## 32-2594: NDUFS2 Recombinant Protein

Alternative Name :
$\mathrm{Cl}-49$,NADH dehydrogenase [ubiquinone] iron-sulfur protein 2,mitochondrial,Complex l-49kD,Cl-49kD,NADH-ubiquinone oxidoreductase 49 kDa subunit,NDUFS2.

## Description

Source : Escherichia Coli. NDUFS2 Human Recombinant produced in E.coli is a single, non-glycosylated polypeptide chain containing 410 amino acids (77-463a.a) and having a molecular mass of 46.5 kDa . NDUFS2 is fused to a 23 amino acid His-tag at N -terminus \& purified by proprietary chromatographic techniques. NDUFS2 is a core subunit of the mitochondrial membrane respiratory chain NADH dehydrogenase (Complex I) which is a part of the minimal assembly required for catalysis. Complex I takes part in the transfer of electrons from NADH to the respiratory chain. Histidine NADH Dehydrogenase Fe-S Protein 2 (NDUFS2) is required for catalytic activity. Imperfections in NDUFS2 are the source of complex I mitochondrial respiratory chain deficiency, which is characterized by many symptoms including liver failure, cardiomyopathy and neurodegeneration.

## Product Info

| Amount : | $10 \mu \mathrm{~g}$ |
| :---: | :---: |
| Purification : | Greater than $80 \%$ as determined by SDS-PAGE. |
| Content : | The NDUFS2 solution ( $0.25 \mathrm{mg} / \mathrm{ml}$ ) contains 20 mM Tris- HCl buffer ( pH 8.0 ), 0.4 M Urea and $10 \%$ glycerol. |
| Storage condition : | Store at $4^{\circ} \mathrm{C}$ if entire vial will be used within $2-4$ weeks. Store, frozen at $-20^{\circ} \mathrm{C}$ for longer periods of time. For long term storage it is recommended to add a carrier protein ( $0.1 \%$ HSA or BSA).Avoid multiple freeze-thaw cycles. |
| Amino Acid : | MGSSHHHHHH SSGLVPRGSH MGSVKNITLN FGPQHPAAHG VLRLVMELSG EMVRKCDPHI |
|  | GLLHRGTEKL IEYKTYLQAL PYFDRLDYVS MMCNEQAYSL AVEKLLNIRP PPRAQWIRVL |
|  | FGEITRLLNH IMAVTTHALD LGAMTPFFWL FEEREKMFEF YERVSGARMH AAYIRPGGVH |
|  | QDLPLGLMDD IYQFSKNFSL RLDELEELLT NNRIWRNRTI DIGVVTAEEA LNYGFSGVML |
|  | RGSGIQWDLR KTQPYDVYDQ VEFDVPVGSR GDCYDRYLCR VEEMRQSLRI IAQCLNKMPP |
|  | GEIKVDDAKV SPPKRAEMKT SMESLIHHFK LYTEGYQVPP GATYTAIEAP KGEFGVYLVS |
|  | DGSSRPYRCK IKAPGFAHLA GLDKMSKGHM LADVVAIIGT QDIVFGEVDR. |



