

Pik-Fang Kho

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

404
citations

933447

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888059

17
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24
all docs

24
docs citations

24
times ranked

903
citing authors

#	ARTICLE	IF	CITATIONS
1	Assessing the Genetic Predisposition of Education on Myopia: A Mendelian Randomization Study. <i>Genetic Epidemiology</i> , 2016, 40, 66-72.	1.3	56
2	Genetically low vitamin D concentrations and myopic refractive error: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2017, 46, 1882-1890.	1.9	47
3	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. <i>International Journal of Cancer</i> , 2021, 148, 307-319.	5.1	35
4	Genome-Wide Association Studies of Endometrial Cancer: Latest Developments and Future Directions. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1095-1102.	2.5	32
5	A multi-level investigation of the genetic relationship between endometriosis and ovarian cancer histotypes. <i>Cell Reports Medicine</i> , 2022, 3, 100542.	6.5	26
6	Assessment and visualization of phenome-wide causal relationships using genetic data: an application to dental caries and periodontitis. <i>European Journal of Human Genetics</i> , 2021, 29, 300-308.	2.8	23
7	Nonsteroidal anti-inflammatory drugs, statins, and pancreatic cancer risk: a population-based case-control study. <i>Cancer Causes and Control</i> , 2016, 27, 1457-1464.	1.8	18
8	Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. <i>Human Genetics</i> , 2021, 140, 1353-1365.	3.8	18
9	Assessing the Role of Selenium in Endometrial Cancer Risk: A Mendelian Randomization Study. <i>Frontiers in Oncology</i> , 2019, 9, 182.	2.8	15
10	p53 major hotspot variants are associated with poorer prognostic features in hereditary cancer patients. <i>Cancer Genetics</i> , 2019, 235-236, 21-27.	0.4	11
11	Phenome-wide screening of GWAS data reveals the complex causal architecture of obesity. <i>Human Genetics</i> , 2021, 140, 1253-1265.	3.8	11
12	Multi-tissue transcriptome-wide association study identifies eight candidate genes and tissue-specific gene expression underlying endometrial cancer susceptibility. <i>Communications Biology</i> , 2021, 4, 1211.	4.4	11
13	Genetic basis to structural grey matter associations with chronic pain. <i>Brain</i> , 2021, 144, 3611-3622.	7.6	10
14	Associations between Genetically Predicted Circulating Protein Concentrations and Endometrial Cancer Risk. <i>Cancers</i> , 2021, 13, 2088.	3.7	10
15	Evidence of Genetic Overlap Between Circadian Preference and Brain White Matter Microstructure. <i>Twin Research and Human Genetics</i> , 2021, 24, 1-6.	0.6	2
16	Expression QTL analysis of glaucoma endophenotypes in the Norfolk Island isolate provides evidence that immune-related genes are associated with optic disc size. <i>Journal of Human Genetics</i> , 2018, 63, 83-87.	2.3	1
17	Genome-wide linkage and association analysis of primary open-angle glaucoma endophenotypes in the Norfolk Island isolate. <i>Molecular Vision</i> , 2017, 23, 660-665.	1.1	1
18	Abstract PR003: Multi-tissue transcriptome-wide association study identifies genetic mechanisms underlying endometrial cancer susceptibility. , 2021, , .		0