



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

YB91183Hu01

Nucleoporin 188kDa (NUP188)

Organism: Homo sapiens (Human)

Instruction manual

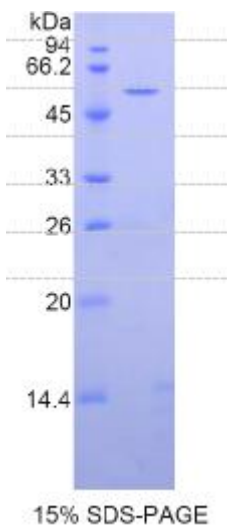
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NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

5th Edition (Revised in January, 2013)

[DESCRIPTION]

Protein Names: Nucleoporin 188kDa

Human NUP188



Synonyms: NUP188, KIAA0169

Species: Human

Size: 100μg

Source: *Escherichia coli*-derived

Subcellular Location: Nucleus, nuclear pore complex.

[PROPERTIES]



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Residues: Thr540~Gly782 (Accession # Q5SRE5),

with two N-terminal Tags, His-tag and GST-tag.

Grade & Purity: >95%, 55kDa as determined by

SDS-PAGE reducing conditions.

Formulation: Supplied as lyophilized form in PBS, pH

7.4, containing 5% sucrose, 0.01% sarcosyl.

Endotoxin Level: <1.0 EU per 1μg (determined by

the LAL method).

Applications: SDS-PAGE; WB; ELISA; IP.

(May be suitable for use in other assays to be determined by the end user.)

Predicted Molecular Mass: 53.8kDa

Predicted isoelectric point: 6.4



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[PREPARATION]

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.

MRNKKFELGL EFPNLPYYID GDVKLTQSMA IIRYADKHN MLGGCPKERA EISMLEGAVL
DIRYGVSRIA YSKDFETLKV DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD
VVLYMDPMCL DAFPKLVCFK KRIEAIQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD
GSTSGSGHHHHHSAGLVPRGSTAIGMKETA AAKFERQHMDSPDLGTLEV
LFQGPLGSEF- T LFTCEIEMLL HVVSTADVIQ HCQRVKPIID LVHKVISTDL SIADCLLPIT
SRIYMLLQRL TTVISPPVDV IASCVNCLTV LAARNPAKVW TDLRHTGFLP FVAHPVSSLS
QMISAEGMNA GGYGNLLMNS EQPQGEYGV T IAFLRLITTL VKGQLGSTQS QGLVPCVMFV
LKEMLP SYHK WRYNSHG VRE QIGCLILELI HAILNLCHET DLHSSHTPSL QFLCICSLAY
TEAGQTVINI MG