



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

YBP121Hu01 100 $\mu$ g

Recombinant Wingless Type MMTV Integration Site Family, Member 16 (WNT16)

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)

## [ PROPERTIES ]

Residues: Asn30~Lys365 (Accession # Q9UBV4), with two  
N-terminal Tags, His-tag and GST-tag.

Host: *E. coli*

Subcellular Location: Secreted, extracellular space,  
extracellular matrix.

Purity: >95%

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

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

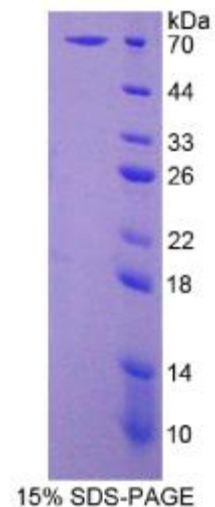
Predicted isoelectric point: 8.6

Predicted Molecular Mass: 70.0kDa

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 target protein is fused with two N-terminal Tags, His-tag and GST-tag, its sequence is listed below.

MSPILGYWKI KGLVQPTRLL LEYLEEKYEE HLYERDEGDK WRNKKFELGL EFPNLPYYID  
GDVKLTQ SMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSR IA YSKDFETLKV  
DFLSKLP EML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK  
KRIEAI PQID KYLKSSKYIA WPLQGWA TF GGGDHPPKSD GSTSGSGHHH HHSAGLVPR  
GSTAIGM KET AAKFERQHM DSPDLGTGGG SGIEGRGSMG YRGSEF-N WWLGIASFG  
VPEKLG CANL PLNSRQKELC KRKPYLLPSI REGARLGIQE CGSQFRHERW NCMITAAATT  
APMGASPL FG YELSSGKTET AFYAVMAAG LVHSVTRSCS AGNMTECSCD TTLQNGGSAS  
EG W H W G G C S D DV Q Y G M W F S R KF L D F P I G N T TG K E N K V L L A MN L H N N E A G R  
QAVAKLMSVD CRCHGVSGSC AVKTCWKTMS SFEKIGHLLK DKYENSIQIS DKTKRKMRRR  
EKDQRKIPIH KDDL LYV NKS PNYCEDKKL GIPGTQGREC NRTSEGADGC NLLCCGRGYN  
THVVRHVERC ECKFIWCCYV RCRRCESMTD VHTCK