

# Tammy Moser Havener

## List of Publications by Year in descending order

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Version: 2024-02-01

19  
papers

310  
citations

933447

10  
h-index

888059

17  
g-index

21  
all docs

21  
docs citations

21  
times ranked

669  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pharmacogenomic characterization of US FDA-approved cytotoxic drugs. <i>Pharmacogenomics</i> , 2011, 12, 1407-1415.	1.3	44
2	Design of a Cyclin G Associated Kinase (GAK)/Epidermal Growth Factor Receptor (EGFR) Inhibitor Set to Interrogate the Relationship of EGFR and GAK in Chordoma. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 4772-4778.	6.4	34
3	A genome-wide association analysis of temozolomide response using lymphoblastoid cell lines shows a clinically relevant association with MGMT. <i>Pharmacogenetics and Genomics</i> , 2012, 22, 796-802.	1.5	32
4	Genetic Variants in <i>CPA6</i> and <i>PRPF31</i> Are Associated With Variation in Response to Metformin in Individuals With Type 2 Diabetes. <i>Diabetes</i> , 2018, 67, 1428-1440.	0.6	32
5	Genetic Variants in <i>HSD17B3</i> , <i>SMAD3</i> , and <i>IPO11</i> Impact Circulating Lipids in Response to Fenofibrate in Individuals With Type 2 Diabetes. <i>Clinical Pharmacology and Therapeutics</i> , 2018, 103, 712-721.	4.7	30
6	Genome-wide association and pharmacological profiling of 29 anticancer agents using lymphoblastoid cell lines. <i>Pharmacogenomics</i> , 2014, 15, 137-146.	1.3	27
7	Synergistic drug combinations and machine learning for drug repurposing in chordoma. <i>Scientific Reports</i> , 2020, 10, 12982.	3.3	27
8	Multivariate methods and software for association mapping in dose-response genome-wide association studies. <i>BioData Mining</i> , 2012, 5, 21.	4.0	17
9	A comparison of association methods for cytotoxicity mapping in pharmacogenomics. <i>Frontiers in Genetics</i> , 2011, 2, 86.	2.3	14
10	Identification and Replication of Loci Involved in Camptothecin-Induced Cytotoxicity Using CEPH Pedigrees. <i>PLoS ONE</i> , 2011, 6, e17561.	2.5	14
11	Common and rare genetic markers of lipid variation in subjects with type 2 diabetes from the ACCORD clinical trial. <i>PeerJ</i> , 2017, 5, e3187.	2.0	11
12	Evaluating the role of admixture in cancer therapy via <i>in vitro</i> drug response and multivariate genome-wide associations. <i>Pharmacogenomics</i> , 2015, 16, 1451-1463.	1.3	8
13	High-throughput screening and genome-wide analyses of 44 anticancer drugs in the 1000 Genomes cell lines reveals an association of the <i>NQO1</i> gene with the response of multiple anticancer drugs. <i>PLoS Genetics</i> , 2021, 17, e1009732.	3.5	6
14	Synergistic Chemotherapy Drug Response Is a Genetic Trait in Lymphoblastoid Cell Lines. <i>Frontiers in Genetics</i> , 2019, 10, 829.	2.3	5
15	Incorporating Concomitant Medications into Genome-Wide Analyses for the Study of Complex Disease and Drug Response. <i>Frontiers in Genetics</i> , 2016, 7, 138.	2.3	2
16	Institutional Profile: UNC Institute for Pharmacogenomics and Individualized Therapy: interdisciplinary research for individual care. <i>Pharmacogenomics</i> , 2010, 11, 13-21.	1.3	1
17	Abstract LB-246: Evaluating the role of admixture in cancer therapy via <i>in vitro</i> drug response and multivariate genome-wide associations. , 2015, , .		1
18	The influence of Neanderthal alleles on cytotoxic response. <i>PeerJ</i> , 2018, 6, e5691.	2.0	1

#	ARTICLE	IF	CITATIONS
19	Race and smoking status associated with paclitaxel drug response in patient-derived lymphoblastoid cell lines. <i>Pharmacogenetics and Genomics</i> , 2021, 31, 48-52.	1.5	0