

YBA555Hu01 100µg

#### Recombinant Nicotinamide Adenine Dinucleotide Phosphate Oxidase 5 (NOX5) 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]	kDa 70		
Residues: Ser456~Phe765	44	-	-
Tags: Two N-terminal Tags, His-tag and T7-tag	33		-
Accession: Q96PH1	26	-	
Host: <i>E. coli</i>	22	-	
Subcellular Location: Membrane. Multi-pass	18		
membrane protein. Endoplasmic reticulum.	2.557.0		
Purity: >95%	14	-	
Endotoxin Level: <1.0EU per 1 $\mu$ g (determined by the LAL	10	2	
method).	10		
Formulation: Supplied as lyophilized form in PBS,			
pH7.4, containing 5% trehalose, 0.01% sarcosyl.			
Predicted isoelectric point: 9.8			
Predicted Molecular Mass: 39.6kDa			
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.



## [ 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.

# [<u>SEQUENCES</u>]

The sequence of the target protein is listed below.

SK V T H LL IK R P P FF H YR P G D Y LY L N IP T IA RY EW H PF T IS S A P E Q KD T I WL H IR S QGQWTNRLYE SFKASDPLGR GSKRLSRSVT MRKSQRSSKG SEILLEKHKF CNIKCYIDGP YGTPTRRIFA SEHAVLIGAG IGITPFASIL QSIMYRHQKR KHTCPSCQHS WIEGVQDNMK LHKVDFIWIN RDQRSFEWFV SLLTKLEMDQ AEEAQYGRFL ELHMYMTSAL GKNDMKAIGL QMALDLLANK EKKDSITGLQ TRTQPGRPDW SKVFQKVAAE KKGKVQVFFC GSPALAKVLK GHCEKFGFRF FQENF