

Susan A Kadlubar

List of Publications by Year in descending order

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28
papers

433
citations

758635

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752256

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28
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28
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28
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1004
citing authors

#	ARTICLE	IF	CITATIONS
1	Suppression of CYP2C9 by MicroRNA hsa-miR-128-3p in Human Liver Cells and Association with Hepatocellular Carcinoma. <i>Scientific Reports</i> , 2015, 5, 8534.	1.6	92
2	Adipocyte hypoxia promotes epithelial-mesenchymal transition-related gene expression and estrogen receptor-negative phenotype in breast cancer cells. <i>Oncology Reports</i> , 2015, 33, 2689-2694.	1.2	40
3	A functional SNP in the 3'UTR of TAP2 gene interacts with microRNA hsa-miR-1270 to suppress the gene expression. <i>Environmental and Molecular Mutagenesis</i> , 2018, 59, 134-143.	0.9	32
4	Potential Role of UGT1A4 Promoter SNPs in Anastrozole Pharmacogenomics. <i>Drug Metabolism and Disposition</i> , 2013, 41, 870-877.	1.7	25
5	Leveraging community-based participatory research capacity to recruit Pacific Islanders into a genetics study. <i>Journal of Community Genetics</i> , 2017, 8, 283-291.	0.5	25
6	Sulfotransferase 1A1 (SULT1A1) gene expression is regulated by members of the NFI transcription factors in human breast cancer cells. <i>BMC Clinical Pathology</i> , 2014, 14, 1.	1.8	23
7	Modulation of ALDH5A1 and SLC22A7 by microRNA hsa-miR-29a-3p in human liver cells. <i>Biochemical Pharmacology</i> , 2015, 98, 671-680.	2.0	21
8	Germline Genetic Variants in TEK, ANGPT1, ANGPT2, MMP9, FGF2 and VEGFA Are Associated with Pathologic Complete Response to Bevacizumab in Breast Cancer Patients. <i>PLoS ONE</i> , 2017, 12, e0168550.	1.1	20
9	Phenotypic CYP2A6 variation and the risk of pancreatic cancer. <i>JOP: Journal of the Pancreas</i> , 2009, 10, 263-70.	1.5	18
10	Sulfation of 4-Hydroxy Toremifene: Individual Variability, Isoform Specificity, and Contribution to Toremifene Pharmacogenomics. <i>Drug Metabolism and Disposition</i> , 2012, 40, 1210-1215.	1.7	17
11	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. <i>Nutrition and Metabolism</i> , 2016, 13, 54.	1.3	17
12	Genome-wide DNA methylation signatures to predict pathologic complete response from combined neoadjuvant chemotherapy with bevacizumab in breast cancer. <i>PLoS ONE</i> , 2020, 15, e0230248.	1.1	13
13	Validation of a genetic risk score for Arkansas women of color. <i>PLoS ONE</i> , 2018, 13, e0204834.	1.1	12
14	A Potential Role for Human UDP-Glucuronosyltransferase 1A4 Promoter Single Nucleotide Polymorphisms in the Pharmacogenomics of Tamoxifen and Its Derivatives. <i>Drug Metabolism and Disposition</i> , 2014, 42, 1392-1400.	1.7	11
15	Change in Mammography Use Following the Revised Guidelines from the U.S. Preventive Services Task Force. <i>Breast Journal</i> , 2017, 23, 164-168.	0.4	10
16	Challenges in recruiting African-American women for a breast cancer genetics study. <i>Hereditary Cancer in Clinical Practice</i> , 2018, 16, 8.	0.6	10
17	Mobile Mammography Screening as an Opportunity to Increase Access of Rural Women to Breast Cancer Research Studies. <i>Breast Cancer: Basic and Clinical Research</i> , 2019, 13, 117822341987629.	0.6	9
18	Development and Evaluation of a Blood Glucose Monitoring YouTube Video for Marshallese Patients Using a Community-Based Participatory Research Approach. <i>Policy, Politics, and Nursing Practice</i> , 2019, 20, 205-215.	0.8	7

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19	Effect of MRP2 and MRP3 Polymorphisms on Anastrozole Glucuronidation and MRP2 and MRP3 Gene Expression in Normal Liver Samples. <i>International Journal of Cancer Research and Molecular Mechanisms</i> , 2015, 1, .	0.2	6
20	A high frequency missense SULT1B1 allelic variant (L145V) selectively expressed in African descendants exhibits altered kinetic properties. <i>Xenobiotica</i> , 2018, 48, 79-88.	0.5	5
21	County poverty levels influence genome-wide DNA methylation profiles in African American and European American women. <i>Translational Cancer Research</i> , 2019, 8, 683-692.	0.4	5
22	Germline Genetic Variants in GATA3 and Breast Cancer Treatment Outcomes in SWOG S8897 Trial and the Pathways Study. <i>Clinical Breast Cancer</i> , 2019, 19, 225-235.e2.	1.1	4
23	Individual- and county-level determinants of high breast cancer incidence rates. <i>Translational Cancer Research</i> , 2019, 8, S323-S333.	0.4	4
24	Genotypic and Allelic Variability in CYP19A1 among Populations of African and European Ancestry. <i>PLoS ONE</i> , 2015, 10, e0117347.	1.1	3
25	CYP19A1 single nucleotide polymorphism associations with CYP19A1, NF κ B1, and IL6 gene expression in human normal colon and normal liver samples. <i>Pharmacogenomics and Personalized Medicine</i> , 2014, 7, 163.	0.4	2
26	Polymorphisms of ERBB2 and breast cancer risk: A meta-analysis of 26 studies involving 35,088 subjects. <i>Journal of Surgical Oncology</i> , 2013, 108, 335-336.	0.8	1
27	Association of DNA-Methylation Profiles With Immune Responses Elicited in Breast Cancer Patients Immunized With a Carbohydrate-Mimicking Peptide: A Pilot Study. <i>Frontiers in Oncology</i> , 2020, 10, 879.	1.3	1
28	Effect of Sulfotransferase Structural Rearrangements in Fulvestrant Sulfation. <i>FASEB Journal</i> , 2012, 26, 850.11.	0.2	0