Joan S Brugge

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Clonal populations of a human TNBC model display significant functional heterogeneity and divergent growth dynamics in distinct contexts. Oncogene, 2022, 41, 112-124.	5.9	6
2	Abstract P5-01-02: Single cell RNA transcriptomics reveals tumor promoting mammary cell subpopulation upon replication stress in <i>BRCA1</i> mutant breast cancer mouse model. Cancer Research, 2022, 82, P5-01-02-P5-01-02.	0.9	0
3	Therapy resistance: opportunities created by adaptive responses to targeted therapies in cancer. Nature Reviews Cancer, 2022, 22, 323-339.	28.4	107
4	Long-term culture, genetic manipulation and xenotransplantation of human normal and breast cancer organoids. Nature Protocols, 2021, 16, 1936-1965.	12.0	97
5	Clinical evaluation of BCL-2/XL levels pre- and post- HER2-targeted therapy. PLoS ONE, 2021, 16, e0251163.	2.5	9
6	Metabolic perturbations sensitize triple-negative breast cancers to apoptosis induced by BH3 mimetics. Science Signaling, 2021, 14, .	3.6	10
7	Cycling cancer persister cells arise from lineages with distinct programs. Nature, 2021, 596, 576-582.	27.8	236
8	Pathologic and molecular responses to neoadjuvant trastuzumab and/or lapatinib from a phase II randomized trial in HER2-positive breast cancer (TRIO-US B07). Nature Communications, 2020, 11, 5824.	12.8	42
9	Transient commensal clonal interactions can drive tumor metastasis. Nature Communications, 2020, 11, 5799.	12.8	30
10	Navitoclax enhances the effectiveness of EGFR-targeted antibody-drug conjugates in PDX models of EGFR-expressing triple-negative breast cancer. Breast Cancer Research, 2020, 22, 132.	5.0	19
11	3D Culture Models with CRISPR Screens Reveal Hyperactive NRF2 as a Prerequisite for Spheroid Formation via Regulation of Proliferation and Ferroptosis. Molecular Cell, 2020, 80, 828-844.e6.	9.7	110
12	Large-Scale Characterization of Drug Responses of Clinically Relevant Proteins in Cancer Cell Lines. Cancer Cell, 2020, 38, 829-843.e4.	16.8	40
13	Fibroblast–tumor cell signaling limits HER2 kinase therapy response via activation of MTOR and antiapoptotic pathways. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 16500-16508.	7.1	23
14	Organoid cultures from normal and cancer-prone human breast tissues preserve complex epithelial lineages. Nature Communications, 2020, 11, 1711.	12.8	134
15	Aging-Associated Alterations in Mammary Epithelia and Stroma Revealed by Single-Cell RNA Sequencing. Cell Reports, 2020, 33, 108566.	6.4	75
16	Synthetic Lethal and Resistance Interactions with BET Bromodomain Inhibitors in Triple-Negative Breast Cancer. Molecular Cell, 2020, 78, 1096-1113.e8.	9.7	114
17	Characterization of Mammary Cells Coâ€expressing Separate Lineage Markers. FASEB Journal, 2020, 34, 1-1.	0.5	0
18	United They Stand, Divided They Fall. Cell Metabolism, 2019, 30, 624-625.	16.2	3

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19	Combined MEK and BCL-2/XL Inhibition Is Effective in High-Grade Serous Ovarian Cancer Patient–Derived Xenograft Models and BIM Levels Are Predictive of Responsiveness. Molecular Cancer Therapeutics, 2019, 18, 642-655.	4.1	39
20	Role for polo-like kinase 4 in mediation of cytokinesis. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 11309-11318.	7.1	30
21	Critical questions in ovarian cancer research and treatment: Report of an American Association for Cancer Research Special Conference. Cancer, 2019, 125, 1963-1972.	4.1	39
22	Neutralization of BCL-2/XL Enhances the Cytotoxicity of T-DM1 <i>In Vivo</i> . Molecular Cancer Therapeutics, 2019, 18, 1115-1126.	4.1	20
23	Deubiquitinases Maintain Protein Homeostasis and Survival of Cancer Cells upon Glutathione Depletion. Cell Metabolism, 2019, 29, 1166-1181.e6.	16.2	121
24	CRB3 and the FERM protein EPB41L4B regulate proliferation of mammary epithelial cells through the release of amphiregulin. PLoS ONE, 2018, 13, e0207470.	2.5	3
25	Cancer Cells Co-opt the Neuronal Redox-Sensing Channel TRPA1 to Promote Oxidative-Stress Tolerance. Cancer Cell, 2018, 33, 985-1003.e7.	16.8	184
26	Starved epithelial cells uptake extracellular matrix for survival. Nature Communications, 2017, 8, 13989.	12.8	91
27	Metabolic changes promote rejection of oncogenic cells. Nature Cell Biology, 2017, 19, 414-415.	10.3	6
28	Rational combination therapy with PARP and MEK inhibitors capitalizes on therapeutic liabilities in <i>RAS</i> mutant cancers. Science Translational Medicine, 2017, 9, .	12.4	174
29	Not just Salk. Science, 2017, 357, 1105-1106.	12.6	4
30	ldentification of cancer genes that are independent of dominant proliferation and lineage programs. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E11276-E11284.	7.1	20
31	Niche-localized tumor cells are protected from HER2-targeted therapy via upregulation of an anti-apoptotic program in vivo. Npj Breast Cancer, 2017, 3, 18.	5.2	18
32	Establishment of Patient-Derived Tumor Xenograft Models of Epithelial Ovarian Cancer for Preclinical Evaluation of Novel Therapeutics. Clinical Cancer Research, 2017, 23, 1263-1273.	7.0	95
33	Systems analysis of apoptotic priming in ovarian cancer identifies vulnerabilities and predictors of drug response. Nature Communications, 2017, 8, 365.	12.8	44
34	Akt regulation of glycolysis mediates bioenergetic stability in epithelial cells. ELife, 2017, 6, .	6.0	55
35	Differential Glutamate Metabolism in Proliferating and Quiescent Mammary Epithelial Cells. Cell Metabolism, 2016, 23, 867-880.	16.2	214
36	Cytokinesis involves a nontranscriptional function of the Hippo pathway effector YAP. Science Signaling, 2016, 9, ra23.	3.6	53

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37	ERK and p38 MAPK Activities Determine Sensitivity to PI3K/mTOR Inhibition via Regulation of MYC and YAP. Cancer Research, 2016, 76, 7168-7180.	0.9	53
38	The Role of Proliferation in Determining Response to Neoadjuvant Chemotherapy in Breast Cancer: A Gene Expression–Based Meta-Analysis. Clinical Cancer Research, 2016, 22, 6039-6050.	7.0	48
39	Coping with the metabolic stress of leaving home. Cell Research, 2016, 26, 757-758.	12.0	5
40	Mutant p53 regulates ovarian cancer transformed phenotypes through autocrine matrix deposition. JCI Insight, 2016, 1, .	5.0	45
41	Moving Closer To Victory. Cold Spring Harbor Symposia on Quantitative Biology, 2016, 81, 281-288.	1.1	0
42	Signal Transduction in Cancer. Cold Spring Harbor Perspectives in Medicine, 2015, 5, a006098-a006098.	6.2	665
43	The enemy of my enemy is my friend. Nature, 2015, 527, 170-171.	27.8	47
44	Characterization of twenty-five ovarian tumour cell lines that phenocopy primary tumours. Nature Communications, 2015, 6, 7419.	12.8	149
45	Meta-analysis of breast cancer expression data using published gene signatures to reveal key cellular processes implicated in chemosensitivity and resistance Journal of Clinical Oncology, 2015, 33, 509-509.	1.6	1
46	Mapping the dynamics of force transduction at cell–cell junctions of epithelial clusters. ELife, 2014, 3, e03282.	6.0	99
47	Oncogene-like induction of cellular invasion from centrosome amplification. Nature, 2014, 510, 167-171.	27.8	360
48	Mesenchymal gene program–expressing ovarian cancer spheroids exhibit enhanced mesothelial clearance. Journal of Clinical Investigation, 2014, 124, 2611-2625.	8.2	110
49	Into the deep: Refocusing on 3D. Nature Cell Biology, 2012, 14, 332-332.	10.3	5
50	In vitro Mesothelial Clearance Assay that Models the Early Steps of Ovarian Cancer Metastasis. Journal of Visualized Experiments, 2012, , .	0.3	36
51	The myosin-II-responsive focal adhesion proteome: a tour de force?. Nature Cell Biology, 2011, 13, 344-346.	10.3	4
52	A New Mutational aktivation in the PI3K Pathway. Cancer Cell, 2007, 12, 104-107.	16.8	230
53	Casting light on focal adhesions. Nature Genetics, 1998, 19, 309-311.	21.4	25