

Kei Taneishi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1665274/publications.pdf>

Version: 2024-02-01

11
papers

269
citations

1478505

6
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

620
citing authors

#	ARTICLE	IF	CITATIONS
1	GLIDA: GPCR-ligand database for chemical genomic drug discovery. <i>Nucleic Acids Research</i> , 2006, 34, D673-D677.	14.5	70
2	CGBVS&DNN: Prediction of Compound&Protein Interactions Based on Deep Learning. <i>Molecular Informatics</i> , 2017, 36, 1600045.	2.5	58
3	Combination of host immune metabolic biomarkers for the PD-1 blockade cancer immunotherapy. <i>JCI Insight</i> , 2020, 5, .	5.0	58
4	Autoencoder-Based Detection of Dynamic Allostery Triggered by Ligand Binding Based on Molecular Dynamics. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 4043-4051.	5.4	32
5	Adjuvant chemotherapy improves survival of patients with high-risk upper urinary tract urothelial carcinoma: a propensity score-matched analysis. <i>BMC Urology</i> , 2017, 17, 110.	1.4	22
6	Development and validation of a set of six adaptable prognosis prediction (SAP) models based on time-series real-world big data analysis for patients with cancer receiving chemotherapy: A multicenter case crossover study. <i>PLoS ONE</i> , 2017, 12, e0183291.	2.5	19
7	Association between UGT1A1*28*28 genotype and lung cancer in the Japanese population. <i>International Journal of Clinical Oncology</i> , 2017, 22, 269-273.	2.2	4
8	Autoencoder-based detection of the residues involved in G protein-coupled receptor signaling. <i>Scientific Reports</i> , 2021, 11, 19867.	3.3	3
9	Establishment of a terminal prognosis prediction model by applying time series analysis to real-world data. <i>Annals of Oncology</i> , 2015, 26, vii99.	1.2	2
10	On computational complexity of graph inference from counting. <i>Natural Computing</i> , 2013, 12, 589-603.	3.0	1
11	Structure-based analyses of gut microbiome-related proteins by neural networks and molecular dynamics simulations. <i>Current Opinion in Structural Biology</i> , 2022, 73, 102336.	5.7	0