## 32-13147: CFB Human, Sf9

## Alternative <br> Name :

CFB, AHUS4, ARMD14, BF, BFD, CFAB, CFBD, FB, FBI12, GBG, H2-Bf, PBF2, Complement Factor B, BFactor, Properdin, Properdin Factor B, C3/C5 Convertase, EC 3.4.21.47, PBF2,Glycine-Rich BetaGlycoprotein, Glycine-Rich Beta Glycoprotein, C3 Proaccelerator, C3 Proactivator, EC 3.4.2, FBI12, H2Bf.

## Description

Source: Sf9, Baculovirus cells.
Sterile Filtered colorless solution.
Complement Factor B, also known as CFB, encodes complement factor B which is a component of the alternative pathway of complement activation. Factor B circulates in the blood as a single chain polypeptide. Once the alternative pathway is activated it is cleaved by complement factor D yielding the noncatalytic chain Ba and the catalytic subunit Bb . The active subunit Bb is a serine protease which connects with C 3 b to form the alternative pathway C 3 convertase. Also, Bb is involved in the proliferation of preactivated B lymphocytes, while Ba inhibits their proliferation.
CFB Human Recombinant produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain containing 245 amino acids (26-259a.a.) and having a molecular mass of 27.3 kDa (Molecular size on SDS-PAGE will appear at approximately 28-40 $\mathrm{kDa})$. CFB is expressed with a 11 amino acid His tag at C-Terminus and purified by proprietary chromatographic techniques.

## Product Info

## Amount : <br> Purification : Content :

## Storage condition :

Amino Acid :
$2 \mu \mathrm{~g} / 10 \mu \mathrm{~g}$
Greater than $95.0 \%$ as determined by SDS-PAGE.
CFB protein solution ( $0.5 \mathrm{mg} / \mathrm{ml}$ ) contains Phosphate Buffered Saline ( pH 7.4 ) 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.
ADPEFTPWSL ARPQGSCSLE GVEIKGGSFR LLQEGQALEY VCPSGFYPYP VQTRTCRSTG SWSTLKTQDQ KTVRKAECRA IHCPRPHDFE NGEYWPRSPY YNVSDEISFH CYDGYTLRGS ANRTCQVNGR WSGQTAICDN GAGYCSNPGI PIGTRKVGSQ YRLEDSVTYH CSRGLTLRGS QRRTCQEGGS WSGTEPSCQD SFMYDTPQEV AEAFLSSLTE TIEGVDAEDG HGPGEQQKRH HHHHH.

