

regulation of lipid metabolic process, GO:0019216

- steroid biosynthetic process, GO:0006694
- oxidoreductase activity, GO:0016627
- lipid homeostasis, GO:0055088
- alpha-linolenic acid metabolic process, GO:0036109
- epoxygenase P450 pathway, GO:0019373
- sterol biosynthetic process, GO:0016126
- isoprenoid biosynthetic process, GO:0008299
- proteasomal protein catabolic process, GO:0010498
- sphingosine biosynthetic process, GO:0046512
- fatty acid beta-oxidation using acyl-CoA dehydrogenase, GO:0033539
- triglyceride homeostasis, GO:0070328
- positive regulation of branching involved in ureteric bud morphogenesis, GO:0090190
- anion transmembrane transporter activity, GO:0008509
- hydrolase activity, GO:0016811
- lipid transporter activity, GO:0005319
- phospholipid catabolic process, GO:0009395
- cholesterol efflux, GO:0033344
- dopamine binding, GO:0035240
- positive regulation of blood pressure, GO:0045777
- aromatase activity, GO:0070330
- phospholipid transporter activity, GO:0005548
- phosphatidylcholine binding, GO:0031210