

32-3373: C1QBP Recombinant Protein

Alternative Name : p32,HABP1,gC1Qr,GC1QBP,SF2p32,gC1Q-R,Complement component 1 Q subcomponent-binding protein mitochondrial,Glycoprotein gC1qBP,C1qBP,GC1q-R protein,Hyaluronan-binding protein 1,Mitochondrial matrix protein p32,p33,C1QBP.

Description

Source : Escherichia Coli. C1QBP Human Recombinant produced in E.Coli is a single, non-glycosylated polypeptide chain containing 210 amino acids and having a molecular mass of 23.9 kDa. C1QBP having the accession number of NP_001203 binds to the globular 'heads' of c1q thus inhibiting c1 activation. C1QBP interacts with a wide range of ligands and is implicated in cell signaling. C1QBP associates with C1r and C1s in order to yield the first component of the serum complement system. C1QBP protein has been identified as the p32 subunit of pre-mRNA splicing factor SF2, as well as a hyaluronic acid-binding protein.C1QBP is a new marker of tumor cells and tumor-associated macrophages/myeloid cells in hypoxic/metabolically deprived areas of tumors.Mitochondrial C1QBP is a critical mediator of p14ARF-induced apoptosis. C1QBP functions as a chemotactic factor for immature dendritic cells, and migration is mediated through ligation of both C1QBP and cC1qR/CR.C1QBP overexpression successfully blocks mRNA accumulation from the adenovirus major late transcription unit (MLTU) and stimulates RNA polymerase II carboxy-terminal domain phosphorylation in virus-infected cells.C1QBP binds with Hepacivirus core protein on CD8+ and CD4+ positive t-cells and inactivates Ick and akt.

Product Info

Amount : 25 µg
Purification : Greater than 95.0% as determined by:(a) Analysis by RP-HPLC.(b) Analysis by SDS-PAGE.
Content : The C1QBP protein solution contains 20mM Tris-HCl pH7.5, 20% glycerol and 1mM DTT.
Storage condition : Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA).Avoid multiple freeze-thaw cycles.
Amino Acid : MLHTDGDKAF VDFLSDEIKE ERKIQKHKTL PKMSGGWELE LNGTEAKLVR KVAGEKITVT FNINNSIPPT FDGEEEPSQG QKVEEQEPEL TSTPNFVVEV IKNDDGKKAL VLDCHYPEDE VGQEDEAESD IFSIREVSFQ STGESEWKDT NYTLNTDSL D WALYDHLMDF LADRGVDNTF ADELVELSTA LEHQEYITFL EDLKSFKVKSQ.