

Kyung In Kim

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

746
citations

1163117

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1125743

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docs citations

13
times ranked

1597
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptional profiling of macrophages in situ in metastatic melanoma reveals localization-dependent phenotypes and function. <i>Cell Reports Medicine</i> , 2022, 3, 100621.	6.5	15
2	Human KIT+ myeloid cells facilitate visceral metastasis by melanoma. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	5
3	Modeling performance of sample collection sites using whole exome sequencing metrics. <i>BioTechniques</i> , 2020, 69, 420-426.	1.8	2
4	A pan-cancer analysis of driver gene mutations, DNA methylation and gene expressions reveals that chromatin remodeling is a major mechanism inducing global changes in cancer epigenomes. <i>BMC Medical Genomics</i> , 2018, 11, 98.	1.5	21
5	IL1 Receptor Antagonist Controls Transcriptional Signature of Inflammation in Patients with Metastatic Breast Cancer. <i>Cancer Research</i> , 2018, 78, 5243-5258.	0.9	119
6	Overfitting, generalization, and MSE in class probability estimation with high-dimensional data. <i>Biometrical Journal</i> , 2014, 56, 256-269.	1.0	6
7	Using single cell sequencing data to model the evolutionary history of a tumor. <i>BMC Bioinformatics</i> , 2014, 15, 27.	2.6	58
8	Probabilistic classifiers with high-dimensional data. <i>Biostatistics</i> , 2011, 12, 399-412.	1.5	27
9	Spatial Clustering of Array CGH Features in Combination with Hierarchical Multiple Testing. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2010, 9, Article40.	0.6	3
10	Effects of dependence in high-dimensional multiple testing problems. <i>BMC Bioinformatics</i> , 2008, 9, 114.	2.6	267
11	SOME COMMENTS ON FALSE DISCOVERY RATE. <i>Journal of Bioinformatics and Computational Biology</i> , 2007, 05, 987-990.	0.8	1
12	CGHcall: calling aberrations for array CGH tumor profiles. <i>Bioinformatics</i> , 2007, 23, 892-894.	4.1	208
13	Estimating the False Discovery Rate Using Nonparametric Deconvolution. <i>Biometrics</i> , 2007, 63, 806-815.	1.4	14