

RPU51556 Human 10µg Recombinant Hyaluronan Binding Protein 2 (HABP2) Organism Species: *Homo sapiens (Human) Instruction manual*

FOR RESEARCH USE ONLY

NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

12th Edition (Revised in Aug, 2016)



[PROPERTIES] Source: Prokaryotic expression Host: E.coli Residues: Phe24~Phe560 Tags: N-terminal His Tag Subcellular Location: Secreted **Purity:** > 90% Traits: Freeze-dried powder Buffer formulation: 100mMNaHCO₃, 500mMNaCl, pH8.3, containing 1mM EDTA, 1mM DTT, 0.01% SKL, 5% Trehalose and Proclin300. Original Concentration: 200µg/mL Applications: Positive Control; Immunogen; SDS-PAGE; WB. (May be suitable for use in other assays to be determined by the end user.) Predicted isoelectric point: 6.0 Predicted Molecular Mass: 63.9kDa Accurate Molecular Mass: 70kDa as determined by SDS-PAGE reducing conditions. Phenomenon explanation: The possible reasons that the actual band size differs from the predicted are as follows: 1. Splice variants: Alternative splicing may create different sized proteins from the same gene. 2. Relative charge: The composition of amino acids may affects the charge of the protein. 3. Post-translational modification: Phosphorylation, glycosylation, methylation etc. 4. Post-translation cleavage: Many proteins are synthesized as pro-proteins, and then cleaved to give the active form. 5. Polymerization of the target protein: Dimerization, multimerization etc.

[USAGE]

Reconstitute in 100mM NaHCO3, 500mM NaCl (pH8.3) to a concentration of 0.1-1.0 mg/mL. Do not vortex.



[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. 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. The loss rate is less than 5% within the

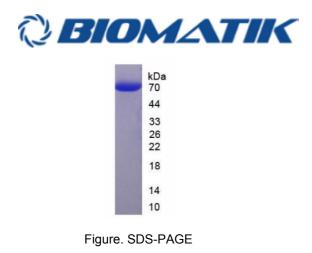
expiration date under appropriate storage condition.

[<u>SEQUENCE</u>]

		FSLMSLL	ESLDPDWTPD	QYDYSYEDYN
QEENTSSTLT	HAENPDWYYT	EDQADPCQPN	PCEHGGDCLV	HGSTFTCSCL
APFSGNKCQK	VQNTCKDNPC	GRGQCLITQS	PPYYRCVCKH	PYTGPSCSQV
VPVCRPNPCQ	NGATCSRHKR	RSKFTCACPD	QFKGKFCEIG	SDDCYVGDGY
SYRGKMNRTV	NQHACLYWNS	HLLLQENYNM	FMEDAETHGI	GEHNFCRNPD
ADEKPWCFIK	VTNDKVKWEY	CDVSACSAQD	VAYPEESPTE	PSTKLPGFDS
CGKTEIAERK	IKRIYGGFKS	TAGKHPWQAS	LQSSLPLTIS	MPQGHFCGGA
LIHPCWVLTA	AHCTDIKTRH	LKVVLGDQDL	KKEEFHEQSF	RVEKIFKYSH
YNERDEIPHN	DIALLKLKPV	DGHCALESKY	VKTVCLPDGS	FPSGSECHIS
GWGVTETGKG	SRQLLDAKVK	LIANTLCNSR	QLYDHMIDDS	MICAGNLQKP
GQDTCQGDSG	GPLTCEKDGT	YYYYGIVSWG	LECGKRPGVY	TQVTKFLNWI
KATIKSESGF				

[IDENTIFICATION_]

Figure. Gene Sequencing (Extract)



[IMPORTANT NOTE]

The kit is designed for research use only, we will not be responsible for any issue if the kit was used in clinical diagnostic or any other procedures.