## 32-13189: CYTIP Human

Cytohesin 1 Interacting Protein, Cytohesin Binder And Regulator, PSCDBP, Pleckstrin Homology, Sec7 And

## Alternative Name:

 Coiled-Coil Domains, Binding Protein,Pleckstrin Homology Sec7 And Coiled-Coil Domains-Binding Protein, Cytohesin-Associated Scaffolding Protein, Cytohesin Binding Protein HE, Cytohesin-Binding Protein HE, Cbp HE, CASP, CYBR, HE Pleckstrin Homology, Sec7 And Coiled/Coil Domains, Binding Protein, Cytohesin-1 Interacting Protein, Cytohesin-Interacting Protein, CYTHIP, B3-1, CYTIP.
## Description

Source: E.coli.
Sterile Filtered colorless solution.
Cytohesin-interacting protein, also known as CYTIP contains 2 leucine zipper domains as well as a putative C-terminal nuclear targeting signal. However, CYTIP does not contain any hydrophobic regions. In addition, CYTIP is expressed weakly in resting NK and T cells. It also modulates the activation of ARF genes by CYTH1.
CYTIP Human Recombinant produced in E.Coli is a single, non-glycosylated polypeptide chain containing 384 amino acids (1-359 a.a) and having a molecular mass of 42.6 kDa .CYTIP is fused to a 25 amino acid His-tag at N-terminus \& purified by proprietary chromatographic techniques.

## Product Info

## Amount:

Purification : Content :

## Storage condition :

Amino Acid :
$5 \mu \mathrm{~g} / 20 \mu \mathrm{~g}$
Greater than $85 \%$ as determined by SDS-PAGE.
CYTIP protein solution ( $1 \mathrm{mg} / \mathrm{ml}$ ) containing 20 mM Tris-HCl (pH 8.0) and $10 \%$ glycerol.
Store at $4^{\circ} \mathrm{C}$ if entire vial will be used within $2-4$ weeks. Store, frozen at $-20^{\circ} \mathrm{C}$ for longer periods of time. For long term storage it is recommended to add a carrier protein $(0.1 \% \mathrm{HSA}$ or BSA).Avoid multiple freeze-thaw cycles.
MGSSHHHHHH SSGLVPPGSH MGSEFMSLQR LLQHSSNGNL ADFCAGPAYS SYSTLTGSLT MDDNRRIQML ADTVATLPRG RKQLALTRSS SLSDFSWSQR KLVTVEKQDN ETFGFEIQSY RPQNQNACSS EMFTLICKIQ EDSPAHCAGL QAGDVLANIN GVSTEGFTYK QVVDLIRSSG NLLTIETLNG TMILKRTELE AKLQVLKQTL KQKWVEYRSL QLQEHRLLHG DAANCPSLEN MDLDELSLFG PLPGPGPALV DRNRLSSESS CKSWLSSMTM DSEDGYQTCV SEDSSRGAFS RQTSTDDECF IPKEGDDFLR RSSSRRNRSI SNTSSGSMSP LWEGNLSSMF GTLPRKSRKG SVRKQLLKFI PGLHRAVEEE ESRF

