

Comprehensive analysis of normal adjacent to tumor tr

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Prostate cancers that $\hat{\epsilon}$ Wnt $\hat{\epsilon}$ ™ respond to abiraterone. <i>Annals of Oncology</i> , 2018, 29, 290-292.	0.6	3
2	Defining a Characteristic Gene Expression Set Responsible for Cancer Stem Cell-Like Features in a Sub-Population of Ewing Sarcoma Cells CADO-ES1. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3908.	1.8	13
3	Inflammatory pathways are upregulated in the nasal epithelium in patients with idiopathic pulmonary fibrosis. <i>Respiratory Research</i> , 2018, 19, 233.	1.4	13
4	Overexpression of TBX3 transcription factor as a potential diagnostic marker for breast cancer. <i>Molecular and Clinical Oncology</i> , 2018, 10, 105-112.	0.4	6
5	Comprehensive analysis of the tumor immune micro-environment in non-small cell lung cancer for efficacy of checkpoint inhibitor. <i>Scientific Reports</i> , 2018, 8, 14576.	1.6	55
6	Hepatocellular Carcinoma Outcome Is Predicted by Expression of Neuronal Calcium Sensor 1. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 1091-1100.	1.1	11
7	Integrin $\hat{\pm}$ 9 Suppresses Hepatocellular Carcinoma Metastasis by Rho GTPase Signaling. <i>Journal of Immunology Research</i> , 2018, 2018, 1-11.	0.9	25
8	Neuropilin-1 expression is associated with lymph node metastasis in breast cancer tissues. <i>Cancer Management and Research</i> , 2018, Volume 10, 1969-1974.	0.9	19
9	Unique Immune Gene Expression Patterns in Bronchoalveolar Lavage and Tumor Adjacent Non-Neoplastic Lung Tissue in Non-Small Cell Lung Cancer. <i>Frontiers in Immunology</i> , 2018, 9, 232.	2.2	10
10	Non-Metastatic Cutaneous Melanoma Induces Chronodisruption in Central and Peripheral Circadian Clocks. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1065.	1.8	33
11	Classification of prostate cancer using a protease activity nanosensor library. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 8954-8959.	3.3	53
12	Intra-individual Gene Expression Variability of Histologically Normal Breast Tissue. <i>Scientific Reports</i> , 2018, 8, 9137.	1.6	5
13	Identification of Coding and Long Noncoding RNAs Differentially Expressed in Tumors and Preferentially Expressed in Healthy Tissues. <i>Cancer Research</i> , 2019, 79, 5167-5180.	0.4	38
14	Alpha-1-acid glycoprotein 1 is upregulated in pancreatic ductal adenocarcinoma and confers a poor prognosis. <i>Translational Research</i> , 2019, 212, 67-79.	2.2	11
15	miRNome Reveals New Insights Into the Molecular Biology of Field Cancerization in Gastric Cancer. <i>Frontiers in Genetics</i> , 2019, 10, 592.	1.1	15
16	Gene expression signature of atypical breast hyperplasia and regulation by SFRP1. <i>Breast Cancer Research</i> , 2019, 21, 76.	2.2	19
17	A cellular and bioinformatics analysis of the SENP1 SUMO isopeptidase in pancreatic cancer. <i>Journal of Gastrointestinal Oncology</i> , 2019, 10, 821-830.	0.6	8
18	The Emerging Role of Cytoskeletal Proteins as Reliable Biomarkers. <i>Proteomics</i> , 2019, 19, e1800483.	1.3	9

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19	Cripto-1 expression in patients with clear cell renal cell carcinoma is associated with poor disease outcome. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019, 38, 378.	3.5	18
20	Gastric Normal Adjacent Mucosa Versus Healthy and Cancer Tissues: Distinctive Transcriptomic Profiles and Biological Features. <i>Cancers</i> , 2019, 11, 1248.	1.7	34
21	Proteomics analysis of the matrisome from MC38 experimental mouse liver metastases. <i>American Journal of Physiology - Renal Physiology</i> , 2019, 317, G625-G639.	1.6	7
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25	Locally advanced rectal cancer transcriptomic-based secretome analysis reveals novel biomarkers useful to identify patients according to neoadjuvant chemoradiotherapy response. <i>Scientific Reports</i> , 2019, 9, 8702.	1.6	14
26	The molecular clock in the skin, its functionality, and how it is disrupted in cutaneous melanoma: a new pharmacological target?. <i>Cellular and Molecular Life Sciences</i> , 2019, 76, 3801-3826.	2.4	28
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38	Alterations of the interactome of Bcl-2 proteins in breast cancer at the transcriptional, mutational and structural level. <i>PLoS Computational Biology</i> , 2019, 15, e1007485.	1.5	42
39	A region-resolved mucosa proteome of the human stomach. <i>Nature Communications</i> , 2019, 10, 39.	5.8	43
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148	Prediction of Biochemical Recurrence-Free Survival of Prostate Cancer Patients Leveraging Multiple Gene Expression Profiles in Tumor Microenvironment. <i>Frontiers in Oncology</i> , 2021, 11, 632571.	1.3	11
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