

# Yajun Yi

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5810634/publications.pdf>

Version: 2024-02-01

13  
papers

416  
citations

840776

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1125743

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citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Genes Required for Enzalutamide Resistance in Castration-Resistant Prostate Cancer Cells <i>In Vitro</i> . <i>Molecular Cancer Therapeutics</i> , 2021, 20, 398-409.	4.1	17
2	Identification of a gene-expression predictor for diagnosis and personalized stratification of lupus patients. <i>PLoS ONE</i> , 2018, 13, e0198325.	2.5	7
3	Loss of FOXA1 Drives Sexually Dimorphic Changes in Urothelial Differentiation and Is an Independent Predictor of Poor Prognosis in Bladder Cancer. <i>American Journal of Pathology</i> , 2015, 185, 1385-1395.	3.8	60
4	Evaluation of public cancer datasets and signatures identifies TP53 mutant signatures with robust prognostic and predictive value. <i>BMC Cancer</i> , 2015, 15, 179.	2.6	15
5	A Murine Model of K-RAS and $\beta$ -Catenin Induced Renal Tumors Expresses High Levels of E2F1 and Resembles Human Wilms Tumor. <i>Journal of Urology</i> , 2015, 194, 1762-1770.	0.4	15
6	NF- $\kappa$ B Gene Signature Predicts Prostate Cancer Progression. <i>Cancer Research</i> , 2014, 74, 2763-2772.	0.9	99
7	FOXA1 deletion in luminal epithelium causes prostatic hyperplasia and alteration of differentiated phenotype. <i>Laboratory Investigation</i> , 2014, 94, 726-739.	3.7	39
8	LIM Domain Only-2 (LMO2) Induces T-Cell Leukemia by Two Distinct Pathways. <i>PLoS ONE</i> , 2014, 9, e85883.	2.5	46
9	SPARCL1 suppresses metastasis in prostate cancer. <i>Molecular Oncology</i> , 2013, 7, 1019-1030.	4.6	32
10	A Data Similarity-Based Strategy for Meta-analysis of Transcriptional Profiles in Cancer. <i>PLoS ONE</i> , 2013, 8, e54979.	2.5	4
11	Web-based interrogation of gene expression signatures using EXALT. <i>BMC Bioinformatics</i> , 2009, 10, 420.	2.6	11
12	Strategy for encoding and comparison of gene expression signatures. <i>Genome Biology</i> , 2007, 8, R133.	8.8	24
13	Coupled analysis of gene expression and chromosomal location. <i>Genomics</i> , 2005, 85, 401-412.	2.9	47