

lyase activity, GO:0016829

- cellular amino acid biosynthetic process, GO:0008652
- gluconeogenesis, GO:0006094
- cellular amino acid metabolic process, GO:0006520
- protein tetramerization, GO:0051262
- peroxisomal matrix, GO:0005782
- pyridoxal phosphate binding, GO:0030170
- purine nucleotide biosynthetic process, GO:0006164
- biosynthetic process, GO:0009058
- tricarboxylic acid cycle, GO:0006099
- presynaptic active zone, GO:0048786
- generation of precursor metabolites and energy, GO:0006091
- pyrimidine nucleotide biosynthetic process, GO:0006221
- tRNA splicing, GO:0006388
- pyruvate metabolic process, GO:0006090
- carboxy-lyase activity, GO:0016831
- carboxylic acid binding, GO:0031406
- negative regulation of G-protein coupled receptor protein signaling pathway, GO:0045744
- transaminase activity, GO:0008483
- carboxylic acid metabolic process, GO:0019752
- purine nucleobase biosynthetic process, GO:0009113
- cyclic nucleotide biosynthetic process, GO:0009190
- fatty-acyl-CoA binding, GO:0000062
- nitrogen compound metabolic process, GO:0006807
- regulation of rhodopsin mediated signaling pathway, GO:0022400
- very long-chain fatty acid biosynthetic process, GO:0042761
- inhibitory synapse, GO:0060077
- dendritic cell chemotaxis, GO:0002407
- amino acid binding, GO:0016597
- NADPH binding, GO:0070402
- polyamine metabolic process, GO:0006595
- molybdopterin cofactor biosynthetic process, GO:0032324
- very long-chain fatty acid metabolic process, GO:0000038
- carbonate dehydratase activity, GO:0004089
- L-phenylalanine catabolic process, GO:0006559
- peptide metabolic process, GO:0006518
- fatty acid alpha-oxidation, GO:0001561
- guanylate cyclase activity, GO:0004383
- guanylate cyclase complex, GO:0008074
- glutamate metabolic process, GO:0006536
- fatty acid beta-oxidation using acyl-CoA oxidase, GO:0033540
- adenylate cyclase activity, GO:0004016
- leukotriene biosynthetic process, GO:0019370
- polyamine biosynthetic process, GO:0006596
- hydrolase activity, GO:0016811
- epithelial cilium movement, GO:0003351
- lipoxygenase pathway, GO:0019372
- phosphodiesterase I activity, GO:0004528
- malate metabolic process, GO:0006108
- phosphorus-oxygen lyase activity, GO:0016849