# Sock Hoai Chan

List of Publications by Year
in descending order

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

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|---|---|---|---|
| 20<br>papers | 365<br>citations | 1039880<br>9<br>h-index | 839398<br>18<br>g-index |
| 22<br>all docs | 22<br>docs citations | 22<br>times ranked | 752<br>citing authors |

| # | 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 CDKN2Aâ€“CDKN2B locus in a patient presenting with Liâ€“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 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Spectrum of Germline Mutations Within Fanconi Anemiaâ€“Associated Genes Across Populations of Varying Ancestry. JNCI Cancer Spectrum, 2021, 5, . | 1.4 | 3 |
| 20 | Advances in Sarcoma Genomics and Therapeutic Management. , 2019, , 609-621. | | 0 |