Susan A Kadlubar

List of Publications by Year
in descending order

28	433	758635 12	752256 20
papers	citations	h-index	g-index

28	28	28	1004
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Association of DNA-Methylation Profiles With Immune Responses Elicited in Breast Cancer Patients Immunized With a Carbohydrate-Mimicking Peptide: A Pilot Study. Frontiers in Oncology, 2020, 10, 879.	1.3	1
2	Genome-wide DNA methylation signatures to predict pathologic complete response from combined neoadjuvant chemotherapy with bevacizumab in breast cancer. PLoS ONE, 2020, 15, e0230248.	1.1	13
3	Mobile Mammography Screening as an Opportunity to Increase Access of Rural Women to Breast Cancer Research Studies. Breast Cancer: Basic and Clinical Research, 2019, 13, 117822341987629.	0.6	9
4	Development and Evaluation of a Blood Glucose Monitoring YouTube Video for Marshallese Patients Using a Community-Based Participatory Research Approach. Policy, Politics, and Nursing Practice, 2019, 20, 205-215.	0.8	7
5	Germline Genetic Variants in GATA3 and Breast Cancer Treatment Outcomes in SWOG S8897 Trial and the Pathways Study. Clinical Breast Cancer, 2019, 19, 225-235.e2.	1.1	4
6	County poverty levels influence genome-wide DNA methylation profiles in African American and European American women. Translational Cancer Research, 2019, 8, 683-692.	0.4	5
7	Individual- and county-level determinants of high breast cancer incidence rates. Translational Cancer Research, 2019, 8, S323-S333.	0.4	4
8	Challenges in recruiting African-American women for a breast cancer genetics study. Hereditary Cancer in Clinical Practice, 2018, 16, 8.	0.6	10
9	A high frequency missense SULT1B1 allelic variant (L145V) selectively expressed in African descendants exhibits altered kinetic properties. Xenobiotica, 2018, 48, 79-88.	0.5	5
10	A functional SNP in the 3â€²â€²UTR of TAP2 gene interacts with microRNA hsaâ€miRâ€1270 to suppress the gene expression. Environmental and Molecular Mutagenesis, 2018, 59, 134-143.	0.9	32
11	Validation of a genetic risk score for Arkansas women of color. PLoS ONE, 2018, 13, e0204834.	1.1	12
12	Leveraging community-based participatory research capacity to recruit Pacific Islanders into a genetics study. Journal of Community Genetics, 2017, 8, 283-291.	0.5	25
13	Change in Mammography Use Following the Revised Guidelines from the U.S. Preventive Services Task Force. Breast Journal, 2017, 23, 164-168.	0.4	10
14	Germline Genetic Variants in TEK, ANGPT1, ANGPT2, MMP9, FGF2 and VEGFA Are Associated with Pathologic Complete Response to Bevacizumab in Breast Cancer Patients. PLoS ONE, 2017, 12, e0168550.	1.1	20
15	2-amino-1-methyl-6-phenylimidazo(4,5-b) pyridine (PhIP) induces gene expression changes in JAK/STAT and MAPK pathways related to inflammation, diabetes and cancer. Nutrition and Metabolism, 2016, 13, 54.	1.3	17
16	Modulation of ALDH5A1 and SLC22A7 by microRNA hsa-miR-29a-3p in human liver cells. Biochemical Pharmacology, 2015, 98, 671-680.	2.0	21
17	Adipocyte hypoxia promotes epithelial-mesenchymal transition-related gene expression and estrogen receptor-negative phenotype in breast cancer cells. Oncology Reports, 2015, 33, 2689-2694.	1.2	40
18	Suppression of CYP2C9 by MicroRNA hsa-miR-128-3p in Human Liver Cells and Association with Hepatocellular Carcinoma. Scientific Reports, 2015, 5, 8534.	1.6	92

#	Article	IF	Citations
19	Genotypic and Allelic Variability in CYP19A1 among Populations of African and European Ancestry. PLoS ONE, 2015, 10, e0117347.	1.1	3
20	Effect of MRP2 and MRP3 Polymorphisms on Anastrozole Glucuronidation and MRP2 and MRP3 Gene Expression in Normal Liver Samples. International Journal of Cancer Research and Molecular Mechanisms, 2015, 1, .	0.2	6
21	CYP19A1 single nucleotide polymorphism associations with CYP19A1, NFκB1, and IL6 gene expression in human normal colon and normal liver samples. Pharmacogenomics and Personalized Medicine, 2014, 7, 163.	0.4	2
22	A Potential Role for Human UDP-Glucuronosyltransferase 1A4 Promoter Single Nucleotide Polymorphisms in the Pharmacogenomics of Tamoxifen and Its Derivatives. Drug Metabolism and Disposition, 2014, 42, 1392-1400.	1.7	11
23	Sulfotransferase 1A1 (SULT1A1) gene expression is regulated by members of the NFI transcription factors in human breast cancer cells. BMC Clinical Pathology, 2014, 14, 1.	1.8	23
24	Polymorphisms of ERBB2 and breast cancer risk: A meta-analysis of 26 studies involving 35,088 subjects. Journal of Surgical Oncology, 2013, 108, 335-336.	0.8	1
25	Potential Role ofUGT1A4Promoter SNPs in Anastrozole Pharmacogenomics. Drug Metabolism and Disposition, 2013, 41, 870-877.	1.7	25
26	Sulfation of 4-Hydroxy Toremifene: Individual Variability, Isoform Specificity, and Contribution to Toremifene Pharmacogenomics. Drug Metabolism and Disposition, 2012, 40, 1210-1215.	1.7	17
27	Effect of Sulfotransferase Structural Rearrangements in Fulvestrant Sulfation. FASEB Journal, 2012, 26, 850.11.	0.2	0
28	Phenotypic CYP2A6 variation and the risk of pancreatic cancer. JOP: Journal of the Pancreas, 2009, 10, 263-70.	1.5	18