TEL:4006-871-227 Web:www.ybio.net Email:shybio@126.com

YBB699Hu01 100µg Angiopoietin Like Protein 3 (ANGPTL3) Organism Species: Homo sapiens (Human) *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: Ser17~Glu460 (Accession # Q9Y5C1), with kDa 94 N-terminal His-Tag. 66.2 45 Host: E. coli 33 Subcellular Location: Secreted. 26 Purity: >95% 20 Endotoxin Level: <1.0EU per 1µg 14.4 (determined by the LAL method). Formulation: Supplied as lyophilized form in PBS, pH7.4 15% SDS-PAGE containing 5% sucrose, 0.01% sarcosyl. Predicted isoelectric point: 6.3 Predicted Molecular Mass: 53.3kDa 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.

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

Storage: Avoid repeated freeze/thaw cycles.

Store at $2-8^{\circ}C$ 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 target protein is fused with N-terminal His-Tag, its sequence is listed below. MGHHHHHHSG SEF-SRID QDNSSFDSLS PEPKSRFAML DDVKILANGL LQLGHGLKDF VHKTKGQIND IFQKLNIFDQ SFYDLSLQTS EIKEEEKELR RTTYKLQVKN EEVKNMSLEL NSKLESLLEE KILLQQKVKY LEEQLTNLIQ NQPETPEHPE VTSLKTFVEK QDNSIKDLLQ TVEDQYKQLN QQHSQIKEIE NQLRRTSIQE PTEISLSSKP RAPRTTPFLQ LNEIRNVKHD GIPAECTTIY NRGEHTSGMY AIRPSNSQVF HVYCDVISGS PWTLIQHRID GSQNFNETWE NYKYGFGRLD GEFWLGLEKI YSIVKQSNYV LRIELEDWKD NKHYIEYSFY LGNHETNYTL H LVA I T G N V P N A I P E N K D LV F S T W D H K A K G H F N C P E G Y S G G W W H D E C G E NNLNGKYNKP RAKSKPERRR GLSWKSQNGR LYSIKSTKML IHPTDSESFE