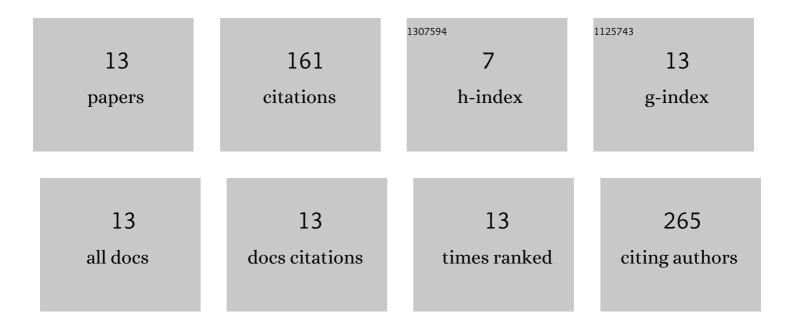
Euna Jeong

List of Publications by Year in descending order

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FUNA LEONG

#	Article	IF	CITATIONS
1	Q-omics: Smart Software for Assisting Oncology and Cancer Research. Molecules and Cells, 2021, 44, 843-850.	2.6	18
2	Analysis of Cross-Association between mRNA Expression and RNAi Efficacy for Predictive Target Discovery in Colon Cancers. Cancers, 2020, 12, 3091.	3.7	12
3	Overexpression and Selective Anticancer Efficacy of in Mutant Lung Cancers. Molecules and Cells, 2019, 42, 804-809.	2.6	23
4	QSurface: fast identification of surface expression markers in cancers. BMC Systems Biology, 2018, 12, 17.	3.0	6
5	β-catenin/TCF activity regulates IGF-1R tyrosine kinase inhibitor sensitivity in colon cancer. Oncogene, 2018, 37, 5466-5475.	5.9	12
6	Patient sample-oriented analysis of gene expression highlights extracellular signatures in breast cancer progression. Biochemical and Biophysical Research Communications, 2017, 487, 307-312.	2.1	6
7	Transcriptome modeling and phenotypic assays for cancer precision medicine. Archives of Pharmacal Research, 2017, 40, 906-914.	6.3	7
8	Lossâ€ofâ€function screens of druggable targetome against cancer stemâ€like cells. FASEB Journal, 2017, 31, 625-635.	0.5	32
9	Cardiac glycosides display selective efficacy for STK11 mutant lung cancer. Scientific Reports, 2016, 6, 29721.	3.3	27
10	Clucose starvation induces mutation and lineage-dependent adaptive responses in a large collection of cancer cell lines. International Journal of Oncology, 2016, 48, 67-72.	3.3	7
11	MACE: mutation-oriented profiling of chemical response and gene expression in cancers. Bioinformatics, 2015, 31, 1508-1514.	4.1	5
12	Analysis of image-based phenotypic parameters for high throughput gene perturbation assays. Computational Biology and Chemistry, 2015, 58, 192-198.	2.3	5
13	Dissecting the global variation of gene expression for the functional interpretation of transcriptome data. Genomics, 2014, 104, 279-286.	2.9	1