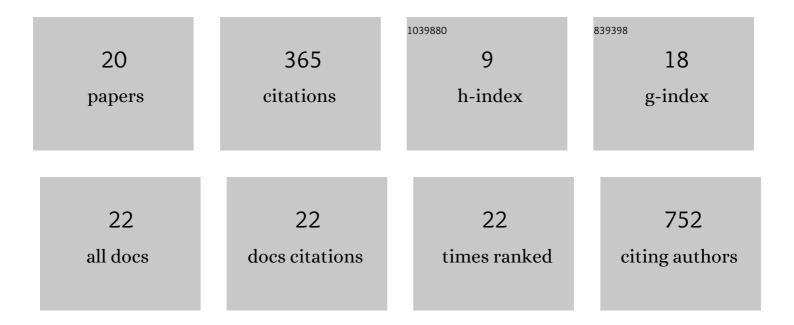
Sock Hoai Chan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6703364/publications.pdf

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SOCK HOAL CHAN

#	Article	IF	CITATIONS
1	Tissue Microbiome Profiling Identifies an Enrichment of Specific Enteric Bacteria in Opisthorchis viverrini Associated Cholangiocarcinoma. EBioMedicine, 2016, 8, 195-202.	2.7	94
2	Germline Mutations in Cancer Predisposition Genes are Frequent in Sporadic Sarcomas. Scientific Reports, 2017, 7, 10660.	1.6	52
3	CDKN2A germline alterations and the relevance of genotype-phenotype associations in cancer predisposition. Hereditary Cancer in Clinical Practice, 2021, 19, 21.	0.6	36
4	Mutation spectrum of POLE and POLD1 mutations in South East Asian women presenting with grade 3 endometrioid endometrial carcinomas. Gynecologic Oncology, 2016, 141, 113-120.	0.6	34
5	Impact of subsidies on cancer genetic testing uptake in Singapore. Journal of Medical Genetics, 2017, 54, 254-259.	1.5	24
6	Germline Pathogenic Variants in Homologous Recombination and DNA Repair Genes in an Asian Cohort of Young-Onset Colorectal Cancer. JNCI Cancer Spectrum, 2018, 2, pky054.	1.4	21
7	Family history assessment significantly enhances delivery of precision medicine in the genomics era. Genome Medicine, 2021, 13, 3.	3.6	19
8	Impact of Variant Reclassification in Cancer Predisposition Genes on Clinical Care. JCO Precision Oncology, 2021, 5, 577-584.	1.5	18
9	Germline mutation contribution to chromosomal instability. Endocrine-Related Cancer, 2017, 24, T33-T46.	1.6	13
10	Clinical relevance of screening checklists for detecting cancer predisposition syndromes in Asian childhood tumours. Npj Genomic Medicine, 2018, 3, 30.	1.7	13
11	Germline hemizygous deletion of CDKN2A–CDKN2B locus in a patient presenting with Li–Fraumeni syndrome. Npj Genomic Medicine, 2016, 1, 16015.	1.7	9
12	Functional analysis of clinical BARD1 germline variants. Journal of Physical Education and Sports Management, 2019, 5, a004093.	0.5	6
13	Whole exome sequencing identifies recessive germline mutations in FAM160A1 in familial NK/T cell lymphoma. Blood Cancer Journal, 2018, 8, 111.	2.8	5
14	Investigation into the origins of an ancient BRCA1 founder mutation identified among Chinese families in Singapore. International Journal of Cancer, 2021, 148, 637-645.	2.3	5
15	Multiple neoplasia in a patient with Gitelman syndrome harboring germline monoallelic MUTYH mutation. Npj Genomic Medicine, 2020, 5, 39.	1.7	3
16	Functional characterisation guides classification of novel BAP1 germline variants. Npj Genomic Medicine, 2020, 5, 50.	1.7	3
17	Biallelic NF1 inactivation in high grade serous ovarian cancers from patients with neurofibromatosis type 1. Familial Cancer, 2020, 19, 353-358.	0.9	3
18	Missense PALB2 germline variant disrupts nuclear localization of PALB2 in a patient with breast cancer. Familial Cancer, 2020, 19, 123-131.	0.9	3

# AK	TICLE	IF	CITATIONS
19 Spe Var	ectrum of Germline Mutations Within Fanconi Anemia–Associated Genes Across Populations of rying Ancestry. JNCI Cancer Spectrum, 2021, 5, .	1.4	3

Advances in Sarcoma Genomics and Therapeutic Management. , 2019, , 609-621.

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