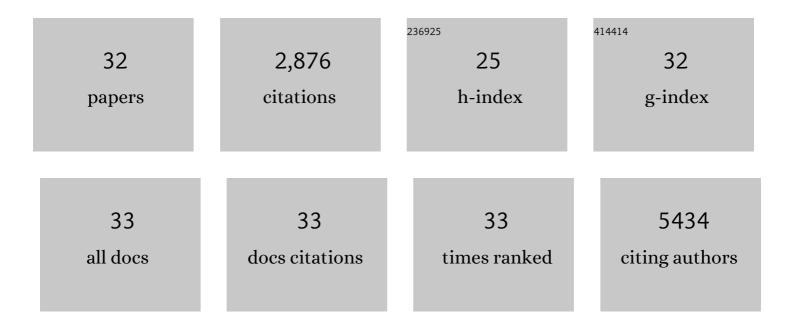
## Narciso Olvera

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12050337/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	lschemia in Tumors Induces Early and Sustained Phosphorylation Changes in Stress Kinase Pathways but Does Not Affect Global Protein Levels. Molecular and Cellular Proteomics, 2014, 13, 1690-1704.	3.8	323
2	Recurrent SMARCA4 mutations in small cell carcinoma of the ovary. Nature Genetics, 2014, 46, 424-426.	21.4	291
3	Identification of Molecular Pathway Aberrations in Uterine Serous Carcinoma by Genome-wide Analyses. Journal of the National Cancer Institute, 2012, 104, 1503-1513.	6.3	231
4	Morphologic patterns associated with BRCA1 and BRCA2 genotype in ovarian carcinoma. Modern Pathology, 2012, 25, 625-636.	5.5	202
5	Uterine Cancer After Risk-Reducing Salpingo-oophorectomy Without Hysterectomy in Women With <i>BRCA</i> Mutations. JAMA Oncology, 2016, 2, 1434.	7.1	189
6	Molecular analysis of high-grade serous ovarian carcinoma with and without associated serous tubal intra-epithelial carcinoma. Nature Communications, 2017, 8, 990.	12.8	169
7	Clinicopathologic Significance of Defective DNA Mismatch Repair in Endometrial Carcinoma. Journal of Clinical Oncology, 2006, 24, 1745-1753.	1.6	152
8	Immune-Active Microenvironment in Small Cell Carcinoma of the Ovary, Hypercalcemic Type: Rationale for Immune Checkpoint Blockade. Journal of the National Cancer Institute, 2018, 110, 787-790.	6.3	123
9	Noninvasive ovarian cancer biomarker detection via an optical nanosensor implant. Science Advances, 2018, 4, eaaq1090.	10.3	121
10	Genomic Complexity and AKT Dependence in Serous Ovarian Cancer. Cancer Discovery, 2012, 2, 56-67.	9.4	109
11	Genetic Analysis of the Early Natural History of Epithelial Ovarian Carcinoma. PLoS ONE, 2010, 5, e10358.	2.5	90
12	Loss of SMARCA4 Expression Is Both Sensitive and Specific for the Diagnosis of Small Cell Carcinoma of Ovary, Hypercalcemic Type. American Journal of Surgical Pathology, 2016, 40, 395-403.	3.7	87
13	Heterogenic Loss of the Wild-Type BRCA Allele in Human Breast Tumorigenesis. Annals of Surgical Oncology, 2007, 14, 2510-2518.	1.5	82
14	Clinicopathologic Analysis of Early-stage Sporadic Ovarian Carcinoma. American Journal of Surgical Pathology, 2004, 28, 147-159.	3.7	77
15	Mutation and expression of the TP53 gene in early stage epithelial ovarian carcinoma. Gynecologic Oncology, 2004, 93, 301-306.	1.4	76
16	Concomitant loss of SMARCA2 and SMARCA4 expression in small cell carcinoma of the ovary, hypercalcemic type. Modern Pathology, 2016, 29, 60-66.	5.5	62
17	BRCA1 Immunohistochemistry in a Molecularly Characterized Cohort of Ovarian High-Grade Serous Carcinomas. American Journal of Surgical Pathology, 2013, 37, 138-146.	3.7	54
18	Increased Progesterone Receptor Expression in Benign Epithelium of BRCA1-Related Breast Cancers. Cancer Research, 2004, 64, 5051-5053.	0.9	51

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#	Article	IF	CITATIONS
19	Genetically Defined, Syngeneic Organoid Platform for Developing Combination Therapies for Ovarian Cancer. Cancer Discovery, 2021, 11, 362-383.	9.4	50
20	Protocol for PTEN Expression by Immunohistochemistry in Formalin-fixed Paraffin-embedded Human Breast Carcinoma. Applied Immunohistochemistry and Molecular Morphology, 2010, 18, 371-374.	1.2	50
21	Massively parallel sequencing analysis of mucinous ovarian carcinomas: genomic profiling and differential diagnoses. Gynecologic Oncology, 2018, 150, 127-135.	1.4	41
22	Clonal relatedness between lobular carcinoma in situ and synchronous malignant lesions. Breast Cancer Research, 2012, 14, R103.	5.0	38
23	Molecular Subtypes of Uterine Leiomyosarcoma and Correlation with Clinical Outcome. Neoplasia, 2015, 17, 183-189.	5.3	33
24	Gene expression profiling of lobular carcinoma in situ reveals candidate precursor genes for invasion. Molecular Oncology, 2015, 9, 772-782.	4.6	32
25	Antibodies Against Specific MUC16 Glycosylation Sites Inhibit Ovarian Cancer Growth. ACS Chemical Biology, 2017, 12, 2085-2096.	3.4	32
26	Tissue preparation for laser capture microdissection and RNA extraction from fresh frozen breast tissue. BioTechniques, 2007, 43, 41-48.	1.8	26
27	Clinicopathologic Analysis of Matched Primary and Recurrent Endometrial Carcinoma. American Journal of Surgical Pathology, 2012, 36, 1771-1781.	3.7	22
28	miR-200c-driven Mesenchymal-To-Epithelial Transition is a Therapeutic Target in Uterine Carcinosarcomas. Scientific Reports, 2017, 7, 3614.	3.3	22
29	Blocking and Randomization to Improve Molecular Biomarker Discovery. Clinical Cancer Research, 2014, 20, 3371-3378.	7.0	18
30	Genetic predisposition to bevacizumab-induced hypertension. Gynecologic Oncology, 2017, 147, 621-625.	1.4	17
31	A pair of datasets for microRNA expression profiling to examine the use of careful study design for assigning arrays to samples. Scientific Data, 2018, 5, 180084.	5.3	5
32	Frequency of <i>BRCA1</i> and <i>BRCA2</i> germline mutations in uterine serous carcinomas and uterine carcinosarcomas. Journal of Clinical Oncology, 2018, 36, e13521-e13521.	1.6	1