George Michalopoulos

List of Publications by Year
in descending order

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