

Andrew H Beck

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9604606/publications.pdf>

Version: 2024-02-01

47
papers

7,676
citations

201674

27
h-index

254184

43
g-index

48
all docs

48
docs citations

48
times ranked

14725
citing authors

#	ARTICLE	IF	CITATIONS
1	Human-interpretable image features derived from densely mapped cancer pathology slides predict diverse molecular phenotypes. <i>Nature Communications</i> , 2021, 12, 1613.	12.8	114
2	A Machine Learning Approach Enables Quantitative Measurement of Liver Histology and Disease Monitoring in NASH. <i>Hepatology</i> , 2021, 74, 133-147.	7.3	101
3	A Machine Learning Approach to Liver Histological Evaluation Predicts Clinically Significant Portal Hypertension in NASH Cirrhosis. <i>Hepatology</i> , 2021, 74, 3146-3160.	7.3	25
4	Report on computational assessment of Tumor Infiltrating Lymphocytes from the International Immuno-Oncology Biomarker Working Group. <i>Npj Breast Cancer</i> , 2020, 6, 16.	5.2	90
5	<i>EN1</i> Is a Transcriptional Dependency in Triple-Negative Breast Cancer Associated with Brain Metastasis. <i>Cancer Research</i> , 2019, 79, 4173-4183.	0.9	47
6	Predicting breast tumor proliferation from whole-slide images: The TUPAC16 challenge. <i>Medical Image Analysis</i> , 2019, 54, 111-121.	11.6	182
7	Application of convolutional neural networks to breast biopsies to delineate tissue correlates of mammographic breast density. <i>Npj Breast Cancer</i> , 2019, 5, 43.	5.2	12
8	SPOP Promotes Nanog Destruction to Suppress Stem Cell Traits and Prostate Cancer Progression. <i>Developmental Cell</i> , 2019, 48, 329-344.e5.	7.0	53
9	Molecular mechanisms linking high body mass index to breast cancer etiology in post-menopausal breast tumor and tumor-adjacent tissues. <i>Breast Cancer Research and Treatment</i> , 2019, 173, 667-677.	2.5	19
10	Using deep convolutional neural networks to identify and classify tumor-associated stroma in diagnostic breast biopsies. <i>Modern Pathology</i> , 2018, 31, 1502-1512.	5.5	145
11	Crowdsourcing scoring of immunohistochemistry images: Evaluating Performance of the Crowd and an Automated Computational Method. <i>Scientific Reports</i> , 2017, 7, 43286.	3.3	31
12	Deep learning-based assessment of tumor-associated stroma for diagnosing breast cancer in histopathology images. , 2017, 2017, 929-932.		27
13	Aspirin Suppresses Growth in PI3K-Mutant Breast Cancer by Activating AMPK and Inhibiting mTORC1 Signaling. <i>Cancer Research</i> , 2017, 77, 790-801.	0.9	96
14	Breast cancer risk factors in relation to estrogen receptor, progesterone receptor, insulin-like growth factor-1 receptor, and Ki67 expression in normal breast tissue. <i>Npj Breast Cancer</i> , 2017, 3, 39.	5.2	27
15	Prostate cancer-associated SPOP mutations confer resistance to BET inhibitors through stabilization of BRD4. <i>Nature Medicine</i> , 2017, 23, 1063-1071.	30.7	240
16	Nanoscale imaging of clinical specimens using pathology-optimized expansion microscopy. <i>Nature Biotechnology</i> , 2017, 35, 757-764.	17.5	182
17	Diagnostic Assessment of Deep Learning Algorithms for Detection of Lymph Node Metastases in Women With Breast Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2017, 318, 2199.	7.4	2,003
18	The molecular basis of breast cancer pathological phenotypes. <i>Journal of Pathology</i> , 2017, 241, 375-391.	4.5	86

#	ARTICLE	IF	CITATIONS
19	Deep learning assessment of tumor proliferation in breast cancer histological images. , 2017, , .		21
20	Alcohol consumption and breast tumor gene expression. Breast Cancer Research, 2017, 19, 108.	5.0	23
21	LINC00520 is induced by Src, STAT3, and PI3K and plays a functional role in breast cancer. Oncotarget, 2016, 7, 81981-81994.	1.8	48
22	Safikhani et al. reply. Nature, 2016, 540, E2-E4.	27.8	22
23	Safikhani et al. reply. Nature, 2016, 540, E6-E8.	27.8	10
24	Safikhani et al. reply. Nature, 2016, 540, E11-E12.	27.8	11
25	DNA defects, epigenetics, and gene expression in cancer-adjacent breast: a study from The Cancer Genome Atlas. Npj Breast Cancer, 2016, 2, 16007.	5.2	33
26	Oncogenic Role of Fusion-circRNAs Derived from Cancer-Associated Chromosomal Translocations. Cell, 2016, 165, 289-302.	28.9	567
27	TNF- α expression, risk factors, and inflammatory exposures in ovarian cancer: evidence for an inflammatory pathway of ovarian carcinogenesis?. Human Pathology, 2016, 54, 82-91.	2.0	45
28	Antibody Therapy Targeting CD47 and CD271 Effectively Suppresses Melanoma Metastasis in Patient-Derived Xenografts. Cell Reports, 2016, 16, 1701-1716.	6.4	56
29	A <i>BRCA1/2</i> Mutational Signature and Survival in Ovarian High-Grade Serous Carcinoma. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1511-1516.	2.5	16
30	The SIRT2 Deacetylase Stabilizes Slug to Control Malignancy of Basal-like Breast Cancer. Cell Reports, 2016, 17, 1302-1317.	6.4	85
31	Region of interest identification and diagnostic agreement in breast pathology. Modern Pathology, 2016, 29, 1004-1011.	5.5	17
32	PharmacoGx: an R package for analysis of large pharmacogenomic datasets. Bioinformatics, 2016, 32, 1244-1246.	4.1	249
33	Progress in Medicine: Experts Take Stock. PLoS Medicine, 2015, 12, e1001933.	8.4	2
34	Etiologic field effect: reappraisal of the field effect concept in cancer predisposition and progression. Modern Pathology, 2015, 28, 14-29.	5.5	172
35	NFAT1 promotes intratumoral neutrophil infiltration by regulating IL8 expression in breast cancer. Molecular Oncology, 2015, 9, 1140-1154.	4.6	59
36	Open Access to Large Scale Datasets Is Needed to Translate Knowledge of Cancer Heterogeneity into Better Patient Outcomes. PLoS Medicine, 2015, 12, e1001794.	8.4	14

#	ARTICLE	IF	CITATIONS
37	MERIT40 Is an Akt Substrate that Promotes Resolution of DNA Damage Induced by Chemotherapy. <i>Cell Reports</i> , 2015, 11, 1358-1366.	6.4	40
38	Noninvasive Imaging of Tumor Burden and Molecular Pathways in Mouse Models of Cancer. <i>Cold Spring Harbor Protocols</i> , 2015, 2015, pdb.top069930.	0.3	28
39	Comprehensive Molecular Portraits of Invasive Lobular Breast Cancer. <i>Cell</i> , 2015, 163, 506-519.	28.9	1,485
40	SPOP Promotes Ubiquitination and Degradation of the ERG Oncoprotein to Suppress Prostate Cancer Progression. <i>Molecular Cell</i> , 2015, 59, 917-930.	9.7	172
41	The Reprogramming of Tumor Stroma by HSF1 Is a Potent Enabler of Malignancy. <i>Cell</i> , 2014, 158, 564-578.	28.9	298
42	Increased rate of atypical squamous cells of undetermined significance and declining high-risk human papillomavirus rates following implementation of ThinPrep Imaging System are associated with increased nuclear chromasia. <i>Journal of the American Society of Cytopathology</i> , 2014, 3, 73-78.	0.5	1
43	Computational Pathology to Discriminate Benign from Malignant Intraductal Proliferations of the Breast. <i>PLoS ONE</i> , 2014, 9, e114885.	2.5	106
44	Chromosomal copy number alterations (CNAs) for risk assessment of ductal carcinoma in situ (DCIS).. <i>Journal of Clinical Oncology</i> , 2014, 32, 565-565.	1.6	0
45	Application of Image-Guided Coring as a new technique for targeting breast tumor tissue in molecular pathology. <i>FASEB Journal</i> , 2013, 27, lb460.	0.5	0
46	Evaluation of a Gene Expression Microarray-based Assay to Determine Tissue Type of Origin on a Diverse Set of 49 Malignancies. <i>American Journal of Surgical Pathology</i> , 2011, 35, 1030-1037.	3.7	12
47	Systematic Analysis of Breast Cancer Morphology Uncovers Stromal Features Associated with Survival. <i>Science Translational Medicine</i> , 2011, 3, 108ra113.	12.4	603