

Full-Length

COX6A1 (Human) Recombinant Protein (P01)

Catalog # H00001337-P01 Size 25 ug, 10 ug

Applications



Specification	
Product Description	Human COX6A1 full-length ORF (AAH07723, 1 a.a 109 a.a.) recombinant protein with GST-tag at N-terminal.
Sequence	MAVVGVSSVSRLLGRSRPQLGRPMSSGAHGEEGSARMWKTLTFFVALPGVAVSMLNVYLKSHH GEHERPEFIAYPHLRIRTKPFPWGDGNHTLFHNPHVNPLPTGYEDE
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	37.73
Interspecies Antigen Sequence	Rat (83)
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCI, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.
Storage Instruction	Store at -80°C. Aliquot to avoid repeated freezing and thawing.
Note	Best use within three months from the date of receipt of this protein.



Applications

- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — COX6A1	
Entrez GenelD	<u>1337</u>
GeneBank Accession#	<u>BC007723</u>
Protein Accession#	AAH07723
Gene Name	COX6A1
Gene Alias	COX6A, COX6AL, MGC104500
Gene Description	cytochrome c oxidase subunit Vla polypeptide 1
Omim ID	<u>602072</u>
Gene Ontology	Hyperlink
Gene Summary	Cytochrome c oxidase (COX), the terminal enzyme of the mitochondrial respiratory chain, catalyz es the electron transfer from reduced cytochrome c to oxygen. It is a heteromeric complex consisti ng of 3 catalytic subunits encoded by mitochondrial genes and multiple structural subunits encode d by nuclear genes. The mitochondrially-encoded subunits function in the electron transfer and the nuclear-encoded subunits may function in the regulation and assembly of the complex. This nuclea r gene encodes polypeptide 1 (liver isoform) of subunit VIa, and polypeptide 1 is found in all non-muscle tissues. Polypeptide 2 (heart/muscle isoform) of subunit VIa is encoded by a different gen e, and is present only in striated muscles. These two polypeptides share 66% amino acid sequen ce identity. It has been reported that there may be several pseudogenes on chromosome 1p31.1 has been documented. [provided by RefSeq
Other Designations	OTTHUMP00000045456 cytochrome C oxidase subunit Vla homolog

Pathway

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- Cardiac muscle contraction
- <u>Metabolic pathways</u>
- Oxidative phosphorylation