respiratory chain, GO:0070469

mitochondrial membrane, GO:0031966 mitochondrial intermembrane space, GO:0005758 mitochondrial respiratory chain complex I assembly, GO:0032981 electron carrier activity, GO:0009055 mitochondrial electron transport, GO:0006120 mitochondrial respiratory chain complex I, GO:0005747 NADH dehydrogenase (ubiquinone) activity, GO:0008137 aerobic respiration, GO:0009060 NAD binding, GO:0051287 hydrogen ion transmembrane transport, GO:1902600 mitochondrial electron transport, GO:0006122 mitochondrial respiratory chain complex III, GO:0005750 cellular respiration, GO:0045333 2 iron, GO:0051537 oxidoreductase activity, GO:0016651 coenzyme binding, GO:0050662 metalloendopeptidase activity, GO:0004222 NADH dehydrogenase activity, GO:0003954 respiratory electron transport chain, GO:0022904 mitochondrial ATP synthesis coupled electron transport, GO:0042775 reactive oxygen species metabolic process, GO:0072593 heme binding, GO:0020037 oxidative phosphorylation, GO:0006119 apoptotic mitochondrial changes, GO:0008637 mitochondrial respiratory chain, GO:0005746 ubiquinol-cytochrome-c reductase activity, GO:0008121 cAMP-mediated signaling, GO:0019933 mitochondrial electron transport, GO:0006123 cytochrome-c oxidase activity, GO:0004129 lactation, GO:0007595