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## 32-13293: LCP2 Human

Alternative Name:

Lymphocyte Cytosolic Protein 2, SH2 Domain-Containing Leukocyte Protein Of 76 KDa, 76 KDa Tyrosine Phosphoprotein, SLP-76 Tyrosine Phosphoprotein, SLP76, Lymphocyte Cytosolic Protein 2 (SH2 Domain Containing Leukocyte Protein Of 76kDa), Lymphocyte Cytosolic Protein 2 (SH2 Domain-Containing Leukocyte Protein Of 76kD), SH2 Domain-Containing Leukocyte Protein Of 76Kd, SLP-76, LCP2.

## **Description**

Source: Sf9, Baculovirus cells. Sterile Filtered colorless solution.

LCP2 (lymphocyte cytosolic protein 2) is an intracellular docking protein with a single SH2 recognition domain. LCP2 is implicated in T-cell antigen receptor mediated signaling. Moreover, LCP2 is phosphorylated by ZAP70 which leads to NF-AT as well as IL2 gene activation. B Cell Linker Protein Deficiency is one of the associated diseases with LCP2.

LCP2 produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain (1-533 a.a.) and fused to a 6 aa His Tag at C-terminus containing a total of 542 amino acids and having a molecular mass of 61.2kDa.LCP2Â is purified by proprietary chromatographic techniques.

## **Product Info**

Amount:  $1 \mu g / 5 \mu g$ 

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

**Content:** LCP2 protein solution (0.25mg/ml) contains 10% glycerol & Phosphate buffered saline (pH7.4).

Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods

**Storage condition:** 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: ADPMALRNVP FRSEVLGWDP DSLADYFKKL NYKDCEKAVK KYHIDGARFL NLTENDIQKF PKLRVPILSK

LSQEINKNEE RRSIFTRKPQ VPRFPEETES HEEDNGGWSS FEEDDYESPN DDQDGEDDGD YESPNEEEEA PVEDDADYEP PPSNDEEALQ NSILPAKPFP NSNSMYIDRP PSGKTPQQPP VPPQRPMAAL PPPPAGRNHS PLPPPQTNHE EPSRSRNHKT AKLPAPSIDR STKPPLDRSL APFDREPFTL GKKPPFSDKP SIPAGRSLGE HLPKIQKPPL PPTTERHERS SPLPGKKPPV PKHGWGPDRR ENDEDDVHQR PLPQPALLPM SSNTFPSRST KPSPMNPLPS SHMPGAFSES NSSFPQSASL PPYFSQGPSN RPPIRAEGRN FPLPLPNKPR PPSPAEEENS LNEEWYVSYI TRPEAEAALR KINODGTFLV RDSSKKTTTN PYVLMVLYKD KVYNIQIRYO KESQVYLLGT

GLRGKEDFLS VSDIIDYFRK MPLLLIDGKN RGSRYQCTLT HAAGYPHHHHÂ HH