

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5863508/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Ameliorative effect of recombinant human lactoferrin on the premature ovarian failure in rats after cyclophosphamide treatments. Journal of Ovarian Research, 2021, 14, 17.	1.3	16
2	High methionyl–tRNA synthetase expression predicts poor prognosis in patients with breast cancer. Journal of Clinical Pathology, 2020, 73, 803-812.	1.0	8
3	High kinesin family member 11 expression predicts poor prognosis in patients with clear cell renal cell carcinoma. Journal of Clinical Pathology, 2019, 72, 354-362.	1.0	28
4	Human breast cancer decellularized scaffolds promote epithelialâ€ŧoâ€mesenchymal transitions and stemness of breast cancer cells in vitro. Journal of Cellular Physiology, 2019, 234, 9447-9456.	2.0	39
5	Decellularized breast matrix as bioactive microenvironment for in vitro threeâ€dimensional cancer culture. Journal of Cellular Physiology, 2019, 234, 3425-3435.	2.0	49
6	Overexpression of FIBCD1 Is Predictive of Poor Prognosis in Gastric Cancer. American Journal of Clinical Pathology, 2018, 149, 474-483.	0.4	10
7	Downregulated SASH1 expression indicates poor clinical prognosis in gastric cancer. Human Pathology, 2018, 74, 83-91.	1.1	14
8	Role of M2 Macrophages in Sepsis-Induced Acute Kidney Injury. Shock, 2018, 50, 233-239.	1.0	52
9	High Spy1 expression predicts poor prognosis in colorectal cancer. Cancer Management and Research, 2018, Volume 10, 2757-2765.	0.9	2
10	Effects of Rab27A and Rab27B on Invasion, Proliferation, Apoptosis, and Chemoresistance in Human Pancreatic Cancer Cells. Pancreas, 2017, 46, 1173-1179.	0.5	30
11	Aldehyde dehydrogenase 1 expression is correlated with poor prognosis in breast cancer. Medicine (United States), 2017, 96, e7171.	0.4	28
12	LAMP1 expression is associated with poor prognosis in breast cancer. Oncology Letters, 2017, 14, 4729-4735.	0.8	27
13	Triple-amiRNA VEGFRs inhibition in pancreatic cancer improves the efficacy of chemotherapy through EMT regulation. Journal of Controlled Release, 2017, 245, 1-14.	4.8	27
14	High expression of PFTK1 in cancer cells predicts poor prognosis in colorectal cancer. Molecular Medicine Reports, 2017, 16, 224-230.	1.1	8
15	Reduced expression of EphA5 is associated with lymph node metastasis, advanced TNM stage, and poor prognosis in colorectal carcinoma. Histology and Histopathology, 2017, 32, 491-497.	0.5	13
16	High TMPRSS11D protein expression predicts poor overall survival in non-small cell lung cancer. Oncotarget, 2017, 8, 12812-12819.	0.8	29
17	High Eg5 expression predicts poor prognosis in breast cancer. Oncotarget, 2017, 8, 62208-62216.	0.8	29
18	High expression of EphA4 is associated with invasion and lymph node metastasis in colorectal carcinomas. International Journal of Clinical and Experimental Pathology, 2017, 10, 9697-9703.	0.5	1

Qin Jin

#	Article	IF	CITATIONS
19	EphA8 is a prognostic marker for epithelial ovarian cancer. Oncotarget, 2016, 7, 20801-20809.	0.8	20
20	Association of High Expression of Groβwith Clinical and Pathological Characteristics of Unfavorable Prognosis in Gastrointestinal Stromal Tumors. Disease Markers, 2015, 2015, 1-8.	0.6	6
21	Plasma miR-185 is decreased in patients with esophageal squamous cell carcinoma and might suppress tumor migration and invasion by targeting RAGE. American Journal of Physiology - Renal Physiology, 2015, 309, G719-G729.	1.6	19