Product Information

## MaxPab®

## NDUFA5 purified MaxPab rabbit polyclonal antibody (D01P)

Catalog \# H00004698-D01P Size 100 ug

## Applications



## Western Blot (Transfected Iysate)

Western Blot analysis of NDUFA5 expression in transfected 293T cell line (H00004698-T02) by NDUFA5 MaxPab polyclonal antibody.

Lane 1: NDUFA5 transfected lysate(13.50 KDa).
Lane 2: Non-transfected lysate.

| Specification |  |
| :---: | :---: |
| Product Description | Rabbit polyclonal antibody raised against a full-length human NDUFA5 protein. |
| Immunogen | NDUFA5 (NP_004991.1, 1 a.a. $\sim 116$ a.a) full-length human protein. |
| Sequence | MAGVLKKTTGLVGLAVCNTPHERLRILYTKILDVLEEIPKNAAYRKYTEQITNEKLAMVKAEPDVKKL EDQLQGGQLEEVILQAEHELNLARKMREWKLWEPLVEEPPADQWKWPI |
| Host | Rabbit |
| Reactivity | Human |
| Quality Control Testing | Antibody reactive against mammalian transfected lysate. |
| Storage Buffer | In 1x PBS, pH 7.4 |
| Storage Instruction | Store at $-20^{\circ} \mathrm{C}$ or lower. Aliquot to avoid repeated freezing and thawing. |

## Applications

Product Information

- Western Blot (Transfected lysate)

Western Blot analysis of NDUFA5 expression in transfected 293T cell line (H00004698-T02) by NDUFA5 MaxPab polyclonal antibody.

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Protocol Download

| Gene Info - NDUFA5 |  |
| :---: | :---: |
| Entrez GenelD | 4698 |
| GeneBank Accession\# | NM 0005000.2 |
| Protein Accession\# | NP_004991.1 |
| Gene Name | NDUFA5 |
| Gene Alias | B13, C-13KD-B, DKFZp781K1356, FLJ12147, NUFM, UQOR13 |
| Gene Description | NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5, 13kDa |
| Omim ID | 601677 |
| Gene Ontology | Hyperlink |
| Gene Summary | The human NDUFA5 gene codes for the B13 subunit of complex I of the respiratory chain, which tr ansfers electrons from NADH to ubiquinone. The high degree of conservation of NDUFA5 extendi ng to plants and fungi indicates its functional significance in the enzyme complex. The protein loca lizes to the inner mitochondrial membrane as part of the 7 component-containing, water soluble "ir on-sulfur protein" (IP) fraction of complex I , although its specific role is unknown. It is assumed to u ndergo post-translational removal of the initiator methionine and N -acetylation of the next amino a cid. The predicted secondary structure is primarily alpha helix, but the carboxy-terminal half of the protein has high potential to adopt a coiled-coil form. The amino-terminal part contains a putative beta sheet rich in hydrophobic amino acids that may serve as mitochondrial import signal. Relate d pseudogenes have also been identified on four other chromosomes. [provided by RefSeq |
| Other Designations | Complex I-13KD-B\|NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 |NADH dehydroge nase (ubiquinone) 1 alpha subcomplex, 5 ( $13 \mathrm{kD}, \mathrm{B} 13$ )type I dehydrogenase\|ubiquinone reductas e |

## Pathway

- Metabolic pathways
- Oxidative phosphorylation


## Disease

- Genetic Predisposition to Disease
- Prostatic Neoplasms
- Tobacco Use Disorder

