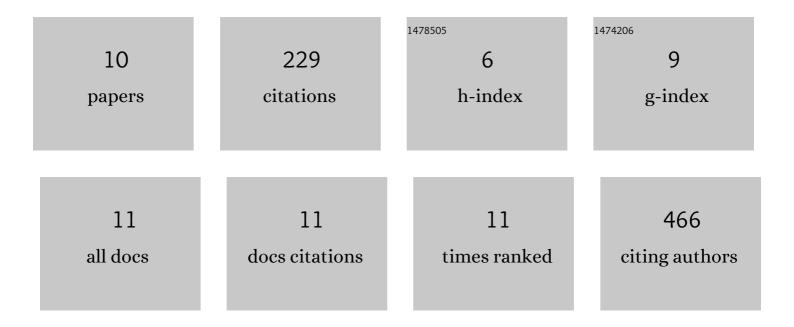
Yeong C Kim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4562421/publications.pdf Version: 2024-02-01



YEONG C KIM

#	Article	IF	CITATIONS
1	Germline variation in <i>BRCA1</i> / <i>2</i> is highly ethnicâ€specific: Evidence from over 30,000 Chinese hereditary breast and ovarian cancer patients. International Journal of Cancer, 2019, 145, 962-973.	5.1	61
2	Zika virus propagation and release in human fetal astrocytes can be suppressed by neutral sphingomyelinase-2 inhibitor GW4869. Cell Discovery, 2018, 4, 19.	6.7	59
3	Prevalence and spectrum of <i>BRCA</i> germline variants in mainland Chinese familial breast and ovarian cancer patients. Oncotarget, 2016, 7, 9600-9612.	1.8	52
4	Ccdc3: A New P63 Target Involved in Regulation Of Liver Lipid Metabolism. Scientific Reports, 2017, 7, 9020.	3.3	18
5	Unique Features of Germline Variation in Five Egyptian Familial Breast Cancer Families Revealed by Exome Sequencing. PLoS ONE, 2017, 12, e0167581.	2.5	17
6	Differences of Variable Number Tandem Repeats in XRCC5 Promoter Are Associated with Increased or Decreased Risk of Breast Cancer in BRCA Gene Mutation Carriers. Frontiers in Oncology, 2016, 6, 92.	2.8	10
7	Erk5 contributes to maintaining the balance of cellular nucleotide levels and erythropoiesis. Cell Cycle, 2015, 14, 3864-3876.	2.6	8
8	Evidences showing wide presence of small genomic aberrations in chronic lymphocytic leukemia. BMC Research Notes, 2010, 3, 341.	1.4	3
9	Single-base LOH can be used as Specific Marker to Classify <i>BRCAx</i> Familial Breast Cancer into More Homogenous Subtypes. Breast Journal, 2017, 23, 479-481.	1.0	1
10	Identification of new susceptibility genes in familial breast cancer by exome sequencing Journal of Clinical Oncology, 2012, 30, 1538-1538.	1.6	0