

**peroxisome, GO:0005777**

fatty acid beta-oxidation, GO:0006635  
protein processing, GO:0016485  
steroid metabolic process, GO:0008202  
regulation of lipid metabolic process, GO:0019216  
cholesterol metabolic process, GO:0008203  
flavin adenine dinucleotide binding, GO:0050660  
peroxisomal matrix, GO:0005782  
release of cytochrome c from mitochondria, GO:0001836  
phosphatidylcholine acyl-chain remodeling, GO:0036151  
oxidoreductase activity, GO:0016616  
mitochondrial fusion, GO:0008053  
fatty-acyl-CoA binding, GO:0000062  
pyridoxal phosphate binding, GO:0030170  
cholesterol biosynthetic process, GO:0006695  
ATPase activity, GO:0042626  
protein tetramerization, GO:0051262  
steroid biosynthetic process, GO:0006694  
clathrin binding, GO:0030276  
oxidoreductase activity, GO:0016627  
xenobiotic metabolic process, GO:0006805  
fatty acid beta-oxidation using acyl-CoA oxidase, GO:0033540  
phosphatidylethanolamine acyl-chain remodeling, GO:0036152  
sterol transporter activity, GO:0015248  
alpha-linolenic acid metabolic process, GO:0036109  
2-oxoglutarate metabolic process, GO:0006103  
Cdc73/Paf1 complex, GO:0016593  
regulation of ARF protein signal transduction, GO:0032012  
carboxylic ester hydrolase activity, GO:0052689  
enoyl-CoA hydratase activity, GO:0004300  
fatty acid beta-oxidation using acyl-CoA dehydrogenase, GO:0033539  
lipid phosphatase activity, GO:0042577  
prostaglandin metabolic process, GO:0006693  
polyamine biosynthetic process, GO:0006596  
phosphatidylinositol acyl-chain remodeling, GO:0036149  
generation of precursor metabolites and energy, GO:0006091  
sterol transport, GO:0015918  
negative regulation of innate immune response, GO:0045824  
isoprenoid biosynthetic process, GO:0008299  
cobalamin metabolic process, GO:0009235  
regulation of cholesterol metabolic process, GO:0090181  
phospholipid dephosphorylation, GO:0046839  
integral component of mitochondrial membrane, GO:0032592  
phospholipase A2 activity, GO:0004623