## Irene M Ong

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6871062/publications.pdf

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41 papers

1,687 citations

394390 19 h-index 315719 38 g-index

41 all docs

41 docs citations

41 times ranked

3230 citing authors

#	Article	IF	CITATIONS
1	A Population-Based Study of Genes Previously Implicated in Breast Cancer. New England Journal of Medicine, 2021, 384, 440-451.	27.0	414
2	PKM2 methylation by CARM1 activates aerobic glycolysis to promote tumorigenesis. Nature Cell Biology, 2017, 19, 1358-1370.	10.3	212
3	Sex―and tissueâ€specific changes in <scp>mTOR</scp> signaling with age in C57 <scp>BL</scp> /6J mice. Aging Cell, 2016, 15, 155-166.	6.7	149
4	PP2A-B′ holoenzyme substrate recognition, regulation and role in cytokinesis. Cell Discovery, 2017, 3, 17027.	6.7	68
5	Gut microbiome populations are associated with structure-specific changes in white matter architecture. Translational Psychiatry, 2018, 8, 6.	4.8	63
6	MASH Explorer: A Universal Software Environment for Top-Down Proteomics. Journal of Proteome Research, 2020, 19, 3867-3876.	3.7	62
7	GATA Factor-Dependent Positive-Feedback Circuit in Acute Myeloid Leukemia Cells. Cell Reports, 2016, 16, 2428-2441.	6.4	59
8	The landscape of antibody binding in SARS-CoV-2 infection. PLoS Biology, 2021, 19, e3001265.	5.6	58
9	Caloric Restriction Engages Hepatic RNA Processing Mechanisms in Rhesus Monkeys. Cell Metabolism, 2018, 27, 677-688.e5.	16.2	56
10	Risk of Breast Cancer Among Carriers of Pathogenic Variants in Breast Cancer Predisposition Genes Varies by Polygenic Risk Score. Journal of Clinical Oncology, 2021, 39, 2564-2573.	1.6	47
11	Integrating Enhancer Mechanisms to Establish a Hierarchical Blood Development Program. Cell Reports, 2017, 20, 2966-2979.	6.4	46
12	Sex and genetic background define the metabolic, physiologic, and molecular response to protein restriction. Cell Metabolism, 2022, 34, 209-226.e5.	16.2	44
13	Polycomb Group Protein YY1 Is an Essential Regulator of Hematopoietic Stem Cell Quiescence. Cell Reports, 2018, 22, 1545-1559.	6.4	36
14	Cotargeting mTORC and EGFR Signaling as a Therapeutic Strategy in HNSCC. Molecular Cancer Therapeutics, 2017, 16, 1257-1268.	4.1	32
15	Tissue factor-factor VIIa complex triggers protease activated receptor 2-dependent growth factor release and migration in ovarian cancer. Gynecologic Oncology, 2017, 145, 167-175.	1.4	30
16	BAF155 methylation drives metastasis by hijacking super-enhancers and subverting anti-tumor immunity. Nucleic Acids Research, 2021, 49, 12211-12233.	14.5	29
17	SOX17 integrates HOXA and arterial programs in hemogenic endothelium to drive definitive lympho-myeloid hematopoiesis. Cell Reports, 2021, 34, 108758.	6.4	27
18	Defining the boundaries and expanding the utility of head and neck cancer patient derived xenografts. Oral Oncology, 2017, 64, 65-72.	1.5	24

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19	PKM2â€TMEM33 axis regulates lipid homeostasis in cancer cells by controlling SCAP stability. EMBO Journal, 2021, 40, e108065.	7.8	24
20	Ultraviolet radiation-induced tumor necrosis factor alpha, which is linked to the development of cutaneous SCC, modulates differential epidermal microRNAs expression. Oncotarget, 2016, 7, 17945-17956.	1.8	22
21	Transcriptional and Functional Programming of Decidual Innate Lymphoid Cells. Frontiers in Immunology, 2019, 10, 3065.	4.8	21
22	Risk of Late-Onset Breast Cancer in Genetically Predisposed Women. Journal of Clinical Oncology, 2021, 39, 3430-3440.	1.6	21
23	The Metabolic Response to a Low Amino Acid Diet is Independent of Diet-Induced Shifts in the Composition of the Gut Microbiome. Scientific Reports, 2019, 9, 67.	3.3	16
24	Systemic Metabolic Alterations Correlate with Islet-Level Prostaglandin E2 Production and Signaling Mechanisms That Predict $\hat{l}^2$ -Cell Dysfunction in a Mouse Model of Type 2 Diabetes. Metabolites, 2021, 11, 58.	2.9	16
25	Determination of tissue contributions to the circulating lipid pool in cold exposure via systematic assessment of lipid profiles. Journal of Lipid Research, 2022, 63, 100197.	4.2	16
26	Development and characterization of patient-derived xenografts from non-small cell lung cancer brain metastases. Scientific Reports, 2021, 11, 2520.	3.3	13
27	Ultraviolet radiation-induced differential microRNA expression in the skin of hairless SKH1 mice, a widely used mouse model for dermatology research. Oncotarget, 2016, 7, 84924-84937.	1.8	12
28	Fibroblast Growth Factor Receptors as Targets for Radiosensitization in Head and Neck Squamous Cell Carcinomas. International Journal of Radiation Oncology Biology Physics, 2020, 107, 793-803.	0.8	10
29	Singleâ $\in$ cell technologies in reproductive immunology. American Journal of Reproductive Immunology, 2019, 82, e13157.	1.2	9
30	Keratin 13 deficiency causes white sponge nevus in mice. Developmental Biology, 2020, 468, 146-153.	2.0	8
31	MR1 overexpression correlates with poor clinical prognosis in glioma patients. Neuro-Oncology Advances, 2021, 3, vdab034.	0.7	7
32	CAMDA 2014: Making sense of RNA-Seq data: From low-level processing to functional analysis. Systems Biomedicine (Austin, Tex ), 2014, 2, 31-40.	0.7	6
33	MixTwice: large-scale hypothesis testing for peptide arrays by variance mixing. Bioinformatics, 2021, 37, 2637-2643.	4.1	6
34	Proteomic and Genomic Methylation Signatures of Idiopathic Subglottic Stenosis. Laryngoscope, 2021, 131, E540-E546.	2.0	5
35	Multiomic analysis reveals decidualâ€specific transcriptional programing of MAIT cells. American Journal of Reproductive Immunology, 2021, 86, e13495.	1.2	5
36	Structure-Leveraged Methods in Breast Cancer Risk Prediction. Journal of Machine Learning Research, 2016, 17, .	62.4	5

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37	SOX17 Is Essential for Integration of Arterial and HOXA Programs in Hemogenic Endothelium. Blood, 2019, 134, 2476-2476.	1.4	3
38	Discriminatory power of common genetic variants in personalized breast cancer diagnosis. , 2016, 9787, .		2
39	Enrichment of melanoma-associated T cells in 6-thioguanine-resistant T cells from metastatic melanoma patients. Melanoma Research, 2020, 30, 52-61.	1.2	2
40	Mediator complex subunit 12 is a gatekeeper of SARS-CoV-2 infection in breast cancer cells. Genes and Diseases, 2022, 9, 5-8.	3.4	2
41	GATA Factor-Dependent Positive-Feedback Circuit Controls Acute Myeloid Leukemia Cell Proliferation: Mechanisms and Targets for Therapeutic Intervention. Blood, 2016, 128, 1706-1706.	1.4	0