

Konstantinos A Kyritsis

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1454060/publications.pdf>

Version: 2024-02-01

11
papers

78
citations

1478505

6
h-index

1474206

9
g-index

12
all docs

12
docs citations

12
times ranked

124
citing authors

#	ARTICLE	IF	CITATIONS
1	InterMineR: an R package for InterMine databases. <i>Bioinformatics</i> , 2019, 35, 3206-3207.	4.1	24
2	The GENetic Syntax Score: a genetic risk assessment implementation tool grading the complexity of coronary artery disease—rationale and design of the GESS study. <i>BMC Cardiovascular Disorders</i> , 2021, 21, 284.	1.7	11
3	A Risk-Stratification Machine Learning Framework for the Prediction of Coronary Artery Disease Severity: Insights From the GESS Trial. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 812182.	2.4	11
4	Dissecting miRNA—Gene Networks to Map Clinical Utility Roads of Pharmacogenomics-Guided Therapeutic Decisions in Cardiovascular Precision Medicine. <i>Cells</i> , 2022, 11, 607.	4.1	11
5	Pharmacogenomic Testing to Guide Personalized Cancer Medicine Decisions in Private Oncology Practice: A Case Study. <i>Frontiers in Oncology</i> , 2020, 10, 521.	2.8	7
6	Uncovering the pharmacological response of novel sesquiterpene derivatives that differentially alter gene expression and modulate the cell cycle in cancer cells. <i>International Journal of Oncology</i> , 2018, 53, 2167-2179.	3.3	6
7	The histone methyltransferase inhibitor A-366 enhances hemoglobin expression in erythroleukemia cells upon co—exposure with chemical inducers in culture. <i>Journal of Biological Research</i> , 2021, 28, 2.	2.1	2
8	Invariable Ribosome Stoichiometry During Murine Erythroid Differentiation: Implications for Understanding Ribosomopathies. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 805541.	3.5	2
9	Analysis of TCGA data of differentially expressed EMT—related genes and miRNAs across various malignancies to identify potential biomarkers. <i>World Academy of Sciences Journal</i> , 2020, 3, .	0.6	1
10	Sequence variation, common tissue expression patterns and learning models: a genome-wide survey of vertebrate ribosomal proteins. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa088.	3.2	1
11	Data on the expression of SRPK1a in mammals. <i>Data in Brief</i> , 2019, 25, 104210.	1.0	0