

XIONG, Momiao Miao

University of Texas, Houston, USA

Selected Scholarly Contributions [Data Provided by **SCOPUS**]

Genome-wide gene and pathway analysis

European Journal of Human Genetics, 2010. Article in Press.

Gene and pathway-based second-wave analysis of genome-wide association studies.

European journal of human genetics : EJHG, 18 (1), pp. 111-117, 2010.

HLA-DPB1 and DPB2 are genetic loci for systemic sclerosis: A genome-wide association study in Koreans with replication in North Americans

Arthritis and Rheumatism, 60 (12), pp. 3807-3814, 2009.

Investigating gene and microRNA expression in glioblastoma

Proceedings - 2009 International Joint Conference on Bioinformatics, Systems Biology and Intelligent Computing, IJCBs 2009, pp. 17-22, 2009.

Gene and pathway-based second-wave analysis of genome-wide association studies

European Journal of Human Genetics, 2009. Article in Press.

Common variants of four bilirubin metabolism genes and their association with serum bilirubin and coronary artery disease in Chinese Han population

Pharmacogenetics and Genomics, 19 (4), pp. 310-318, 2009.

Association of polymorphisms in four bilirubin metabolism genes with serum bilirubin in three Asian populations

Human Mutation, 30 (4), pp. 609-615, 2009.

Discussion: Why do we test multiple traits in genetic association studies

Journal of the Korean Statistical Society, 38 (1), pp. 17-23, 2009.

Mutual information for testing gene-environment interaction

PLoS ONE, 4 (2), art. no. e4578, 2009.

New choices of search directions used in constructing quadratic interpolation models

Proceedings of the 7th International Conference on Machine Learning and Cybernetics, ICMLC, 4, pp. 2320-2325, 2008.

Multifactor effects and evidence of potential interaction between complement factor HY402H and LOC387715 A69S in age-related macular degeneration

PLoS ONE, 3 (12), art. no. e3833, 2008.

Structure-based graph distance measures of high degree of precision

Pattern Recognition, 41 (12), pp. 3547-3561, 2008.

Extended Kalman filter for estimation of parameters in nonlinear state-space models of biochemical networks

PLoS ONE, 3 (11), art. no. e3758, 2008.

Network quotients: Structural skeletons of complex systems

Physical Review E - Statistical, Nonlinear, and Soft Matter Physics, 78 (4), art. no. 046102, 2008.

Emergence of symmetry in complex networks

Physical Review E - Statistical, Nonlinear, and Soft Matter Physics, 77 (6), art. no. 066108, 2008.

Composite measure of linkage disequilibrium for testing interaction between unlinked loci

European Journal of Human Genetics, 16 (5), pp. 644-651, 2008.

Symmetry-based structure entropy of complex networks

Physica A: Statistical Mechanics and its Applications, 387 (11), pp. 2611-2619, 2008.

Differential dynamic properties of scleroderma fibroblasts in response to perturbation of environmental stimuli

PLoS ONE, 3 (2), art. no. e1693, 2008.