

# George Michalopoulos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11327495/publications.pdf>

Version: 2024-02-01

15  
papers

1,992  
citations

687363

13  
h-index

996975

15  
g-index

15  
all docs

15  
docs citations

15  
times ranked

3493  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene Expression Alterations in Prostate Cancer Predicting Tumor Aggression and Preceding Development of Malignancy. <i>Journal of Clinical Oncology</i> , 2004, 22, 2790-2799.	1.6	674
2	Gene expression profiles of prostate cancer reveal involvement of multiple molecular pathways in the metastatic process. <i>BMC Cancer</i> , 2007, 7, 64.	2.6	421
3	Gene expression analysis of prostate cancers. <i>Molecular Carcinogenesis</i> , 2002, 33, 25-35.	2.7	216
4	Glutathione Peroxidase 3, Deleted or Methylated in Prostate Cancer, Suppresses Prostate Cancer Growth and Metastasis. <i>Cancer Research</i> , 2007, 67, 8043-8050.	0.9	205
5	Differences in gene expression in prostate cancer, normal appearing prostate tissue adjacent to cancer and prostate tissue from cancer free organ donors. <i>BMC Cancer</i> , 2005, 5, 45.	2.6	126
6	Myopodin, a Synaptopodin Homologue, Is Frequently Deleted in Invasive Prostate Cancers. <i>American Journal of Pathology</i> , 2001, 159, 1603-1612.	3.8	59
7	CSR1 Suppresses Tumor Growth and Metastasis of Prostate Cancer. <i>American Journal of Pathology</i> , 2006, 168, 597-607.	3.8	50
8	Identification of recurrent fusion genes across multiple cancer types. <i>Scientific Reports</i> , 2019, 9, 1074.	3.3	46
9	Whole-Genome Methylation Sequencing Reveals Distinct Impact of Differential Methylations on Gene Transcription in Prostate Cancer. <i>American Journal of Pathology</i> , 2013, 183, 1960-1970.	3.8	44
10	High throughput screening of methylation status of genes in prostate cancer using an oligonucleotide methylation array. <i>Carcinogenesis</i> , 2004, 26, 471-479.	2.8	43
11	Genome-Wide Methylation Analysis of Prostate Tissues Reveals Global Methylation Patterns of Prostate Cancer. <i>American Journal of Pathology</i> , 2013, 182, 2028-2036.	3.8	40
12	Genome Abnormalities Precede Prostate Cancer and Predict Clinical Relapse. <i>American Journal of Pathology</i> , 2012, 180, 2240-2248.	3.8	33
13	Investigating Multi-cancer Biomarkers and Their Cross-predictability in the Expression Profiles of Multiple Cancer Types. <i>Biomarker Insights</i> , 2009, 4, BML.S930.	2.5	16
14	Detection of fusion transcripts in the serum samples of patients with hepatocellular carcinoma. <i>Oncotarget</i> , 2019, 10, 3352-3360.	1.8	10
15	Transcriptome and Exome Analyses of Hepatocellular Carcinoma Reveal Patterns to Predict Cancer Recurrence in Liver Transplant Patients. <i>Hepatology Communications</i> , 2022, 6, 710-727.	4.3	9