## Mahlet G Tadesse

List of Publications by Year in descending order

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45 papers 1,522 citations

471509 17 h-index 330143 37 g-index

47 all docs

47 docs citations

47 times ranked

2325 citing authors

#	Article	IF	CITATIONS
1	Bayesian statistics and modelling. Nature Reviews Methods Primers, 2021, 1, .	21.2	419
2	Bayesian Variable Selection in Clustering High-Dimensional Data. Journal of the American Statistical Association, 2005, 100, 602-617.	3.1	190
3	Incorporating biological information into linear models: A Bayesian approach to the selection of pathways and genes. Annals of Applied Statistics, 2011, 5, 1978-2002.	1.1	119
4	Circulating early- and mid-pregnancy microRNAs and risk of gestational diabetes. Diabetes Research and Clinical Practice, $2017, 132, 1-9$ .	2.8	89
5	Construct Validity and Factor Structure of the Pittsburgh Sleep Quality Index and Epworth Sleepiness Scale in a Multi-National Study of African, South East Asian and South American College Students. PLoS ONE, 2014, 9, e116383.	2.5	69
6	LCâ€MS/MSâ€based serum proteomics for identification of candidate biomarkers for hepatocellular carcinoma. Proteomics, 2015, 15, 2369-2381.	2.2	66
7	GC-MS Based Plasma Metabolomics for Identification of Candidate Biomarkers for Hepatocellular Carcinoma in Egyptian Cohort. PLoS ONE, 2015, 10, e0127299.	2.5	60
8	Pathway and Network Approaches for Identification of Cancer Signature Markers from Omics Data. Journal of Cancer, 2015, 6, 54-65.	2.5	44
9	Psychometric properties and factor structure of the General Health Questionnaire as a screening tool for anxiety and depressive symptoms in a multi-national study of young adults. Journal of Affective Disorders, 2015, 187, 197-202.	4.1	43
10	Metabolomic Characterization of Hepatocellular Carcinoma in Patients with Liver Cirrhosis for Biomarker Discovery. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 675-683.	2.5	40
11	Biological network inference using low order partial correlation. Methods, 2014, 69, 266-273.	3.8	35
12	Placental Genome and Maternal-Placental Genetic Interactions: A Genome-Wide and Candidate Gene Association Study of Placental Abruption. PLoS ONE, 2014, 9, e116346.	2.5	29
13	Assessing validity of a depression screening instrument in the absence of a gold standard. Annals of Epidemiology, 2014, 24, 527-531.	1.9	28
14	Candidate Gene and MicroRNA Expression in Fetal Membranes and Preterm Delivery Risk. Reproductive Sciences, 2016, 23, 731-737.	2.5	28
15	Early Pregnancy Maternal Blood DNA Methylation in Repeat Pregnancies and Change in Gestational Diabetes Mellitus Status—A Pilot Study. Reproductive Sciences, 2015, 22, 904-910.	2.5	25
16	Daytime Sleepiness, Circadian Preference, Caffeine Consumption and Khat Use among College Students in Ethiopia. Journal of Sleep Disorders Treatment & Care, 2014, 03, .	0.1	22
17	Daytime sleepiness, circadian preference, caffeine consumption and use of other stimulants among Thai college students. Journal of Public Health and Epidemiology, 2014, 8, 202-210.	0.3	21
18	Identification of race-associated metabolite biomarkers for hepatocellular carcinoma in patients with liver cirrhosis and hepatitis C virus infection. PLoS ONE, 2018, 13, e0192748.	2.5	19

#	Article	IF	CITATIONS
19	Adherence to Adjuvant Endocrine Therapy in Insured Black and White Breast Cancer Survivors: Exploring Adherence Measures in Patient Data. Journal of Managed Care & Decialty Pharmacy, 2019, 25, 578-586.	0.9	16
20	Race and Patient-reported Symptoms in Adherence to Adjuvant Endocrine Therapy: A Report from the Women's Hormonal Initiation and Persistence Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 699-709.	2.5	16
21	Maternal pre-pregnancy body mass index and circulating microRNAs in pregnancy. Obesity Research and Clinical Practice, 2017, 11, 464-474.	1.8	15
22	Genetic variations related to maternal whole blood mitochondrial DNA copy number: a genome-wide and candidate gene study. Journal of Maternal-Fetal and Neonatal Medicine, 2017, 30, 2433-2439.	1.5	15
23	Identification of Differentially Expressed Genes in High-Density Oligonucleotide Arrays Accounting for the Quantification Limits of the Technology. Biometrics, 2003, 59, 542-554.	1.4	13
24	Initiation of Adjuvant Endocrine Therapy in Black and White Women With Breast Cancer. Clinical Breast Cancer, 2018, 18, 337-346.e1.	2.4	13
25	Circadian clock-related genetic risk scores and risk of placental abruption. Placenta, 2015, 36, 1480-1486.	1.5	10
26	Multivariate Bayesian Variable Selection Exploiting Dependence Structure Among Outcomes: Application to Air Pollution Effects on DNA Methylation. Biometrics, 2017, 73, 232-241.	1.4	10
27	Mixture of inhomogeneous matrix models for speciesâ€rich ecosystems. Environmetrics, 2015, 26, 39-51.	1.4	8
28	Physical activity and epigenetic biomarkers in maternal blood during pregnancy. Epigenomics, 2018, 10, 1383-1395.	2.1	8
29	Evaluation of normalization methods for analysis of LC-MS data. , 2012, , .		7
30	Evaluation of hierarchical models for integrative genomic analyses. Bioinformatics, 2016, 32, 738-746.	4.1	6
31	Identification of miRNA-mRNA associations in hepatocellular carcinoma using hierarchical integrative model. BMC Medical Genomics, 2020, 13, 56.	1.5	6
32	Bayesian Variable Selection Methods for Matched Case-Control Studies. International Journal of Biostatistics, 2017, 13, .	0.7	5
33	Integrative Analysis of DNA Methylation and microRNA Expression Reveals Mechanisms of Racial Heterogeneity in Hepatocellular Carcinoma. Frontiers in Genetics, 2021, 12, 708326.	2.3	5
34	Placental genetic variations in circadian clock-related genes increase the risk of placental abruption. International Journal of Molecular Epidemiology and Genetics, 2016, 7, 32-40.	0.4	5
35	A Bayesian Hierarchical Model for the Analysis of Affymetrix Arrays. Annals of the New York Academy of Sciences, 2004, 1020, 41-48.	3.8	4
36	Normalization of LC-MS data using Gaussian process. , 2012, , .		3

#	Article	IF	CITATIONS
37	Maternal-fetal genetic interactions, imprinting, and risk of placental abruption. Journal of Maternal-Fetal and Neonatal Medicine, 2022, 35, 3473-3482.	1.5	3
38	Risk of Common Mental Disorders in Relation to Symptoms of Obstructive Sleep Apnea Syndrome among Ethiopian College Students. Journal of Sleep Disorders Treatment & Care, 2015, 04, .	0.1	2
39	Analysis of LC-MS Data Using probabilitic-based mixture regression modelsAnalyse von LC-MS-Daten mit wahrscheinlichkeitsbasierter Mischung von Regressionsmodellen. Automatisierungstechnik, 2009, 57, 453-465.	0.8	1
40	An integrated pharmacogenomic analysis of doxorubic in response using genotype information on DMET genes. , 2013, , .		1
41	Bayesian Alignment Model for LC-MS Data. , 2011, , .		0
42	Integration of multiple data sources for identifying functional modules using Bayesian network. , 2012, , .		0
43	Variability assessment of LC-MS experiments and its application to experimental design and difference detection. , 2012, , .		0
44	Assessment of GC-MS in Detecting Changes in the Levels of Metabolites Using a Spike-in Experiment in Human Plasma. Metabolomics: Open Access, 2016, 6, .	0.1	0
45	Bayesian variable selection for high dimensional predictors and self-reported outcomes. BMC Medical Informatics and Decision Making, 2020, 20, 212.	3.0	0