

Alexander S Brodsky

List of Publications by Year in descending order

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33
papers

3,597
citations

394286

19
h-index

395590

33
g-index

33
all docs

33
docs citations

33
times ranked

5821
citing authors

#	ARTICLE	IF	CITATIONS
1	Somatic mutations in collagens are associated with a distinct tumor environment and overall survival in gastric cancer. <i>BMC Cancer</i> , 2022, 22, 139.	1.1	9
2	Insights from the Menstrual Cycle in Pulmonary Arterial Hypertension. <i>Annals of the American Thoracic Society</i> , 2021, 18, 218-228.	1.5	15
3	Effects of High Fat Versus Normal Diet on Extracellular Vesicle-Induced Angiogenesis in a Swine Model of Chronic Myocardial Ischemia. <i>Journal of the American Heart Association</i> , 2021, 10, e017437.	1.6	17
4	Senescent Tissue-Resident Mesenchymal Stromal Cells Are an Internal Source of Inflammation in Human Osteoarthritic Cartilage. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 725071.	1.8	11
5	Mesenchymal Stem Cell Extracellular Vesicles Reverse Sugen/Hypoxia Pulmonary Hypertension in Rats. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2020, 62, 577-587.	1.4	54
6	Stromal ColX \pm 1 expression correlates with tumor-infiltrating lymphocytes and predicts adjuvant therapy outcome in ER-positive/HER2-positive breast cancer. <i>BMC Cancer</i> , 2019, 19, 1036.	1.1	4
7	ColX \pm 1 is a stromal component that colocalizes with elastin in the breast tumor extracellular matrix. <i>Journal of Pathology: Clinical Research</i> , 2019, 5, 40-52.	1.3	3
8	Stromal Clusterin Expression Predicts Therapeutic Response to Neoadjuvant Chemotherapy in Triple Negative Breast Cancer. <i>Clinical Breast Cancer</i> , 2018, 18, e373-e379.	1.1	9
9	Fatty acid-binding protein 1 is preferentially lost in microsatellite instable colorectal carcinomas and is immune modulated via the interferon β pathway. <i>Modern Pathology</i> , 2017, 30, 123-133.	2.9	18
10	TAF4b Regulates Oocyte-Specific Genes Essential for Meiosis. <i>PLoS Genetics</i> , 2016, 12, e1006128.	1.5	24
11	Collagen type III \pm 1 as a useful diagnostic immunohistochemical marker for fibroepithelial lesions of the breast. <i>Human Pathology</i> , 2016, 57, 176-181.	1.1	14
12	Identification of stromal ColX \pm 1 and tumor-infiltrating lymphocytes as putative predictive markers of neoadjuvant therapy in estrogen receptor-positive/HER2-positive breast cancer. <i>BMC Cancer</i> , 2016, 16, 274.	1.1	42
13	Medullary carcinoma of the colon: a distinct morphology reveals a distinctive immunoregulatory microenvironment. <i>Modern Pathology</i> , 2016, 29, 528-541.	2.9	60
14	Cisplatin Resistant Spheroids Model Clinically Relevant Survival Mechanisms in Ovarian Tumors. <i>PLoS ONE</i> , 2016, 11, e0151089.	1.1	39
15	Notch3 Overexpression Promotes Anoikis Resistance in Epithelial Ovarian Cancer via Upregulation of COL4A2. <i>Molecular Cancer Research</i> , 2015, 13, 78-85.	1.5	36
16	Expression Profiling of Primary and Metastatic Ovarian Tumors Reveals Differences Indicative of Aggressive Disease. <i>PLoS ONE</i> , 2014, 9, e94476.	1.1	66
17	Oxysterols synergize with statins by inhibiting SREBP-2 in ovarian cancer cells. <i>Gynecologic Oncology</i> , 2014, 135, 333-341.	0.6	24
18	Prognostic microRNA expression signature from examination of colorectal primary and metastatic tumors. <i>Anticancer Research</i> , 2014, 34, 3957-67.	0.5	21

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19	Identification of Ovarian Cancer Metastatic miRNAs. <i>PLoS ONE</i> , 2013, 8, e58226.	1.1	78
20	T090137 Inhibits Cisplatin-Induced Apoptosis in Ovarian Cancer Cells. <i>International Journal of Gynecological Cancer</i> , 2011, 21, 1350-1356.	1.2	8
21	Integrated genomics of ovarian xenograft tumor progression and chemotherapy response. <i>BMC Cancer</i> , 2011, 11, 308.	1.1	10
22	Estrogen Coordinates Translation and Transcription, Revealing a Role for NRSF in Human Breast Cancer Cells. <i>Molecular Endocrinology</i> , 2010, 24, 1120-1135.	3.7	20
23	Exon expression profiling reveals stimulus-mediated exon use in neural cells. <i>Genome Biology</i> , 2007, 8, R159.	13.9	36
24	Genome-wide analysis of estrogen receptor binding sites. <i>Nature Genetics</i> , 2006, 38, 1289-1297.	9.4	1,227
25	Genomic localization of RNA binding proteins reveals links between pre-mRNA processing and transcription. <i>Genome Research</i> , 2006, 16, 912-921.	2.4	51
26	Chromosome-Wide Mapping of Estrogen Receptor Binding Reveals Long-Range Regulation Requiring the Forkhead Protein FoxA1. <i>Cell</i> , 2005, 122, 33-43.	13.5	1,208
27	Genomic mapping of RNA polymerase II reveals sites of co-transcriptional regulation in human cells. <i>Genome Biology</i> , 2005, 6, R64.	13.9	78
28	Analysis of RNA-protein interactions by flow cytometry. <i>Current Opinion in Molecular Therapeutics</i> , 2003, 5, 235-40.	2.8	4
29	A Microbead-based System for Identifying and Characterizing RNA-Protein Interactions by Flow Cytometry. <i>Molecular and Cellular Proteomics</i> , 2002, 1, 922-929.	2.5	14
30	Base flexibility in HIV-2 TAR RNA mapped by solution ¹⁵ N, ¹³ C NMR relaxation. <i>Journal of Molecular Biology</i> , 2002, 317, 263-278.	2.0	71
31	Pre-mRNA processing factors are required for nuclear export. <i>Rna</i> , 2000, 6, 1737-1749.	1.6	161
32	NMR evidence for a base triple in the HIV-2 TAR C-G.C+ mutant- argininamide complex. <i>Nucleic Acids Research</i> , 1998, 26, 1991-1995.	6.5	28
33	Solution structure of the HIV-2 TAR-argininamide complex 1 Edited by I. Tinoco. <i>Journal of Molecular Biology</i> , 1997, 267, 624-639.	2.0	137