Hugo Horlings

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9471089/publications.pdf

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71 papers 16,654 citations

66234 42 h-index 76769 74 g-index

76 all docs 76 docs citations

76 times ranked 26945 citing authors

#	Article	IF	CITATIONS
1	Long non-coding RNA HOTAIR reprograms chromatin state to promote cancer metastasis. Nature, 2010, 464, 1071-1076.	13.7	4,648
2	A Functional Genetic Approach Identifies the PI3K Pathway as a Major Determinant of Trastuzumab Resistance in Breast Cancer. Cancer Cell, 2007, 12, 395-402.	7.7	1,471
3	Extensive and coordinated transcription of noncoding RNAs within cell-cycle promoters. Nature Genetics, 2011, 43, 621-629.	9.4	1,080
4	An Integrative Genomic and Proteomic Analysis of PIK3CA, PTEN, and AKT Mutations in Breast Cancer. Cancer Research, 2008, 68, 6084-6091.	0.4	916
5	Dysfunctional CD8 T Cells Form a Proliferative, Dynamically Regulated Compartment within Human Melanoma. Cell, 2019, 176, 775-789.e18.	13.5	760
6	Recurrent fusion of <i>MYB</i> and <i>NFIB</i> transcription factor genes in carcinomas of the breast and head and neck. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 18740-18744.	3.3	711
7	Immune induction strategies in metastatic triple-negative breast cancer to enhance the sensitivity to PD-1 blockade: the TONIC trial. Nature Medicine, 2019, 25, 920-928. Assessing Tumor-Infiltrating Lymphocytes in Solid Tumors: A Practical Review for Pathologists and	15.2	589
8	Proposal for a Standardized Method from the International Immuno-Oncology Biomarkers Working Group: Part 2: TILs in Melanoma, Gastrointestinal Tract Carcinomas, Non–Small Cell Lung Carcinoma and Mesothelioma, Endometrial and Ovarian Carcinomas, Squamous Cell Carcinoma of the Head and Neck, Genitourinary Carcinomas, and Primary Brain Tumors. Advances in Anatomic Pathology, 2017, 24,	2.4	530
9	311-335. Identification of CMTM6 and CMTM4 as PD-L1 protein regulators. Nature, 2017, 549, 106-110.	13.7	501
10	Assessing Tumor-infiltrating Lymphocytes in Solid Tumors: A Practical Review for Pathologists and Proposal for a Standardized Method From the International Immunooncology Biomarkers Working Group: Part 1: Assessing the Host Immune Response, TILs in Invasive Breast Carcinoma and Ductal Carcinoma In Situ, Metastatic Tumor Deposits and Areas for Further Research. Advances in Anatomic Pathology, 2017, 24, 235-251.	2.4	469
11	Cancer-Associated Mutations in Endometriosis without Cancer. New England Journal of Medicine, 2017, 376, 1835-1848.	13.9	451
12	Integrative molecular profiling of triple negative breast cancers identifies amplicon drivers and potential therapeutic targets. Oncogene, 2010, 29, 2013-2023.	2.6	385
13	MicroRNA Sequence and Expression Analysis in Breast Tumors by Deep Sequencing. Cancer Research, 2011, 71, 4443-4453.	0.4	331
14	Abrogation of BRAF ^{V600E} -induced senescence by PI3K pathway activation contributes to melanomagenesis. Genes and Development, 2012, 26, 1055-1069.	2.7	229
15	A case of meningoencephalitis by the relapsing fever spirochaete Borrelia miyamotoi in Europe. Lancet, The, 2013, 382, 658.	6.3	224
16	Genomic consequences of aberrant DNA repair mechanisms stratify ovarian cancer histotypes. Nature Genetics, 2017, 49, 856-865.	9.4	220
17	Genetic regulators of large-scale transcriptional signatures in cancer. Nature Genetics, 2006, 38, 421-430.	9.4	204
18	Gene Expression Profiling to Identify the Histogenetic Origin of Metastatic Adenocarcinomas of Unknown Primary. Journal of Clinical Oncology, 2008, 26, 4435-4441.	0.8	176

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19	High prevalence of oncogenic MYD88 and CD79B mutations in diffuse large B-cell lymphomas presenting at immune-privileged sites. Blood Cancer Journal, 2013, 3, e139-e139.	2.8	164
20	Identification of a pharmacologically tractable Fra- 1 /ADORA2B axis promoting breast cancer metastasis. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 5139-5144.	3.3	150
21	Retrospective analysis of metastatic behaviour of breast cancer subtypes. Breast Cancer Research and Treatment, 2015, 150, 547-557.	1.1	141
22	Synchronous Endometrial and Ovarian Carcinomas: Evidence of Clonality. Journal of the National Cancer Institute, 2015, 108, djv428.	3.0	128
23	Prediction of BRCA1-association in hereditary non-BRCA1/2 breast carcinomas with array-CGH. Breast Cancer Research and Treatment, 2009, 116, 479-489.	1.1	124
24	The tale of TILs in breast cancer: A report from The International Immuno-Oncology Biomarker Working Group. Npj Breast Cancer, 2021, 7, 150.	2.3	112
25	Mucinous and neuroendocrine breast carcinomas are transcriptionally distinct from invasive ductal carcinomas of no special type. Modern Pathology, 2009, 22, 1401-1414.	2.9	110
26	Pitfalls in assessing stromal tumor infiltrating lymphocytes (sTILs) in breast cancer. Npj Breast Cancer, 2020, 6, 17.	2.3	106
27	Molecular profiles of progesterone receptor loss in human breast tumors. Breast Cancer Research and Treatment, 2009, 114, 287-299.	1.1	94
28	High prevalence of oncogenic MYD88 and CD79B mutations in primary testicular diffuse large B-cell lymphoma. Leukemia, 2014, 28, 719-720.	3.3	91
29	Spatial immunophenotypes predict response to anti-PD1 treatment and capture distinct paths of T cell evasion in triple negative breast cancer. Nature Communications, 2021, 12, 5668.	5.8	91
30	Report on computational assessment of Tumor Infiltrating Lymphocytes from the International Immuno-Oncology Biomarker Working Group. Npj Breast Cancer, 2020, 6, 16.	2.3	90
31	Microarray-Based Determination of Estrogen Receptor, Progesterone Receptor, and HER2 Receptor Status in Breast Cancer. Clinical Cancer Research, 2009, 15, 7003-7011.	3.2	87
32	Cancer-immune interactions in ER-positive breast cancers: PI3K pathway alterations and tumor-infiltrating lymphocytes. Breast Cancer Research, 2019, 21, 90.	2.2	81
33	Assessment of PD-L1 expression across breast cancer molecular subtypes, in relation to mutation rate, <i>BRCA1</i> -like status, tumor-infiltrating immune cells and survival. Oncolmmunology, 2018, 7, e1509820.	2.1	80
34	latrogenic endometriosis harbors somatic cancer-driver mutations. Human Reproduction, 2019, 34, 69-78.	0.4	73
35	Reversal of pre-existing NGFR-driven tumor and immune therapy resistance. Nature Communications, 2020, 11, 3946.	5.8	71
36	Genomic profiling of histological special types of breast cancer. Breast Cancer Research and Treatment, 2013, 142, 257-269.	1.1	64

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37	Sorafenib synergizes with metformin in NSCLC through AMPK pathway activation. International Journal of Cancer, 2015, 136, 1434-1444.	2.3	64
38	Integration of DNA Copy Number Alterations and Prognostic Gene Expression Signatures in Breast Cancer Patients. Clinical Cancer Research, 2010, 16, 651-663.	3.2	61
39	Identification of breast cancer cell subtypes sensitive to ATG4B inhibition. Oncotarget, 2016, 7, 66970-66988.	0.8	58
40	ESR1 gene amplification in breast cancer: a common phenomenon?. Nature Genetics, 2008, 40, 807-808.	9.4	53
41	Divergent effects of insulin-like growth factor-1 receptor expression on prognosis of estrogen receptor positive versus triple negative invasive ductal breast carcinoma. Breast Cancer Research and Treatment, 2011, 129, 725-736.	1.1	53
42	TERT promoter mutation in adult granulosa cell tumor of the ovary. Modern Pathology, 2018, 31, 1107-1115.	2.9	49
43	External Validation of Adjuvant! Online Breast Cancer Prognosis Tool. Prioritising Recommendations for Improvement. PLoS ONE, 2011, 6, e27446.	1.1	38
44	Epithelial-to-mesenchymal transition status of primary breast carcinomas and its correlation with metastatic behavior. Breast Cancer Research and Treatment, 2019, 174, 649-659.	1.1	37
45	Characteristics and Outcome of <i>AKT1</i> i>E17K-Mutant Breast Cancer Defined through AACR Project GENIE, a Clinicogenomic Registry. Cancer Discovery, 2020, 10, 526-535.	7.7	36
46	Integration of Clinical and Gene Expression Data Has a Synergetic Effect on Predicting Breast Cancer Outcome. PLoS ONE, 2012, 7, e40358.	1.1	35
47	Frequent NFIB-associated Gene Rearrangement in Adenoid Cystic Carcinoma of the Vulva. International Journal of Gynecological Pathology, 2017, 36, 289-293.	0.9	32
48	The NF-κB Pathway Promotes Tamoxifen Tolerance and Disease Recurrence in Estrogen Receptor–Positive Breast Cancers. Molecular Cancer Research, 2020, 18, 1018-1027.	1.5	31
49	Comprehensive evaluation of methods to assess overall and cell-specific immune infiltrates in breast cancer. Breast Cancer Research, 2019, 21, 151.	2.2	30
50	Adultâ€type granulosa cell tumor of the ovary: a <scp><i>FOXL2</i></scp> â€centric disease. Journal of Pathology: Clinical Research, 2021, 7, 243-252.	1.3	27
51	Integrated molecular pathway analysis informs a synergistic combination therapy targeting PTEN/PI3K and EGFR pathways for basal-like breast cancer. BMC Cancer, 2016, 16, 587.	1.1	26
52	Prognostic factors in patients with oligometastatic breast cancer – A systematic review. Cancer Treatment Reviews, 2020, 91, 102114.	3.4	24
53	Characterization of Oligometastatic Disease in a Real-World Nationwide Cohort of 3447 Patients With de Novo Metastatic Breast Cancer. JNCI Cancer Spectrum, 2021, 5, pkab010.	1.4	21
54	Development and internal validation of a prognostic model to predict recurrence free survival in patients with adult granulosa cell tumors of the ovary. Gynecologic Oncology, 2014, 134, 498-504.	0.6	20

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55	Correlation Between Surrogate End Points and Overall Survival in a Multi-institutional Clinicogenomic Cohort of Patients With Non–Small Cell Lung or Colorectal Cancer. JAMA Network Open, 2021, 4, e2117547.	2.8	20
56	Application of a risk-management framework for integration of stromal tumor-infiltrating lymphocytes in clinical trials. Npj Breast Cancer, 2020, 6, 15.	2.3	16
57	A High-Dimensional Window into the Micro-Environment of Triple Negative Breast Cancer. Cancers, 2021, 13, 316.	1.7	16
58	Using Somatic Mutations to Guide Treatment Decisions. JAMA Oncology, 2015, 1, 275.	3.4	15
59	Functional Profiling of FSH and Estradiol in Ovarian Granulosa Cell Tumors. Journal of the Endocrine Society, 2020, 4, bvaa034.	0.1	13
60	Association of the germline TP53 R72P and MDM2 SNP309 variants with breast cancer survival in specific breast tumor subgroups. Breast Cancer Research and Treatment, 2011, 130, 599-608.	1.1	9
61	Predicting clinical benefit from everolimus in patients with advanced solid tumors, the CPCT-03 study. Oncotarget, 2017, 8, 55582-55592.	0.8	9
62	Low grade serous carcinoma of the peritoneum in a BRCA1 carrier previously diagnosed with a $\hat{a} \in \mathbb{C}$ consists a serous tubal intra-epithelial carcinoma $\hat{a} \in \mathbb{C}$ (STIC) on risk reducing surgery. Gynecologic Oncology Reports, 2015, 12, 72-74.	0.3	8
63	Interobserver Agreement of PD-L1/SP142 Immunohistochemistry and Tumor-Infiltrating Lymphocytes (TILs) in Distant Metastases of Triple-Negative Breast Cancer: A Proof-of-Concept Study. A Report on Behalf of the International Immuno-Oncology Biomarker Working Group. Cancers, 2021, 13, 4910.	1.7	8
64	LUNG TUMOR LOCATION AND LYMPHOCYTE INFILTRATION IN MICE ARE GENETICALLY DETERMINED. Experimental Lung Research, 2005, 31, 513-525.	0.5	5
65	Low probability of disease cure in advanced ovarian carcinomas before the PARP inhibitor era. British Journal of Cancer, 2022, 127, 79-83.	2.9	5
66	Left-sided native valve Staphylococcus aureus endocarditis. Netherlands Journal of Medicine, 2010, 68, 341-7.	0.6	4
67	Differential Survival and Therapy Benefit of Patients with Breast Cancer Are Characterized by Distinct Epithelial and Immune Cell Microenvironments. Clinical Cancer Research, 2022, 28, 960-971.	3.2	4
68	Spatial interplay of lymphocytes and fibroblasts in estrogen receptor-positive HER2-negative breast cancer. Npj Breast Cancer, 2022, 8, 56.	2.3	3
69	135TiP GELATO-trial: Assessing the efficacy of carboplatin and atezolizumab in metastatic lobular breast cancer. Annals of Oncology, 2020, 31, S61.	0.6	2
70	IHC-based Ki67 as response biomarker to tamoxifen in breast cancer window trials enrolling premenopausal women. Npj Breast Cancer, 2021, 7, 138.	2.3	1
71	Translating the Genomic Architecture of Breast Cancer into Clinical Applications. Science Translational Medicine, 2010, 2, 38ps32.	5.8	0