

## 32-4459: Recombinant Human PIN2-Interacting Protein 1

**Alternative Name** PINX1,LPTL,LPTS,MGC8850,FLJ20565,Pin2-interacting protein X1,TRF1-interacting protein 1,Liver-related putative tumor suppressor,Protein 67-11-3.

### Description

Source : Escherichia Coli. Recombinant Human PINX1 produced in E.Coli is a single, non-glycosylated polypeptide chain containing 348 amino acids (1-328 a.a) and having a molecular mass of 39.1 kDa. PINX1 is fused to 20 amino acid His Tag at N-terminus and purified by proprietary chromatographic techniques. PINX1 is a common expressed protein that localizes to nucleoli and telomere speckles. PINX1 contains a Telomerase Inhibiting Domain that is caable of binding MCRS1, TERT and TRF1. PINX1 has been shown to be a potent telomerase inhibitor and putative tumor suppressor. PINX1 is recruited to chromosome periphery by Nucleolin, their complex is necessary for faithful chromosome congression. PINX1 regulates the nucleolar accumulation and telomeric association of TRF1. PINX1 is involved in gastric cancer development. PINX1 expression is a sign of gastric cancer development. Constitutive expression of PINX1 attributes to telomere maintenance by telomerase and tumorigenicity in cancer cells.

### Product Info

<b>Amount :</b>	20 µg
<b>Purification :</b>	Greater than 95.0% as determined by analysis by SDS-PAGE.
<b>Content :</b>	The PINX1 protein contains 20mM Tris-HCl buffer pH-8, 1mM DTT and 10% glycerol.
<b>Storage condition :</b>	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.
<b>Amino Acid :</b>	MGSSHHHHHH SSGLVPRGSH MSMLAERRRK QKWAVDPQNT AWSNDDSKFG QRMLEKMGWS KGKGLGAQEH GATDHIKQVQ KNNHLGLGAT INNEDNWIAH QDDFNQLLAE LNTCHGQETT DSSDKKEKKS FSLEEKSKIS KNRVHYMKFT KGKDLSSRSK TDLDCIFGKR QSKKTPEGDA SPSTPEENET TTTSAFTIQE YFAKRMAALK NKPQVPVPGS DISETQVERK RGKKINKEAT GKDVESYLQP KAKRHTEGKP ERAEAQERVA KKSAPAEQ LRGPCWDQSS KASAQDAGDH VQPPEGRDFT LKPKKRRGKK KLQKPVEIAE DATLEETLVK KKKKKDSK.