fatty acid beta-oxidation response to using acyl-CoA endoplasmic endoplasmic regulation of cellu**regulation of** endoplasmic SRP-dependent cotranslational dentificacellular translational initiation reticulum cellulariosynthetic cotranslational protein transport unfolded protein amino acidess positive response metabolic process translation protein targeting metabolic metabolic transport integrin-mediated pathway to membrane process process signaling pathway vesicle-mediated vesicle-mediated translation transport, Golgi to transport endoplasmic reticulum protein import into cecepred extabilization negative mitochondrial matrix endoplasmic protein insertion regulation reticulum to Golgi into mitochondrial homeostasis formation of mitochondrial vesicle-mediated inner membrane of G2/M cytoplasmic translational transport translation transition of nuclear-transcribed mRNA elongation ion homeostasi translational fatty acid homeostasis catabolic process. mitotic cell cycle elongation complex nonsense-mediated decay extracellular translational matrix ribosomal large subunit assembly termination epithelial regulation o regulation of protein refolding by RNA mRNA stability large subunit assembly oolymerase folding mesoderm ubiquitin-independent autophagosome protein catabolic catabolic process mRNA splicing, anaphase-promoting mitochondrial respiratory maturation formation complex-dependent via spliceosome chain complex I assembly protein folding catabolic process