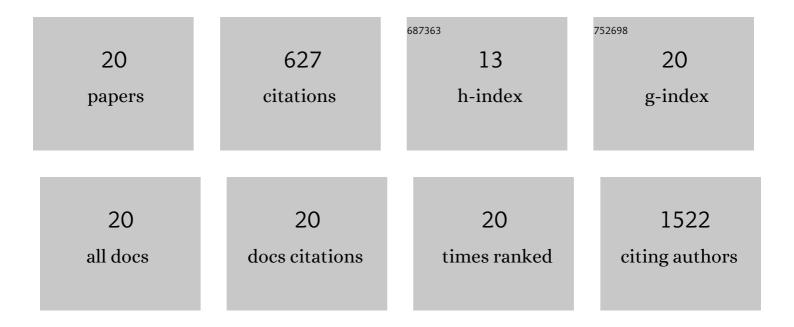
Kanthida Kusonmano

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The genomic landscape and evolution of endometrial carcinoma progression and abdominopelvic metastasis. Nature Genetics, 2016, 48, 848-855.	21.4	174
2	Loss of progesterone receptor links to high proliferation and increases from primary to metastatic endometrial cancer lesions. European Journal of Cancer, 2014, 50, 3003-3010.	2.8	73
3	Molecular profiling of endometrial carcinoma precursor, primary and metastatic lesions suggests different targets for treatment in obese compared to non-obese patients. Oncotarget, 2015, 6, 1327-1339.	1.8	50
4	HER2 expression patterns in paired primary and metastatic endometrial cancer lesions. British Journal of Cancer, 2018, 118, 378-387.	6.4	43
5	Clinicopathologic and molecular markers in cervical carcinoma: a prospective cohort study. American Journal of Obstetrics and Gynecology, 2017, 217, 432.e1-432.e17.	1.3	38
6	ATAD2 overexpression links to enrichment of B-MYB-translational signatures and development of aggressive endometrial carcinoma. Oncotarget, 2015, 6, 28440-28452.	1.8	37
7	High Phospho-Stathmin(Serine38) Expression Identifies Aggressive Endometrial Cancer and Suggests an Association with PI3K Inhibition. Clinical Cancer Research, 2013, 19, 2331-2341.	7.0	35
8	Informatics for Metabolomics. Advances in Experimental Medicine and Biology, 2016, 939, 91-115.	1.6	28
9	Switch in FOXA1 Status Associates with Endometrial Cancer Progression. PLoS ONE, 2014, 9, e98069.	2.5	22
10	Diverse Microbial Community Profiles of Propionate-Degrading Cultures Derived from Different Sludge Sources of Anaerobic Wastewater Treatment Plants. Microorganisms, 2020, 8, 277.	3.6	21
11	PIK3CA exon9 mutations associate with reduced survival, and are highly concordant between matching primary tumors and metastases in endometrial cancer. Scientific Reports, 2017, 7, 10240.	3.3	19
12	<i>PIK3CA</i> Amplification Associates with Aggressive Phenotype but Not Markers of AKT-MTOR Signaling in Endometrial Carcinoma. Clinical Cancer Research, 2019, 25, 334-345.	7.0	17
13	Increased angiogenesis is associated with a 32-gene expression signature and 6p21 amplification in aggressive endometrial cancer. Oncotarget, 2015, 6, 10634-10645.	1.8	15
14	Microbial community structure in aquifers associated with arsenic: analysis of 16S rRNA and arsenite oxidase genes. PeerJ, 2021, 9, e10653.	2.0	12
15	Identification of highly connected and differentially expressed gene subnetworks in metastasizing endometrial cancer. PLoS ONE, 2018, 13, e0206665.	2.5	11
16	Dip in the gene pool: Metagenomic survey of natural coccolithovirus communities. Virology, 2014, 466-467, 129-137.	2.4	10
17	Genetic Aberration Analysis in Thai Colorectal Adenoma and Early-Stage Adenocarcinoma Patients by Whole-Exome Sequencing. Cancers, 2019, 11, 977.	3.7	10
18	Arsenic speciation, the abundance of arsenite-oxidizing bacteria and microbial community structures in groundwater, surface water, and soil from a gold mine. Journal of Environmental Science and Health - Part A Toxic/Hazardous Substances and Environmental Engineering, 2021, 56, 769-785.	1.7	6

#	Article	IF	CITATIONS
19	Aneuploidy related transcriptional changes in endometrial cancer link low expression of chromosome 15q genes to poor survival. Oncotarget, 2017, 8, 9696-9707.	1.8	4
20	Gene Expression Analysis Through Network Biology: Bioinformatics Approaches. Advances in Biochemical Engineering/Biotechnology, 2016, 160, 15-32.	1.1	2