

32-13257: Hemopexin Human, Sf9

Alternative Name : Hemopexin, Beta-1B-Glycoprotein, HX, Beta-1B-glycoprotein.

Description

Source: Sf9, Baculovirus cells.

Sterile filtered colorless solution.

Hemopexin (or haemopexin) is a plasma protein that binds heme with the highest affinity of any known protein. Hemopexin is generally expressed in liver, and belongs to acute phase reactants, the synthesis of which is induced after inflammation. Heme is potentially very toxic because of its ability to intercalate into lipid membrane and to generate hydroxyl radicals. Hemopexin's function of scavenging the heme released or lost by the turnover of heme proteins such as hemoglobin defends the body from the oxidative damage that free heme can cause. Additionally, hemopexin discharges its bound ligand for internalisation upon interacting with a specific receptor located on the surface of liver cells. This hemopexin function is in order to preserve the body's iron. Hemopexin's levels in the serum are an indication of how much heme is present in the blood. Low Hemopexin levels show that there is a lot of it in the serum. For that reason, low hemopexin levels indicate that there has been consid

Hemopexin produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain containing 448 amino acids (24-462a.a.) and having a molecular mass of 50.4kDa. (Molecular size on SDS-PAGE will appear at approximately 50-70kDa). Hemopexin is expressed with a 9 amino acid His tag at C-Terminus and purified by proprietary chromatographic techniques.

Product Info

Amount : 2 µg / 10 µg

Purification : Greater than 90.0% as determined by SDS-PAGE.

Content : Hemopexin protein solution (0.5mg/ml) contains 10% glycerol & Phosphate Buffered Saline (pH 7.4).

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 : ADPTPLPPTS AHGNVAEGET KPDPDVTERC SDGWSFDATT LDDNGTMLFF KGEFVWKSHK
WDRELISERW KNFPSPVDAA FRQGHNSVFL IKGDKVWVYP PEKKEKGYPK LLQDEFPGIP SPLDAAVECH
RGECQAEGVL FFQGDREWFV DLATGTMKER SWPAVGNCS ALRWLGYYC FQGNQFLRFD
PVRGEVPPRY PRDVRDYFMP CPGRGHGHRN GTGHGNSTHH GPEYMRCSPH LVLSALTSDN
HGATYAFSGT HYWRLDTSRD GWHSWPIAHQ WPQGPSAVDA AFSWEEKLYL VQGTQVYVFL
TKGGYTLVSG YPKRLEKEVG TPHGIILDSV DAAFCPGSS RLHIMAGRRL WWLDLKSGAQ ATWTLPWPH
EKVDGALCME KSLGPNSCSA NGPGLYLIHG PNLYCYS DVE KLNAAKALPQ PQNVTSLLGC THHHHHHH.