Data sharing statement: Data generated in this manuscript constitutes mRNA and proteomics information. Authors follow the NIH Grants Policy on Sharing of Unique Research Resources including the "Sharing of Biomedical Research Resources: Principles and Guidelines for Recipients of NIH Grants and Contracts" issued in December, 1999.

Dr. Lu Lu is responsible for coordinating data sharing through GeneNetwork (GN) at the: http://www.genenetwork.org/webqtl/main.py. GN is a group of linked data sets and tools used to study complex networks of genes, molecules, and higher order gene function and phenotypes. GN combines more than 25 years of legacy data generated by hundreds of scientists together with sequence data (SNPs) and massive transcriptome data sets (expression genetic or quantitative trait locus data sets). GN connected to numerous links to the UCSC and Ensembl Genome Browsers, PubMed, Entrez Gene, GNF Expression Atlas, ABI Panther, and WebGestalt provide users with rapid interpretive information about genomic regions, published phenotypes and genes.

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