

mitochondrial intermembrane space, GO:0005758

- integral component of mitochondrial inner membrane, GO:0031305
- copper ion binding, GO:0005507
- oxidative phosphorylation, GO:0006119
- regulation of androgen receptor signaling pathway, GO:0060765
- nucleotide biosynthetic process, GO:0009165
- antioxidant activity, GO:0016209
- porphyrin-containing compound biosynthetic process, GO:0006779
- coenzyme A biosynthetic process, GO:0015937
- response to reactive oxygen species, GO:0000302
- NAD+ binding, GO:0070403
- cardiolipin binding, GO:1901612
- response to activity, GO:0014823
- response to iron ion, GO:0010039
- response to muscle activity, GO:0014850
- protein channel activity, GO:0015266
- nucleoside diphosphate kinase activity, GO:0004550
- positive regulation of transcription regulatory region DNA binding, GO:2000679
- response to L-ascorbic acid, GO:0033591
- negative regulation of cholesterol biosynthetic process, GO:0045541
- membrane depolarization, GO:0051899
- peptide metabolic process, GO:0006518
- nucleoside metabolic process, GO:0009116
- protein phosphatase 2B binding, GO:0030346
- cellular response to gonadotropin stimulus, GO:0071371
- peroxidase activity, GO:0004601
- protein homotrimerization, GO:0070207
- molybdopterin cofactor binding, GO:0043546
- myelination in peripheral nervous system, GO:0022011