Supplemental Tables

Protein Carbonylation in a Murine Model for Early Alcoholic Liver Disease

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Table S1. Composite list of proteins identified in this study. The assigned protein accession number, protein name, molecular weight, protein fraction, treatment group, unique spectra, total spectra and sequence coverage are indicated.

Table S2. DAVID analysis demonstrates key metabolic pathways and processes implicated in the pathogenesis of ALD. Count pertains to the number of proteins contained in the enriched category; % reveals the percentage of proteins identified in our list that were enriched in the given term. Only terms with a fold-enrichment greater than 5.0 and a Benjamini corrected *P* value less than 0.01 were included.