Jessada Thutkawkorapin

List of Publications by Year
in descending order

17 papers	298 citations	1478505 6 h-index	940533 16 g-index
17 all docs	17 docs citations	17 times ranked	844 citing authors

#	Article	IF	Citations
1	SweGen: a whole-genome data resource of genetic variability in a cross-section of the Swedish population. European Journal of Human Genetics, 2017, 25, 1253-1260.	2.8	148
2	Wholeâ€genome sequencing of synchronous thyroid carcinomas identifies aberrant DNA repair in thyroid cancer dedifferentiation. Journal of Pathology, 2020, 250, 183-194.	4.5	40
3	Exome sequencing in one family with gastric- and rectal cancer. BMC Genetics, 2016, 17, 41.	2.7	31
4	Exome sequencing in 51 early onset nonâ€familial CRC cases. Molecular Genetics & Genomic Medicine, 2019, 7, e605.	1.2	17
5	Frequent GU wobble pairings reduce translation efficiency in Plasmodium falciparum. Scientific Reports, 2017, 7, 723.	3.3	14
6	PHIP- a novel candidate breast cancer susceptibility locus on 6q14.1. Oncotarget, 2017, 8, 102769-102782.	1.8	9
7	pyCancerSig: subclassifying human cancer with comprehensive single nucleotide, structural and microsatellite mutational signature deconstruction from whole genome sequencing. BMC Bioinformatics, 2020, 21, 128.	2.6	7
8	Cancer risk susceptibility loci in a Swedish population. Oncotarget, 2017, 8, 110300-110310.	1.8	7
9	Genetic analyses supporting colorectal, gastric, and prostate cancer syndromes. Genes Chromosomes and Cancer, 2019, 58, 775-782.	2.8	6
10	Linkage analysis revealed risk loci on 6p21 and 18p11.2-q11.2 in familial colon and rectal cancer, respectively. European Journal of Human Genetics, 2019, 27, 1286-1295.	2.8	4
11	Two novel colorectal cancer risk loci in the region on chromosome 9q22.32. Oncotarget, 2018, 9, 11170-11179.	1.8	4
12	Massive parallel sequencing in a family with rectal cancer. Hereditary Cancer in Clinical Practice, 2021, 19, 23.	1.5	3
13	A search for modifying genetic factors in CHEK2:c.1100delC breast cancer patients. Scientific Reports, 2021, 11, 14763.	3.3	3
14	Sequencing for germline mutations in Swedish breast cancer families reveals novel breast cancer risk genes. Scientific Reports, 2021, 11, 14737.	3.3	2
15	Colorectal cancer risk susceptibility loci in a Swedish population. Molecular Carcinogenesis, 2021, , .	2.7	2
16	A Swedish Genome-Wide Haplotype Association Analysis Identifies a Novel Breast Cancer Susceptibility Locus in 8p21.2 and Characterizes Three Loci on Chromosomes 10, 11 and 16. Cancers, 2022, 14, 1206.	3.7	1
17	Identification of known and novel familial cancer genes in Swedish colorectal cancer families. International Journal of Cancer, 2021, 149, 627-634.	5.1	0