

YBD546Ra01 100µg Recombinant Cytochrome P450 11B1 (CYP11B1) Organism Species: Rattus norvegicus (Rat) Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

9th Edition (Revised in Jul, 2013)

## [ <u>PROPERTIES</u> ]

Residues: Metl~Val498 (Accession # P15393), with two N-terminal Tags, His-tag and T7-tag. Host: *E. coli* Subcellular Location: Mitochondrion membrane. Purity: >95% 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: 9.6 Predicted Molecular Mass: 61.2kDa 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.





## [ <u>STORAGE AND STABILITY</u> ]

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  $37^{\circ}$ 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 target protein is fused with two N-terminal Tags, His-tag and T7-tag, its sequence is listed below.

MGSSHHHHHH SSGLVPRGSH MASMTGGQQM GRGSEF- MALRVTADVW LARPWQCLHR TRALGTTAKV APKTLKPFEA IPQYSRNKWL KMIQILREQG QENLHLEMHQ AFQELGPIFR HSAGGAQIVS VMLPEDAEKL HQVESILPHR MPLEPWVAHR ELRGLRRGVF LLNGADWRFN RLQLNPNMLS PKAIQSFVPF VDVVARDFVE NLKKRMLENV HGSMSINIQS NMFNYTMEAS HFVISGERLG LTGHDLKPES VTFTHALHSM FKSTTQLMFL PKSLTRWTST RVWKEHFDSW DIISEYVTKC IKNVYRELAE GRQQSWSVIS EMVAQSTLSM DAIHANSMEL IAGSVDTTAI SLVMTLFELA RNPDVQQALR QESLAAEASI VANPQKAMSD LPLLRAALKE TLRLYPVGSF VERIVHSDLV LQNYHVPAGT FVIIYLYSMG RNPAVFPRPE RYMPQRWLER KRSFQHLAFG FGVRQCLGRR LAEVEMLLLL HHMLKTFQVE TLRQEDMQMV FRFLLMPSSS PFLTFRPV