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YBR876Ra01 100µg

Recombinant Coenzyme Q6 Homolog, Monooxygenase (COQ6)

Organism Species: Rattus norvegicus (Rat)

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: Met1~Arg476 Tags: Two N-terminal Tags, His-tag and T7-tag Accession: Q68FU7 Host: *E. coli* Subcellular Location: Golgi apparatus. Cell projection. Purity: >90% Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl. Predicted isoelectric point: 6.7



Predicted Molecular Mass: 55.2kDa

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

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

[<u>USAGE</u>]

Reconstitute in sterile PBS, pH7.2-pH7.4.



[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.

[<u>SEQUENCES</u>]

The sequence of the target protein is listed below.

MA A R I G P M A G L L C V R W W S TA Q L A A R G G P LV A C R R W T S S S T D S V Y D V V V S G GGMVGSAMAC ALGHDIHFHD KKILLLEAGP KKTLEKLSET YSNRVSSISL GSTTLLSSFG AWDHICNMRC KAFRRMQVWD SCSEALIMFD KDNLDDMGYI VENDVIMHAI TKQLEAVADR VKVLYESKAV GYAWPGPFSL ADSSPWVHIT LGDGSTLQTK LLIGADGHNS GVRQAAGIQN VGWNYDQSAV VATLHLSEAT ENNVAWQRFL PSGPIALLPL SDTLSSLVWS TSHAHAAELV SMHEEEFVDA INSAFWSDVH HTDFVDSASA MVHHAVALLK PTKVSARQLP PSVAKVDAKS RALFPLGLGH AAEYVRPRVA LIGDAAHRVH PLAGQGVNMG FGDISSLIHY LSTAAFNGKD LGSMSHLTGY ETDRQRHNTA LLAATDLLKR LYSTSTTPLV LLRTWGLQAT NAVSPLKEQI MAFASR