

CITATION REPORT

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A gene expression-based immune signature for lung adenocarcinoma prognosis

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Cancer Immunology, Immunotherapy, 2020, 69, 1881-1890.

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#	Paper	IF	Citations
13	Tumor Profiling at the Service of Cancer Therapy. <i>Frontiers in Oncology</i> , 2020 , 10, 595613	5.3	2
12	A Novel Early-Stage Lung Adenocarcinoma Prognostic Model Based on Feature Selection With Orthogonal Regression. <i>Frontiers in Cell and Developmental Biology</i> , 2020 , 8, 620746	5.7	
11	Elastic Net Models Based on DNA Copy Number Variations Predicts Clinical Features, Expression Signatures, and Mutations in Lung Adenocarcinoma. <i>Frontiers in Genetics</i> , 2021 , 12, 668040	4.5	
10	The comparison of limma and DESeq2 in gene analysis. <i>E3S Web of Conferences</i> , 2021 , 271, 03058	0.5	1
9	Identification of theranostic factors for patients developing metastasis after surgery for early-stage lung adenocarcinoma. <i>Theranostics</i> , 2021 , 11, 3661-3675	12.1	7
8	Potential Genes Associated with the Survival of Lung Adenocarcinoma Were Identified by Methylation. <i>Computational and Mathematical Methods in Medicine</i> , 2020 , 2020, 7103412	2.8	2
7	Analysis of the tumor immune environment identifies an immune gene set-based prognostic signature in non-small cell lung cancer.. <i>Annals of Translational Medicine</i> , 2022 , 10, 15	3.2	0
6	An advanced non-small cell lung cancer patient with and mutations, and PD-L1 positive, benefited from immunotherapy: a case report.. <i>Annals of Translational Medicine</i> , 2022 , 10, 381	3.2	1
5	Prognostic Value of Genomic Instability of mA-Related lncRNAs in Lung Adenocarcinoma.. <i>Frontiers in Cell and Developmental Biology</i> , 2022 , 10, 707405	5.7	0
4	Human endogenous retrovirus-H long terminal repeat-associating 2: The next immune checkpoint for antitumour therapy.. <i>EBioMedicine</i> , 2022 , 79, 103987	8.8	2
3	Data_Sheet_1.PDF. 2021 ,		
2	Chronic exposure to waterpipe smoke elicits immunomodulatory and carcinogenic effects in the lung.. <i>Cancer Prevention Research</i> , 2022 ,	3.2	
1	Class Discovery, Comparison, and Prediction Methods for RNA-Seq Data. 2022 , 2060-2084		0