



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

YB91967Po01 100 μ g

Apolipoprotein A4 (APOA4)

Organism: *Sus scrofa*; Porcine (Pig)

Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY

NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

7th Edition (Revised in May, 2013)

[PROPERTIES]

Residues: Glu21~Gly382 (Accession # 046409),
with N-terminal His-Tag.

Host: *E. coli*

Subcellular Location: Secreted.

Purity: >95%

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

Formulation: Supplied as lyophilized form in PBS,
pH7.4, containing 5% sucrose, 0.01% sarcosyl.

Predicted isoelectric point: 5.8

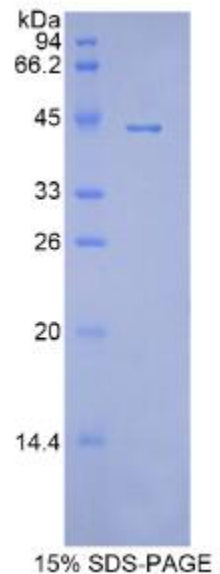
Predicted Molecular Mass: 42.8kDa

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 PBS, pH7.2-pH7.4.





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

[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 target protein is fused with N-terminal His-Tag, its sequence is listed below.

MGHHHHHSG SEF-EVNADQVATV MWDYFSQLGS NAKKAVEHLQ KSELTQQLNT
L F Q D K L G E V N T Y T E D L Q K K L V P F A T E L H E R L T K D S E K L K E E I R R E L E E L R
ARLLPHATEVSQKIGDNVRE LQQLGPFTG GLRTQVNTQV QQLQRQLKPY AERMESVLRQ
NIRNLEASVA PYADEFKAKI DQNV EELKGS LTPYAEELKA KIDQNV EELR RSLAPYAQDV
QEKLNHQLEG LAFQMKKQAE ELKAKISANA DELRQKLPV AENVHGLKG NTEGLQKSLL
ELRSHLDQQV EEFRLKVEPY GETFNKALVQ QVEDLRQKLG PLAGDVEGHL SFLEKDLRDK
VNTFFSTLKEEASQGSQAL PAQEKAQAPL EG