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

YBA249Hu01 100µg Recombinant Nischarin (NISCH) 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 ]	-	kDa 70
Residues: Leu1246~Ser1346 (Accession # Q9Y2I1), with		44
two N-terminal Tags, His-tag and GST-tag.	-	33
Host: E. coli	-	26
Subcellular Location: Cell membrane.	-	22
	-	18
Cytoplasm. Early endosome. Recycling endosome.		10
Purity: >95%		14
Endotoxin Level: <1.0EU per 1µg		
(determined by the LAL method).		10
Formulation: Supplied as lyophilized form in PBS, pH7.4, 15% SDS-PAGE		
containing 5% trehalose, 0.01% sarcosyl.		
Predicted isoelectric point: 6.0		
Predicted Molecular Mass: 43.8kDa		
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.



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

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

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 GDVKLTQSMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSRIA YSKDFETLKV DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK KRIEAIPQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD GSTSGSGHHH HHHSAGLVPR GS TA I G M K E T AA A K F E R O H M DS P D L G T G G G SG I E G R G S M G YR G S E F - LT G S T PMQVVTCLTR DSYLTHCFLQ HLMVVLSSLE RTPSPEPVDK DFYSEFGNKT TGKMENYELI HSSRVKFTYP SEEEIGDLTF TVAQKMAEPE KAPALS