

A

GM969 shXPA vs. scrambled	XPA deficient vs. XPA complemented
GO0046870 CADMIUM ION BINDING	GO0000786 NUCLEOSOME
GO0003735 STRUCTURAL CONSTITUENT OF RIBOSOME	GO0006334 NUCLEOSOME ASSEMBLY
GO0005840 RIBOSOME	GO0007001 CHROMOSOME ORGANIZATION AND BIOGENESIS (
GO0030529 RIBONUCLEOPROTEIN COMPLEX	* GO0005743 MITOCHONDRIAL INNER MEMBRANE
GO0006334 NUCLEOSOME ASSEMBLY	* GO0008137 NADH DEHYDROGENASE (UBIQUINONE) ACTIVITY
GO0007001 CHROMOSOME ORGANIZATION AND BIOGENESIS (GO0005694 CHROMOSOME
GO0005694 CHROMOSOME	* GO0005739 MITOCHONDRION
GO0015935 SMALL RIBOSOMAL SUBUNIT	GO0046870 CADMIUM ION BINDING
GO0000786 NUCLEOSOME	GO0005507 COPPER ION BINDING
GO0005507 COPPER ION BINDING	* GO0004129 CYTOCHROME C OXIDASE ACTIVITY
* GO0005743 MITOCHONDRIAL INNER MEMBRANE	* GO0005746 MITOCHONDRIAL RESPIRATORY CHAIN
* GO0005739 MITOCHONDRION	* GO0006120 MITOCHONDRIAL ELECTRON TRANSPORT NADH T
* GO0005746 MITOCHONDRIAL RESPIRATORY CHAIN	GO0005840 RIBOSOME
* GO0006120 MITOCHONDRIAL ELECTRON TRANSPORT NADH T	GO0030529 RIBONUCLEOPROTEIN COMPLEX
* GO0004129 CYTOCHROME C OXIDASE ACTIVITY	GO0003735 STRUCTURAL CONSTITUENT OF RIBOSOME
* GO0008137 NADH DEHYDROGENASE (UBIQUINONE) ACTIVITY	GO0015935 SMALL RIBOSOMAL SUBUNIT
GO0051028 MRNA TRANSPORT	GO0006810 TRANSPORT
GO0004672 PROTEIN KINASE ACTIVITY	GO0007264 SMALL GTPASE MEDIATED SIGNAL TRANSDUCTIO
GO0005575 CELLULAR COMPONENT	GO0005575 CELLULAR COMPONENT
GO0006468 PROTEIN AMINO ACID PHOSPHORYLATION	GO0051028 MRNA TRANSPORT
GO0015031 PROTEIN TRANSPORT	GO0005515 PROTEIN BINDING
GO0000166 NUCLEOTIDE BINDING	GO0004672 PROTEIN KINASE ACTIVITY
GO0006810 TRANSPORT	GO0015031 PROTEIN TRANSPORT
GO0007264 SMALL GTPASE MEDIATED SIGNAL TRANSDUCTIO	GO0006468 PROTEIN AMINO ACID PHOSPHORYLATION
GO0005515 PROTEIN BINDING	GO0000166 NUCLEOTIDE BINDING









