

**respiratory chain, GO:0070469**

- mitochondrial respiratory chain complex I assembly, GO:0032981
- mitochondrial intermembrane space, GO:0005758
- mitochondrial electron transport, GO:0006120
- mitochondrial respiratory chain complex I, GO:0005747
- NADH dehydrogenase (ubiquinone) activity, GO:0008137
- electron carrier activity, GO:0009055
- aerobic respiration, GO:0009060
- NADH dehydrogenase activity, GO:0003954
- hydrogen ion transmembrane transport, GO:1902600
- mitochondrial electron transport, GO:0006122
- oxidative phosphorylation, GO:0006119
- mitochondrial respiratory chain complex III, GO:0005750
- ubiquinol-cytochrome-c reductase activity, GO:0008121
- cardiovascular system development, GO:0072358
- cytochrome-c oxidase activity, GO:0004129
- cellular respiration, GO:0045333
- mitochondrial electron transport, GO:0006123
- mitochondrial respiratory chain, GO:0005746
- mitochondrial respiratory chain complex IV, GO:0005751
- respiratory electron transport chain, GO:0022904
- oxidoreductase activity, GO:0016651
- coenzyme binding, GO:0050662
- mitochondrial respiratory chain complex III assembly, GO:0034551
- response to nicotine, GO:0035094
- quinone binding, GO:0048038
- response to mercury ion, GO:0046689
- cerebellum development, GO:0021549