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YBJ551Ra01 100µg

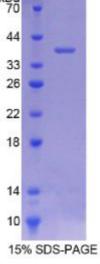
Recombinant Biliverdin Reductase A (BLVRA)

Organism Species: Rattus norvegicus (Rat)

Instruction manual

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

[PROPERTIES] Residues: Gly10~His279 Tags: Two N-terminal Tags, His-tag and T7-tag Accession: P46844 Host: E. coli Subcellular Location: Cytoplasm. **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: 5.4 Predicted Molecular Mass: 34.4kDa



Accurate Molecular Mass: 38kDa as determined by SDS-PAGE reducing conditions. Applications: SDS-PAGE; WB; ELISA; IP.

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

The possible reasons that the actual band size differs from the predicted are as follows: Note:

- 1. Splice variants: Alternative splicing may create different sized proteins from the same gene.
- 2. Relative charge: The composition of amino acids may affects the charge of the protein.
- 3. Post-translational modification: Phosphorylation, glycosylation, methylation etc.
- 4. Post-translation cleavage: Many proteins are synthesized as pro-proteins, and then cleaved to give the active form.

10th Edition (Revised in Jan, 2014)



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[USAGE]

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.

[SEQUENCES]

The sequence of the target protein is listed below.

G VVVVGVGRAG SVRLRDLKDP RSAAFLNLIG FVSRRELGSL DEVRQISLED ALRSQEIDVA YICSESSSHE DYIRQFLQAG KHVLVEYPMT LSFAAAQELW ELAAQKGRVL HEEHVELLME EFEFLRREVL GKELLKGSLR FTASPLEEER FGFPAFSGIS RLTWLVSLFG ELSLISATLE ERKEDQYMKM TVQLETQNKG LLSWIEEKGP GLKRNRYVNF QFTSGSLEEV PSVGVNKNIF LKDQDIFVQK LLDQVSAEDL AAEKKRIMH