Supplementary table 1 legend:

The table lists proteins with significant differential expression (IRI/Con ratio ≥1.5 or ≤0.67, *p* < 0.05), including protein accession numbers, gene names, fold changes, *p*-values, and functional annotations (KEGG pathways, biological processes). Subcellular localization and structural domains are indicated. IRI, ischemia-reperfusion injury; CON, control group; MW, molecular weight; KEGG, kyoto encyclopedia of genes and genomes; KOG, eukaryotic krthologous groups; Pfam, protein families database.