

# Sandra L Romero-Cordoba

## List of Publications by Year in descending order

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Version: 2024-02-01

40  
papers

3,208  
citations

394421

19  
h-index

377865

34  
g-index

42  
all docs

42  
docs citations

42  
times ranked

7247  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comprehensive omic characterization of breast cancer in Mexican-Hispanic women. <i>Nature Communications</i> , 2021, 12, 2245.	12.8	14
2	Regulation of anti-tumorigenic pathways by the combinatory treatment of calcitriol and TGF- $\beta$ 2 in PC-3 and DU145 cells. <i>Journal of Steroid Biochemistry and Molecular Biology</i> , 2021, 209, 105831.	2.5	4
3	Mexican <i>Ganoderma Lucidum</i> Extracts Decrease Lipogenesis Modulating Transcriptional Metabolic Networks and Gut Microbiota in C57BL/6 Mice Fed with a High-Cholesterol Diet. <i>Nutrients</i> , 2021, 13, 38.	4.1	15
4	miR-9-Mediated Inhibition of EFEMP1 Contributes to the Acquisition of Pro-Tumoral Properties in Normal Fibroblasts. <i>Cells</i> , 2020, 9, 2143.	4.1	13
5	Biological Landscape of Triple Negative Breast Cancers Expressing CTLA-4. <i>Frontiers in Oncology</i> , 2020, 10, 1206.	2.8	21
6	A lncRNA landscape in breast cancer reveals a potential role for AC009283.1 in proliferation and apoptosis in HER2-enriched subtype. <i>Scientific Reports</i> , 2020, 10, 13146.	3.3	24
7	MiR-302b as a Combinatorial Therapeutic Approach to Improve Cisplatin Chemotherapy Efficacy in Human Triple-Negative Breast Cancer. <i>Cancers</i> , 2020, 12, 2261.	3.7	12
8	Deep Into Breast Cancer Heterogeneity to Increase Immunotherapeutic Effectiveness. <i>JCO Precision Oncology</i> , 2020, 4, 1267-1268.	3.0	2
9	Decoding Immune Heterogeneity of Triple Negative Breast Cancer and Its Association with Systemic Inflammation. <i>Cancers</i> , 2019, 11, 911.	3.7	40
10	WNT signaling modulates PD-L1 expression in the stem cell compartment of triple-negative breast cancer. <i>Oncogene</i> , 2019, 38, 4047-4060.	5.9	137
11	Wide Profiling of Circulating MicroRNAs in Spinocerebellar Ataxia Type 7. <i>Molecular Neurobiology</i> , 2019, 56, 6106-6120.	4.0	12
12	MicroRNAs in Tumor Cell Metabolism: Roles and Therapeutic Opportunities. <i>Frontiers in Oncology</i> , 2019, 9, 1404.	2.8	53
13	Atypical <i>RAS</i> Mutations in Metastatic Colorectal Cancer. <i>JCO Precision Oncology</i> , 2019, 3, 1-11.	3.0	1
14	Expression of long non-coding RNA ENSG00000226738 (LncKLHDC7B) is enriched in the immunomodulatory triple-negative breast cancer subtype and its alteration promotes cell migration, invasion, and resistance to cell death. <i>Molecular Oncology</i> , 2019, 13, 909-927.	4.6	29
15	Population and breast cancer patients' analysis reveals the diversity of genomic variation of the BRCA genes in the Mexican population. <i>Human Genomics</i> , 2019, 13, 3.	2.9	16
16	Loss of function of miR-342-3p results in MCT1 over-expression and contributes to oncogenic metabolic reprogramming in triple negative breast cancer. <i>Scientific Reports</i> , 2018, 8, 12252.	3.3	75
17	The PDZ-Binding Motif of HPV16-E6 Oncoprotein Modulates the Keratinization and Stemness Transcriptional Profile <i>In Vivo</i> . <i>BioMed Research International</i> , 2017, 2017, 1-9.	1.9	3
18	Molecular features of influenza A (H1N1)pdm09 prevalent in Mexico during winter seasons 2012-2014. <i>PLoS ONE</i> , 2017, 12, e0180419.	2.5	7

#	ARTICLE	IF	CITATIONS
19	Abstract 5437: miR-302b as adjuvant therapeutic tool to improve chemotherapy efficacy in human triple-negative breast cancer. , 2017, , .		0
20	Insights into the Regulatory Role of Non-coding RNAs in Cancer Metabolism. <i>Frontiers in Physiology</i> , 2016, 7, 342.	2.8	38
21	Exosome-mediated delivery of miR-9 induces cancer-associated fibroblast-like properties in human breast fibroblasts. <i>Cell Death and Disease</i> , 2016, 7, e2312-e2312.	6.3	232
22	Expression and prognostic significance of the autoimmune regulator gene in breast cancer cells. <i>Cell Cycle</i> , 2016, 15, 3220-3229.	2.6	16
23	Changes in global gene expression profiles induced by HPV 16 E6 oncoprotein variants in cervical carcinoma C33-A cells. <i>Virology</i> , 2016, 488, 187-195.	2.4	29
24	Abstract A47: A microRNA signature identifies subtypes of triple-negative breast cancer and reveals miR-342-3p as regulator of a lactate metabolic pathway through silencing monocarboxylate transporter 1. <i>Cancer Research</i> , 2016, 76, A47-A47.	0.9	2
25	Abstract 3924: AIRE is expressed and associated with good prognosis in breast cancer. , 2016, , .		0
26	Calcitriol increases Dicer expression and modifies the microRNAs signature in SiHa cervical cancer cells. <i>Biochemistry and Cell Biology</i> , 2015, 93, 376-384.	2.0	24
27	Altered Expression of Circulating MicroRNA in Plasma of Patients with Primary Osteoarthritis and In Silico Analysis of Their Pathways. <i>PLoS ONE</i> , 2014, 9, e97690.	2.5	85
28	MicroRNAs transported by exosomes in body fluids as mediators of intercellular communication in cancer. <i>OncoTargets and Therapy</i> , 2014, 7, 1327.	2.0	125
29	miRNA biogenesis: Biological impact in the development of cancer. <i>Cancer Biology and Therapy</i> , 2014, 15, 1444-1455.	3.4	205
30	mRNA and miRNA expression patterns associated to pathways linked to metal mixture health effects. <i>Gene</i> , 2014, 533, 508-514.	2.2	54
31	Landscape of genomic alterations in cervical carcinomas. <i>Nature</i> , 2014, 506, 371-375.	27.8	708
32	Abstract 4370: miRNA profiles identify different subgroups of triple negative tumors and reveal novel miRNA-mRNA interactions in breast cancer tumorigenesis. , 2014, , .		0
33	Seasonal and pandemic influenza H1N1 viruses induce differential expression of SOCS-1 and RIG-I genes and cytokine/chemokine production in macrophages. <i>Cytokine</i> , 2013, 62, 151-159.	3.2	34
34	Gene expression profile regulated by the HPV16 E7 oncoprotein and estradiol in cervical tissue. <i>Virology</i> , 2013, 447, 155-165.	2.4	30
35	Williams neural stem cells new model for insight into microRNA dysregulation. <i>Frontiers in Bioscience - Elite</i> , 2013, E5, 1057-1073.	1.8	2
36	Abstract 4604: Landscape of human and viral genomic alterations in cervical carcinomas.. , 2013, , .		0

#	ARTICLE	IF	CITATIONS
37	Sequence analysis of mutations and translocations across breast cancer subtypes. Nature, 2012, 486, 405-409.	27.8	1,107
38	Identification and Pathway Analysis of microRNAs with No Previous Involvement in Breast Cancer. PLoS ONE, 2012, 7, e31904.	2.5	39
39	Abstract A25: Differences in microRNA expression patterns in breast cancer subtypes. Cancer Research, 2012, 72, A25-A25.	0.9	0
40	Abstract PL07-01: Molecular profiling of breast cancer in Mexico: Identification of novel therapeutic targets through whole genome sequencing analysis.. , 2012, , .		0