Michael L Gatza

List of Publications by Citations

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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.

40
papers

3,718
citations

43
papers

4,649
ext. papers

24
h-index

9.9
avg, IF

4.55
L-index

#	Paper	IF	Citations
40	Comprehensive Molecular Portraits of Invasive Lobular Breast Cancer. <i>Cell</i> , 2015 , 163, 506-19	56.2	1055
39	Proteogenomics connects somatic mutations to signalling in breast cancer. <i>Nature</i> , 2016 , 534, 55-62	50.4	938
38	A pathway-based classification of human breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 6994-9	11.5	258
37	An integrated genomics approach identifies drivers of proliferation in luminal-subtype human breast cancer. <i>Nature Genetics</i> , 2014 , 46, 1051-9	36.3	158
36	Tumor mutational burden is a determinant of immune-mediated survival in breast cancer. Oncolmmunology, 2018, 7, e1490854	7.2	129
35	Parkin targets HIF-11 for ubiquitination and degradation to inhibit breast tumor progression. <i>Nature Communications</i> , 2017 , 8, 1823	17.4	100
34	Exosomal miRNA confers chemo resistance via targeting Cav1/p-gp/M2-type macrophage axis in ovarian cancer. <i>EBioMedicine</i> , 2018 , 38, 100-112	8.8	100
33	NRF2 Induction Supporting Breast Cancer Cell Survival Is Enabled by Oxidative Stress-Induced DPP3-KEAP1 Interaction. <i>Cancer Research</i> , 2017 , 77, 2881-2892	10.1	91
32	Using a stem cell-based signature to guide therapeutic selection in cancer. <i>Cancer Research</i> , 2011 , 71, 1772-80	10.1	91
31	A genomic strategy to elucidate modules of oncogenic pathway signaling networks. <i>Molecular Cell</i> , 2009 , 34, 104-14	17.6	91
30	Cellular transformation by the HTLV-I Tax protein, a jack-of-all-trades. <i>Oncogene</i> , 2003 , 22, 5141-9	9.2	80
29	Genetic heterogeneity of Myc-induced mammary tumors reflecting diverse phenotypes including metastatic potential. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 16387-92	11.5	67
28	Analysis of tumor environmental response and oncogenic pathway activation identifies distinct basal and luminal features in HER2-related breast tumor subtypes. <i>Breast Cancer Research</i> , 2011 , 13, R62	8.3	50
27	Type III TGF-Ireceptor enhances colon cancer cell migration and anchorage-independent growth. <i>Neoplasia</i> , 2011 , 13, 758-70	6.4	45
26	New Mechanisms of Resistance to MEK Inhibitors in Melanoma Revealed by Intravital Imaging. <i>Cancer Research</i> , 2018 , 78, 542-557	10.1	43
25	Cross-species DNA copy number analyses identifies multiple 1q21-q23 subtype-specific driver genes for breast cancer. <i>Breast Cancer Research and Treatment</i> , 2015 , 152, 347-56	4.4	37
24	Amplification of SOX4 promotes PI3K/Akt signaling in human breast cancer. <i>Breast Cancer Research and Treatment</i> , 2017 , 162, 439-450	4.4	34

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23	FOXO transcription factors control E2F1 transcriptional specificity and apoptotic function. <i>Cancer Research</i> , 2013 , 73, 6056-67	10.1	33
22	Activating PIK3CA Mutations Induce an Epidermal Growth Factor Receptor (EGFR)/Extracellular Signal-regulated Kinase (ERK) Paracrine Signaling Axis in Basal-like Breast Cancer. <i>Molecular and Cellular Proteomics</i> , 2015 , 14, 1959-76	7.6	31
21	Ubiquitination of HTLV-I Tax in response to DNA damage regulates nuclear complex formation and nuclear export. <i>Retrovirology</i> , 2007 , 4, 95	3.6	31
20	Mediator kinase CDK8/CDK19 drives YAP1-dependent BMP4-induced EMT in cancer. <i>Oncogene</i> , 2018 , 37, 4792-4808	9.2	31
19	Impact of transforming viruses on cellular mutagenesis, genome stability, and cellular transformation. <i>Environmental and Molecular Mutagenesis</i> , 2005 , 45, 304-25	3.2	30
18	SIGNATURE: a workbench for gene expression signature analysis. <i>BMC Bioinformatics</i> , 2011 , 12, 443	3.6	25
17	SMAD4 Suppresses WNT-Driven Dedifferentiation and Oncogenesis in the Differentiated Gut Epithelium. <i>Cancer Research</i> , 2018 , 78, 4878-4890	10.1	24
16	Interaction of E2F7 transcription factor with E2F1 and C-terminal-binding protein (CtBP) provides a mechanism for E2F7-dependent transcription repression. <i>Journal of Biological Chemistry</i> , 2013 , 288, 24.	5 8 149	24
15	Multi-Omic Data Interpretation to Repurpose Subtype Specific Drug Candidates for Breast Cancer. <i>Frontiers in Genetics</i> , 2019 , 10, 420	4.5	21
14	Genotoxic stress and cellular stress alter the subcellular distribution of human T-cell leukemia virus type 1 tax through a CRM1-dependent mechanism. <i>Journal of Virology</i> , 2006 , 80, 6657-68	6.6	21
13	The Modulatory Role of MicroRNA-873 in the Progression of KRAS-Driven Cancers. <i>Molecular Therapy - Nucleic Acids</i> , 2019 , 14, 301-317	10.7	17
12	Recurrence analysis on prostate cancer patients with Gleason score 7 using integrated histopathology whole-slide images and genomic data through deep neural networks. <i>Journal of Medical Imaging</i> , 2018 , 5, 047501	2.6	14
11	Integrative proteogenomic analyses of human tumours identifies ADNP as a novel oncogenic mediator of cell cycle progression in high-grade serous ovarian cancer with poor prognosis. <i>EBioMedicine</i> , 2019 , 50, 191-202	8.8	9
10	Emerging Role of SOX Proteins in Breast Cancer Development and Maintenance. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2019 , 24, 213-230	2.4	8
9	Aggressive Mammary Cancers Lacking Lymphocytic Infiltration Arise in Irradiated Mice and Can Be Prevented by Dietary Intervention. <i>Cancer Immunology Research</i> , 2020 , 8, 217-229	12.5	6
8	CPT1A and fatty acid Ebxidation are essential for tumor cell growth and survival in hormone receptor-positive breast cancer. <i>NAR Cancer</i> , 2021 , 3, zcab035	5.2	5
7	Recent Advances in Integrative Multi-Omics Research in Breast and Ovarian Cancer. <i>Journal of Personalized Medicine</i> , 2021 , 11,	3.6	5
6	FOXA1 and adaptive response determinants to HER2 targeted therapy in TBCRC 036. <i>Npj Breast Cancer</i> , 2021 , 7, 51	7.8	4

5	Differentiation among prostate cancer patients with Gleason score of 7 using histopathology whole-slide image and genomic data. <i>Proceedings of SPIE</i> , 2018 , 10579,	1.7	3
4	SOX4 and SMARCA4 cooperatively regulate PI3k signaling through transcriptional activation of TGFBR2. <i>Npj Breast Cancer</i> , 2021 , 7, 40	7.8	3
3	Multi-omic Dissection of Oncogenically Active Epiproteomes Identifies Drivers of Proliferative and Invasive Breast Tumors. <i>IScience</i> , 2019 , 17, 359-378	6.1	2
2	Nuclear Receptor-Mediated Metabolic Reprogramming and the Impact on HR+ Breast Cancer. <i>Cancers</i> , 2021 , 13,	6.6	1
1	SMAD4 is critical in suppression of BRAF-V600E serrated tumorigenesis. <i>Oncogene</i> , 2021 , 40, 6034-604	18 9.2	О