RPD312Hu01 100μg

Recombinant Cytochrome P450 Reductase (CPR)

Organism Species: Homo sapiens (Human)

Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY

NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

10th Edition (Revised in Jan, 2014)

[PROPERTIES]

Residues: Gly77~Phe518

Tags: Two N-terminal Tags, His-tag and T7-tag

Accession: P16435

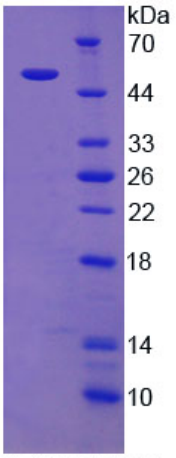
Host: *E. coli*

Subcellular Location: Endoplasmic reticulum membrane; Peripheral membrane protein.

Purity: >95%

Endotoxin Level: <1.0EU per 1μg (determined by the LAL method).

Formulation: Supplied as lyophilized form in 20mM Tris, 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% sarcosyl, 5% trehalose, and preservative.



15% SDS-PAGE

Predicted isoelectric point: 5.0

Predicted Molecular Mass: 53.4kDa

Applications: SDS-PAGE; WB; ELISA; IP.

(May be suitable for use in other assays to be determined by the end user.)

[USAGE]

Reconstitute in sterile ddH_2O.

[STORAGE AND STABILITY]

Storage: Avoid repeated freeze/thaw cycles.

Store at 2-8°C for one month.

Aliquot and store at -80°C for 12 months.

Stability Test: The thermal stability is described by the loss rate of the target protein. The loss rate was determined by accelerated thermal degradation test, that is, incubate the protein at 37°C for 48h, and no obvious degradation and precipitation were observed. (Referring from China Biological Products Standard, which was calculated by the Arrhenius equation.) The loss of this protein is less than 5% within the expiration date under appropriate storage condition.

[SEQUENCES]

The sequence of the target protein is listed below.

GRNI IVFYGSQTGT AEEFANRLSK DAHRYGMRGM SADPEEYDLA DLSSLPEIDN ALVVFCMATY GEGDPTDNAQ DFYDWLQETD VDLSGVKFAV FGLGNKTYEH FNAMGKYVDK RLEQLGAQRI FELGLGDDDG NLEEDFITWR EQFWPAVCEH FGVEATGEES SIRQYELVVH TDIDAAKVYM GEMGRLKSYE NQKPPFDAKN PFLAAVTTNR KLNQGTERHL MHLELDISDS KIRYESGDHV AVYPANDSAL VNQLGKILGA DLDVVMSLNN LDEESNKKHP FPCPTSYRTA LTYYLDITNP PRTNVLYELA QYASEPSEQE LLRKMASSSG EGKELYLSWV VEARRHILAI LQDCPSLRPP IDHLCELLPR LQARYYSIAS SSKVHPNSVH ICAVVVEYET KAGRINKGVA TNWLRAKEPA GENGGRALVP MFVRKSQF