

Anna Ehinger

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2024412/publications.pdf>

Version: 2024-02-01

56
papers

4,589
citations

236925

25
h-index

243625

44
g-index

62
all docs

62
docs citations

62
times ranked

8753
citing authors

#	ARTICLE	IF	CITATIONS
1	Pan-cancer analysis of whole genomes. <i>Nature</i> , 2020, 578, 82-93.	27.8	1,966
2	Extensive transduction of nonrepetitive DNA mediated by L1 retrotransposition in cancer genomes. <i>Science</i> , 2014, 345, 1251343.	12.6	348
3	Origins and functional consequences of somatic mitochondrial DNA mutations in human cancer. <i>ELife</i> , 2014, 3, .	6.0	318
4	Whole-genome sequencing of triple-negative breast cancers in a population-based clinical study. <i>Nature Medicine</i> , 2019, 25, 1526-1533.	30.7	218
5	The path to a better biomarker: application of a risk management framework for the implementation of PD-L1 and TILs as immunology biomarkers in breast cancer clinical trials and daily practice. <i>Journal of Pathology</i> , 2020, 250, 667-684.	4.5	142
6	Spatial deconvolution of HER2-positive breast cancer delineates tumor-associated cell type interactions. <i>Nature Communications</i> , 2021, 12, 6012.	12.8	140
7	The Sweden Cancerome Analysis Network - Breast (SCAN-B) Initiative: a large-scale multicenter infrastructure towards implementation of breast cancer genomic analyses in the clinical routine. <i>Genome Medicine</i> , 2015, 7, 20.	8.2	129
8	The tale of TILs in breast cancer: A report from The International Immuno-Oncology Biomarker Working Group. <i>Npj Breast Cancer</i> , 2021, 7, 150.	5.2	112
9	Pitfalls in assessing stromal tumor infiltrating lymphocytes (sTILs) in breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 17.	5.2	106
10	Clinical Value of RNA Sequencing-Based Classifiers for Prediction of the Five Conventional Breast Cancer Biomarkers: A Report From the Population-Based Multicenter Sweden Cancerome Analysis Network Breast Initiative. <i>JCO Precision Oncology</i> , 2018, 2, 1-18.	3.0	101
11	Processed pseudogenes acquired somatically during cancer development. <i>Nature Communications</i> , 2014, 5, 3644.	12.8	86
12	Refinement of breast cancer molecular classification by miRNA expression profiles. <i>BMC Genomics</i> , 2019, 20, 503.	2.8	75
13	Analytical validation of a standardised scoring protocol for Ki67 immunohistochemistry on breast cancer excision whole sections: an international multicentre collaboration. <i>Histopathology</i> , 2019, 75, 225-235.	2.9	74
14	G protein-coupled estrogen receptor 1 (GPER, GPR 30) in normal human endometrium and early pregnancy decidua. <i>Molecular Human Reproduction</i> , 2010, 16, 743-751.	2.8	55
15	Comprehensive molecular comparison of BRCA1 hypermethylated and BRCA1 mutated triple negative breast cancers. <i>Nature Communications</i> , 2020, 11, 3747.	12.8	53
16	Expression of HIF-1 α is related to a poor prognosis and tamoxifen resistance in contralateral breast cancer. <i>PLoS ONE</i> , 2019, 14, e0226150.	2.5	52
17	Histological grade provides significant prognostic information in addition to breast cancer subtypes defined according to St Gallen 2013. <i>Acta Oncologica</i> , 2017, 56, 68-74.	1.8	51
18	The G protein-coupled estrogen receptor 1 (GPER/GPR30) does not predict survival in patients with ovarian cancer. <i>Journal of Ovarian Research</i> , 2012, 5, 9.	3.0	47

#	ARTICLE	IF	CITATIONS
19	Cross comparison and prognostic assessment of breast cancer multigene signatures in a large population-based contemporary clinical series. <i>Scientific Reports</i> , 2019, 9, 12184.	3.3	39
20	Differential localization and expression of urokinase plasminogen activator (uPA), its receptor (uPAR), and its inhibitor (PAI-1) mRNA and protein in endometrial tissue during the menstrual cycle. <i>Molecular Human Reproduction</i> , 2004, 10, 655-663.	2.8	38
21	Prediction of Lymph Node Metastasis in Breast Cancer by Gene Expression and Clinicopathological Models: Development and Validation within a Population-Based Cohort. <i>Clinical Cancer Research</i> , 2019, 25, 6368-6381.	7.0	37
22	The mutational landscape of the <sc>SCAN</sc> â€œrealâ€ world primary breast cancer transcriptome. <i>EMBO Molecular Medicine</i> , 2020, 12, e12118.	6.9	36
23	Minimizing inequality in access to precision medicine in breast cancer by real-time population-based molecular analysis in the SCAN-B initiative. <i>British Journal of Surgery</i> , 2018, 105, e158-e168.	0.3	32
24	Variability in Breast Cancer Biomarker Assessment and the Effect on Oncological Treatment Decisions: A Nationwide 5-Year Population-Based Study. <i>Cancers</i> , 2021, 13, 1166.	3.7	31
25	Changing clinical presentation of angiosarcomas after breast cancer: from late tumors in edematous arms to earlier tumors on the thoracic wall. <i>Breast Cancer Research and Treatment</i> , 2010, 122, 883-887.	2.5	30
26	Contralateral breast cancer can represent a metastatic spread of the first primary tumor: determination of clonal relationship between contralateral breast cancers using next-generation whole genome sequencing. <i>Breast Cancer Research</i> , 2015, 17, 102.	5.0	30
27	An Open-Source, Automated Tumor-Infiltrating Lymphocyte Algorithm for Prognosis in Triple-Negative Breast Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 5557-5565.	7.0	26
28	Prognostic implications of the expression levels of different immunoglobulin heavy chain-encoding RNAs in early breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 28.	5.2	25
29	Agreement between molecular subtyping and surrogate subtype classification: a contemporary population-based study of ER-positive/HER2-negative primary breast cancer. <i>Breast Cancer Research and Treatment</i> , 2019, 178, 459-467.	2.5	23
30	Molecular analyses of triple-negative breast cancer in the young and elderly. <i>Breast Cancer Research</i> , 2021, 23, 20.	5.0	23
31	Preexisting Somatic Mutations of Estrogen Receptor Alpha (<i>ESR1</i>) in Early-Stage Primary Breast Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab028.	2.9	20
32	Distinct mechanisms of resistance to fulvestrant treatment dictate level of ER independence and selective response to CDK inhibitors in metastatic breast cancer. <i>Breast Cancer Research</i> , 2021, 23, 26.	5.0	19
33	Systematically higher Ki67 scores on core biopsy samples compared to corresponding resection specimen in breast cancer: a multi-operator and multi-institutional study. <i>Modern Pathology</i> , 2022, 35, 1362-1369.	5.5	18
34	Histamine uptake by human endometrial cells expressing the organic cation transporter EMT and the vesicular monoamine transporter-2. <i>Molecular Human Reproduction</i> , 2006, 12, 483-489.	2.8	13
35	Prior Adjuvant Tamoxifen Treatment in Breast Cancer Is Linked to Increased AIB1 and HER2 Expression in Metachronous Contralateral Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0150977.	2.5	9
36	Prognosis, stage and oestrogen receptor status of contralateral breast cancer in relation to characteristics of the first tumour, prior endocrine treatment and radiotherapy. <i>European Journal of Cancer</i> , 2015, 51, 2304-2313.	2.8	8

#	ARTICLE	IF	CITATIONS
37	Breast metastases from pancreatic and ovarian carcinoma. <i>Diagnostic Cytopathology</i> , 1999, 21, 154-155.	1.0	7
38	Difficulties in diagnostics of lung tumours in biopsies: an interpathologist concordance study evaluating the international diagnostic guidelines. <i>Journal of Clinical Pathology</i> , 2022, 75, 302-309.	2.0	7
39	The Prognostic Role of Intratumoral Stromal Content in Lobular Breast Cancer. <i>Cancers</i> , 2022, 14, 941.	3.7	5
40	Stability of oestrogen and progesterone receptor antigenicity in formalin-fixed paraffin-embedded breast cancer tissue over time. <i>Apmis</i> , 2018, 126, 746-754.	2.0	4
41	Development of Training Materials for Pathologists to Provide Machine Learning Validation Data of Tumor-Infiltrating Lymphocytes in Breast Cancer. <i>Cancers</i> , 2022, 14, 2467.	3.7	4
42	Acute pancreatitis evoked by small-cell lung carcinoma metastases and detected by endoscopic ultrasound. <i>Endoscopy</i> , 2012, 44, E45-E46.	1.8	3
43	Paracrine Stimulation of Capillary Endothelial Cell Migration by Endometrial Tissue Involves Epidermal Growth Factor and Is Mediated Via Up-Regulation of the Urokinase Plasminogen Activator Receptor. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2001, 86, 1724-1730.	3.6	3
44	Defining the mutational landscape of 3,217 primary breast cancer transcriptomes through large-scale RNA-seq within the Sweden Cancerome Analysis Network: Breast Project (SCAN-B; NCT03430492).. <i>Journal of Clinical Oncology</i> , 2020, 38, 518-518.	1.6	2
45	Abstract P4-09-03: On the development and clinical value of RNA-sequencing-based classifiers for prediction of the five conventional breast cancer biomarkers: A report from the population-based multicenter SCAN-B study. , 2018, , .		1
46	Abstract CT074: Pre-existing ESR1 mutations in early-stage primary breast cancer predict failure of endocrine therapy and poor survival. , 2020, , .		1
47	Abstract P6-08-43: Histological grade provides significant prognostic information in the discrimination between luminal A-like and luminal B-like HER-2 normal subtypes of breast cancer according to St Gallen 2013. , 2015, , .		0
48	Abstract P1-07-17: The SCAN-B study: 5-year summary of a large-scale population-based prospective breast cancer translational genomics platform covering a wide geography of Sweden (NCT02306096).. , 2017, , .		0
49	Abstract P2-02-09: Breast cancer subtype distribution and circulating tumor DNA in response to neoadjuvant chemotherapy: Experiences from a preoperative cohort within SCAN-B. , 2018, , .		0
50	Abstract P2-03-01: Analytical validation of a standardized scoring protocol for Ki67 assessed on breast excision whole sections: An international multicenter collaboration. , 2018, , .		0
51	Abstract P1-06-01: Putting multigene signatures to the test: Prognostic assessment in population-based contemporary clinical breast cancer. , 2018, , .		0
52	Abstract P3-02-02: Concordance between immunohistochemical and gene-expression based subtyping of early breast cancer using core needle biopsies and surgical specimens - experiences from SCAN-B. , 2018, , .		0
53	Abstract P1-18-10: Preoperative treatment of HER2-positive breast cancer in South Sweden. A retrospective, comprehensive survey of neo-adjuvant treated HER2-positive breast cancer in the SCAN-B project 2010-2017. , 2020, , .		0
54	Abstract P5-02-01: Analytical validation and prognostic potential of an automated digital scoring protocol for Ki67: An International Ki67 Working Group study. , 2020, , .		0

#	ARTICLE	IF	CITATIONS
55	Abstract P2-08-11: How reliable are biomarkers assessed on a core needle biopsy? A study of paired core needle biopsies and surgical specimens in early breast cancer. Cancer Research, 2022, 82, P2-08-11-P2-08-11.	0.9	0
56	Abstract 460: Tools for collecting pathologist annotations and understanding interobserver variability. Cancer Research, 2022, 82, 460-460.	0.9	0