# Local DAG for enriched terms in GO Biological Process

Nodes sized according to Binomial Fold Enrichment



**2.27** regulation of membrane permeability

**2.59** protein-lipid complex subunit organization

**2.17** adherens junction organization

**2.05** secondary alcohol metabolic process

**2.28** cellular response to topologically incorrect protein

**2.37** **2.04** response to cadmium ion / homeostasis

**2.11** regulation of generation of precursor metabolites and energy

**2.06** cholesterol metabolic process

**2.25** cellular response to unfolded protein

**2.54** **2.25** **2.48** regulation of triglyceride metabolic process / regulation of lipid metabolic process / regulation of mitochondrial membrane permeability

**2.35** endoplasmic reticulum unfolded protein response

**2.08** monocarboxylic acid catabolic process

**2.15** **2.13** regulation of gluconeogenesis / regulation of glucose metabolic process

**2.1** fatty acid catabolic process

**2.01** alpha-amino acid catabolic process

**2.09** regulation of ATP metabolic process

Node size legend: ( <1 ) ( 2 ) ( 3 ) ( 4 ) ( >5 )