Tamar Geiger

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.

6,814 80 72 32 h-index g-index citations papers 80 9,621 14.2 5.95 avg, IF L-index ext. citations ext. papers

#	Paper Paper	IF	Citations
7 2	Disrupted neural correlates of anesthesia and sleep reveal early circuit dysfunctions in Alzheimer models <i>Cell Reports</i> , 2022 , 38, 110268	10.6	1
71	Nucleoporin-93 reveals a common feature of aggressive breast cancers: robust nucleocytoplasmic transport of transcription factors <i>Cell Reports</i> , 2022 , 38, 110418	10.6	0
70	Phosphoproteomics reveals novel modes of function and inter-relationships among PIKKs in response to genotoxic stress. <i>EMBO Journal</i> , 2021 , 40, e104400	13	14
69	IRS1 phosphorylation underlies the non-stochastic probability of cancer cells to persist during EGFR inhibition therapy <i>Nature Cancer</i> , 2021 , 2, 1055-1070	15.4	0
68	Proteogenomics of glioblastoma associates molecular patterns with survival. <i>Cell Reports</i> , 2021 , 34, 108	8 787 6	6
67	Distinct extracellular-matrix remodeling events precede symptoms of inflammation. <i>Matrix Biology</i> , 2021 , 96, 47-68	11.4	6
66	Anti-tumour immunity induces aberrant peptide presentation in melanoma. <i>Nature</i> , 2021 , 590, 332-337	50.4	28
65	Serine Biosynthesis Is a Metabolic Vulnerability in IDH2-Driven Breast Cancer Progression. <i>Cancer Research</i> , 2021 , 81, 1443-1456	10.1	4
64	Clinical Proteomics of Metastatic Melanoma Reveals Profiles of Organ Specificity and Treatment Resistance. <i>Clinical Cancer Research</i> , 2021 , 27, 2074-2086	12.9	3
63	Metastasis-Entrained Eosinophils Enhance Lymphocyte-Mediated Antitumor Immunity. <i>Cancer Research</i> , 2021 , 81, 5555-5571	10.1	3
62	Nascent Ribo-Seq measures ribosomal loading time and reveals kinetic impact on ribosome density. <i>Nature Methods</i> , 2021 , 18, 1068-1074	21.6	2
61	Spontaneous regression of micro-metastases following primary tumor excision: a critical role for primary tumor secretome. <i>BMC Biology</i> , 2020 , 18, 163	7.3	1
60	NF- B -miR-155 axis activation mediates ovulation-induced oncogenic effects in fallopian tube epithelium. <i>Carcinogenesis</i> , 2020 , 41, 1703-1712	4.6	2
59	Across the Globe: Proteogenomic Landscapes of Lung Cancer. Cell, 2020, 182, 9-11	56.2	
58	Proteomic patterns associated with response to breast cancer neoadjuvant treatment. <i>Molecular Systems Biology</i> , 2020 , 16, e9443	12.2	15
57	PDZD8 interacts with Protrudin and Rab7 at ER-late endosome membrane contact sites associated with mitochondria. <i>Nature Communications</i> , 2020 , 11, 3645	17.4	26
56	Spatiotemporal Proteomic Analysis of Stress Granule Disassembly Using APEX Reveals Regulation by SUMOylation and Links to ALS Pathogenesis. <i>Molecular Cell</i> , 2020 , 80, 876-891.e6	17.6	44

(2018-2020)

55	Pleiotropic tumor suppressor functions of WWOX antagonize metastasis. <i>Signal Transduction and Targeted Therapy</i> , 2020 , 5, 43	21	11
54	Proteomics of Melanoma Response to Immunotherapy Reveals Mitochondrial Dependence. <i>Cell</i> , 2019 , 179, 236-250.e18	56.2	107
53	Simultaneous Integration of Multi-omics Data Improves the Identification of Cancer Driver Modules. <i>Cell Systems</i> , 2019 , 8, 456-466.e5	10.6	21
52	Mitochondrial Regulation of the Hippocampal Firing Rate Set Point and Seizure Susceptibility. <i>Neuron</i> , 2019 , 102, 1009-1024.e8	13.9	51
51	Specific inhibition of splicing factor activity by decoy RNA oligonucleotides. <i>Nature Communications</i> , 2019 , 10, 1590	17.4	37
50	Proteomic and genomic signatures of repeat instability in cancer and adjacent normal tissues. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 16987-16996	5 11.5	6
49	Systematic Detection of Amino Acid Substitutions in Proteomes Reveals Mechanistic Basis of Ribosome Errors and Selection for Translation Fidelity. <i>Molecular Cell</i> , 2019 , 75, 427-441.e5	17.6	36
48	MicroRNAs Affect Complement Regulator Expression and Mitochondrial Activity to Modulate Cell Resistance to Complement-Dependent Cytotoxicity. <i>Cancer Immunology Research</i> , 2019 , 7, 1970-1983	12.5	6
47	Microvesicle Proteomic Profiling of Uterine Liquid Biopsy for Ovarian Cancer Early Detection. <i>Molecular and Cellular Proteomics</i> , 2019 , 18, 865-875	7.6	25
46	The landscape of tiered regulation of breast cancer cell metabolism. <i>Scientific Reports</i> , 2019 , 9, 17760	4.9	9
45	Cell shape alteration during adipogenesis is associated with coordinated matrix cues. <i>Journal of Cellular Physiology</i> , 2019 , 234, 3850-3863	7	23
44	UBQLN4 Represses Homologous Recombination and Is Overexpressed in Aggressive Tumors. <i>Cell</i> , 2019 , 176, 505-519.e22	56.2	68
43	Identification of nucleolar protein NOM1 as a novel nuclear IGF1R-interacting protein. <i>Molecular Genetics and Metabolism</i> , 2019 , 126, 259-265	3.7	4
42	S101, an Inhibitor of Proliferating T Cells, Rescues Mice From Superantigen-Induced Shock. <i>Journal of Infectious Diseases</i> , 2018 , 217, 288-297	7	1
41	IGF1R signaling drives antiestrogen resistance through PAK2/PIX activation in luminal breast cancer. <i>Oncogene</i> , 2018 , 37, 1869-1884	9.2	19
40	Nuclear poly(A)-binding protein 1 is an ATM target and essential for DNA double-strand break repair. <i>Nucleic Acids Research</i> , 2018 , 46, 730-747	20.1	6
39	The Proteome of Prostate Cancer Bone Metastasis Reveals Heterogeneity with Prognostic Implications. <i>Clinical Cancer Research</i> , 2018 , 24, 5433-5444	12.9	44
38	Clinical Proteomics of Breast Cancer Reveals a Novel Layer of Breast Cancer Classification. <i>Cancer Research</i> , 2018 , 78, 6001-6010	10.1	38

37	Proteomic analysis of polyribosomes identifies splicing factors as potential regulators of translation during mitosis. <i>Nucleic Acids Research</i> , 2017 , 45, 5945-5957	20.1	23
36	Mutant eIF2B leads to impaired mitochondrial oxidative phosphorylation in vanishing white matter disease. <i>Journal of Neurochemistry</i> , 2017 , 141, 694-707	6	26
35	Next-Generation Proteomics and Its Application to Clinical Breast Cancer Research. <i>American Journal of Pathology</i> , 2017 , 187, 2175-2184	5.8	21
34	Plasma Biomarker Identification and Quantification by Microparticle Proteomics. <i>Methods in Molecular Biology</i> , 2017 , 1619, 477-486	1.4	2
33	Reduced changes in protein compared to mRNA levels across non-proliferating tissues. <i>BMC Genomics</i> , 2017 , 18, 305	4.5	43
32	Breast cancer tumorigenicity is dependent on high expression levels of NAF-1 and the lability of its Fe-S clusters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 10890-5	11.5	49
31	The Perseus computational platform for comprehensive analysis of (prote)omics data. <i>Nature Methods</i> , 2016 , 13, 731-40	21.6	3257
30	The Proteome of Primary Prostate Cancer. <i>European Urology</i> , 2016 , 69, 942-52	10.2	97
29	System-wide Clinical Proteomics of Breast Cancer Reveals Global Remodeling of Tissue Homeostasis. <i>Cell Systems</i> , 2016 , 2, 172-84	10.6	63
28	Proteomic maps of breast cancer subtypes. <i>Nature Communications</i> , 2016 , 7, 10259	17.4	178
27	TMPRSS2-ERG fusion protein regulates insulin-like growth factor-1 receptor (IGF1R) gene expression in prostate cancer: involvement of transcription factor Sp1. <i>Oncotarget</i> , 2016 , 7, 51375-513	9 3 ·3	17
26	Elucidation of Signaling Pathways from Large-Scale Phosphoproteomic Data Using Protein Interaction Networks. <i>Cell Systems</i> , 2016 , 3, 585-593.e3	10.6	42
25	Tumor macrophages are pivotal constructors of tumor collagenous matrix. <i>Journal of Experimental Medicine</i> , 2016 , 213, 2315-2331	16.6	174
24	The focal adhesion protein PINCH-1 associates with EPLIN at integrin adhesion sites. <i>Journal of Cell Science</i> , 2015 , 128, 1023-33	5.3	17
23	Myelin-associated glycoprotein gene mutation causes Pelizaeus-Merzbacher disease-like disorder. <i>Brain</i> , 2015 , 138, 2521-36	11.2	37
22	Lam6 Regulates the Extent of Contacts between Organelles. <i>Cell Reports</i> , 2015 , 12, 7-14	10.6	140
21	Senescent cells communicate via intercellular protein transfer. <i>Genes and Development</i> , 2015 , 29, 791-8	8 02 2.6	82
20	Regulation of Elg1 activity by phosphorylation. <i>Cell Cycle</i> , 2015 , 14, 3689-97	4.7	10

19	Proteomics-level analysis of myelin formation and regeneration in a mouse model for Vanishing White Matter disease. <i>Journal of Neurochemistry</i> , 2015 , 134, 513-26	6	21
18	Down-regulation of LATS kinases alters p53 to promote cell migration. <i>Genes and Development</i> , 2015 , 29, 2325-30	12.6	49
17	Phosphoproteomic analysis reveals Smarcb1 dependent EGFR signaling in Malignant Rhabdoid tumor cells. <i>Molecular Cancer</i> , 2015 , 14, 167	42.1	11
16	MiR-375 promotes redifferentiation of adult human Itells expanded in vitro. <i>PLoS ONE</i> , 2015 , 10, e0122	1908	48
15	Uncovering Hidden Layers of Cell Cycle Regulation through Integrative Multi-omic Analysis. <i>PLoS Genetics</i> , 2015 , 11, e1005554	6	34
14	Proteomics of microparticles with SILAC Quantification (PROMIS-Quan): a novel proteomic method for plasma biomarker quantification. <i>Molecular and Cellular Proteomics</i> , 2015 , 14, 1127-36	7.6	32
13	Tumor-derived osteopontin reprograms normal mammary fibroblasts to promote inflammation and tumor growth in breast cancer. <i>Cancer Research</i> , 2015 , 75, 963-73	10.1	98
12	Rescue of embryonic stem cells from cellular transformation by proteomic stabilization of mutant p53 and conversion into WT conformation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 7006-11	11.5	36
11	Predicting cancer-specific vulnerability via data-driven detection of synthetic lethality. <i>Cell</i> , 2014 , 158, 1199-1209	56.2	167
10	A dynamic interface between vacuoles and mitochondria in yeast. <i>Developmental Cell</i> , 2014 , 30, 95-102	10.2	266
9	Genome-wide identification and quantification of protein synthesis in cultured cells and whole tissues by puromycin-associated nascent chain proteomics (PUNCH-P). <i>Nature Protocols</i> , 2014 , 9, 751-60	18.8	52
8	PUNCH-P for global translatome profiling: Methodology, insights and comparison to other techniques. <i>Translation</i> , 2013 , 1, e27516		11
7	Opening the floodgates: proteomics and the integrin adhesome. <i>Current Opinion in Cell Biology</i> , 2012 , 24, 562-8	9	78
6	Proteomic portrait of human breast cancer progression identifies novel prognostic markers. <i>Cancer Research</i> , 2012 , 72, 2428-39	10.1	107
5	Use of stable isotope labeling by amino acids in cell culture as a spike-in standard in quantitative proteomics. <i>Nature Protocols</i> , 2011 , 6, 147-57	18.8	232
4	Super-SILAC mix for quantitative proteomics of human tumor tissue. <i>Nature Methods</i> , 2010 , 7, 383-5	21.6	431
3	Proteomics on an Orbitrap benchtop mass spectrometer using all-ion fragmentation. <i>Molecular and Cellular Proteomics</i> , 2010 , 9, 2252-61	7.6	189
2	ModulOmics: Integrating Multi-Omics Data to Identify Cancer Driver Modules		1

Distinct extracellular-matrix remodeling events precede symptoms of inflammation

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