell carcinomas represent their human counterparts at the molecular level. , 2015, , .		0
29	Molecular Homology and Difference between Spontaneous Canine Mammary Cancer and Human Breast Cancer. Cancer Research, 2014, 74, 5045-5056.	0.4	110