

# Pedro Nicolau-Neto

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

16  
papers

103  
citations

6  
h-index

9  
g-index

18  
ext. papers

137  
ext. citations

5  
avg, IF

2.1  
L-index

#	Paper	IF	Citations
16	GLIPR1 and SPARC expression profile reveals a signature associated with prostate Cancer Brain metastasis. <i>Molecular and Cellular Endocrinology</i> , <b>2021</b> , 528, 111230	4.4	1
15	MET overexpression and intratumor heterogeneity in esophageal squamous cell carcinoma. <i>Brazilian Journal of Medical and Biological Research</i> , <b>2021</b> , 54, e10877	2.8	2
14	Lipid droplet biogenesis and COX-2 pathway activation are triggered by Barrett's esophagus and adenocarcinoma, but not esophageal squamous cell carcinoma risk factors. <i>Scientific Reports</i> , <b>2021</b> , 11, 981	4.9	1
13	5-Aza-2'-deoxycytidine induces a greater inflammatory change, at the molecular levels, in normoxic than hypoxic tumor microenvironment. <i>Molecular Biology Reports</i> , <b>2021</b> , 48, 1161-1169	2.8	1
12	IL6 and BCL3 Expression Are Potential Biomarkers in Esophageal Squamous Cell Carcinoma. <i>Frontiers in Oncology</i> , <b>2021</b> , 11, 722417	5.3	1
11	Transcriptome Analysis Identifies ALCAM Overexpression as a Prognosis Biomarker in Laryngeal Squamous Cell Carcinoma. <i>Cancers</i> , <b>2020</b> , 12,	6.6	6
10	Transcriptome analysis of breast cancer cell line exposed to hypoxia-mimetic chemical CoCl <sub>2</sub> or hypoxic microenvironment. <i>Gene Reports</i> , <b>2020</b> , 20, 100686	1.4	
9	Multi-cancer V-ATPase molecular signatures: A distinctive balance of subunit C isoforms in esophageal carcinoma. <i>EBioMedicine</i> , <b>2020</b> , 51, 102581	8.8	9
8	The <i>CDKN1A</i> Gene Acts as Tumor Suppressor in Breast Cancer. <i>Cancers</i> , <b>2019</b> , 11,	6.6	7
7	UBE2C Is a Transcriptional Target of the Cell Cycle Regulator FOXM1. <i>Genes</i> , <b>2018</b> , 9,	4.2	28
6	Esophageal squamous cell carcinoma transcriptome reveals the effect of on patient outcome through novel PIK3R3 mediated activation of PI3K signaling pathway. <i>Oncotarget</i> , <b>2018</b> , 9, 16634-16647 <sup>3-3</sup>		16
5	Mutations, Differential Gene Expression, and Chimeric Transcripts in Esophageal Squamous Cell Carcinoma Show High Heterogeneity. <i>Translational Oncology</i> , <b>2018</b> , 11, 1283-1291	4.9	5
4	Prostate cancer molecular profiling: the Achilles heel for the implementation of precision medicine. <i>Cell Biology International</i> , <b>2017</b> , 41, 1239-1245	4.5	5
3	hypermethylation and decreased expression in esophageal squamous cell carcinoma and histologically normal tumor surrounding esophageal cells. <i>Clinical Epigenetics</i> , <b>2017</b> , 9, 130	7.7	10
2	Association between long interspersed nuclear element-1 methylation levels and relapse in Wilms tumors. <i>Clinical Epigenetics</i> , <b>2017</b> , 9, 128	7.7	7
1	Nicotinic cholinergic receptors in esophagus: Early alteration during carcinogenesis and prognostic value. <i>World Journal of Gastroenterology</i> , <b>2016</b> , 22, 7146-56	5.6	3