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YBA740Hu01 100µg

Recombinant Prostaglandin I Synthase (PTGIS)

Organism Species: Homo sapiens (Human)

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~Pro500

Tags: Two N-terminal Tags, His-tag and T7-tag

Accession: Q16647

Host: *E. coli*

Subcellular Location: Endoplasmic reticulum membrane.

Single-pass membrane protein.

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: 6.8

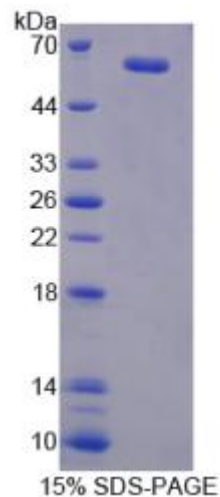
Predicted Molecular Mass: 60.8kDa

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.





[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.

MAWAALLGLL AALLLLLLLS RRRTRRPGEP PLDLGSIPWL GYALDFGKDA ASFLTRMKEK
HGDIFTILVG GRYVTVLLDP HSYDAVVWEP RTRLDFHAYA IFLMERIFDV QLPHYSPSDE
KARMKLTLLH RELQALTEAM YTNLHAVLLG DATEAGSGWH EMGLLDFSYS FLLRAGYLTL
YGIEALPRTH ESQAQDRVHS ADVFHTFRQL DRLLPKLARG SLSVGDKDHM CSVKSRLWKL
LSPARLARRA HRSKWLESYL LHLEEMGVSE EMQARALVLQ LWATQGNMGP AAFWLLLFL
KNPEALAAVR GELESILWQA EQPVSQTTTL PQKVL DSTPV LDSVLSESLR LTAAPFITRE
VVVDLAMPMA DGREFNLRRG DRLLLFPFLS PQRDPEIYTD PEVFKYNRFL NPDGSEKKDF
YKDGKRLKNY NMPWGAGHNN CLGRSYAVNS IKQFVFLVLV HLDLELINAD VEIPEFDLSR
YGFGLMQPEH DVPVRYRIRP