

## 32-13804: THBS4 Mouse

<b>Format :</b>	The THBS4 solution (0.25mg/1ml) contains phosphate buffered saline (pH7.4), 0.1mM PMSF and 20% glycerol.
<b>Alternative Name :</b>	THBS4, Thbs4, Thbs-4, Thrombospondin-4, Thrombospondin4, thrombospondin 4, Tsp4, TSP-4, TSP, TSP4, TS.

### Description

Source: HEK293 Cells.

Physical Appearance: Sterile filtered colorless solution.

Biological Activity: null

Thrombospondin-4 (THBS4) is a member of the thrombospondin protein family. This family members are adhesive glycoproteins that mediate cell-to-cell and cell-to-matrix interactions. THBS4 forms a pentamer and binds to heparin and calcium. Among them THBS4 binds a variety of matrix proteins including Collagens I, II, III, V, Laminin-1, Matrilin-2 and Fibronectin. THBS4 is up-regulated in the spinal cord following peripheral nerve injury where it contributes to presynaptic hypersensitivity and hyperalgesia and is also up-regulated in muscle following denervation.

THBS4 Mouse Recombinant produced in HEK293 Cells is a single, glycosylated polypeptide chain containing 943 amino acids (27-963a.a) and having a molecular mass of 104.3 kDa. THBS4 is expressed with a 6 amino acid His tag at C-Terminus and purified by proprietary chromatographic techniques.

### Product Info

<b>Amount :</b>	10 µg / 2 µg
<b>Purification :</b>	Greater than 85.0% as determined by SDS-PAGE.
<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>	QATPQVFDLL PSSSQRLNPS ALQPVLTDP LHEVYLISTF KLQSKSSATI FGLYSSSDNS KYFEFTVMGR LNKAILRYLK NDGKIHLVVF NNLQLADGRR HRVLLRLSNL QRGDGSVELY LDCAQADSVR NLPRAFSGLT QNPESIELRT FQRKPQDFLE ELKLVVRGSL FQVASLQDCF LQSEPLAAT STGDFNRQFL GQMTQLNQLL GEVKDLLRQQ VKETSFLRNT IAECQACGPL SFQSPTNTL VPIAPPAPT RPTRHCDSSP CFRGVRCTDT RDGFQCGPCP DGYTGNGITC SDVDECKYHP CYPGVRVNL APGFRCDACP VGFTGPMVQG VGINFATNK QVCTDVDECQ NGACVLNSIC INTLGSYRCG PCKPGYTG DQ TRGCKTERSC RNPEQNPCSV HAQCIEERQG DVTCVCGVGW AGDGYVCGKD VDIDSYDDEE LPCSARNCKK DNCKYVPNSG QEDADRDGIG DACDEDADGD GILNEQDNCV LTHNIDQRNS DKDIFGDACD NCRMVLNNDQ KDTDGDGRGD ACDDMDGDG IKNILDNCPR VPNRDQQDRD GDDVGDACDS CPDVSNPNQS DVDNDLVGDS CDTNQDSGDG GHQDSTDNCP TVINSSQLDT DKDYGIGDEC DDDNDNGIPD LVPPGPDNCR LVPNPAQEDS NNDGVGDICE ADFDQDQVID HIDVCPENAE ITLTDFRAYQ TVVLDPEGDA QIDPNWVVLN QGMEIVQTMN SDPGLAVGYT AFNGVDFEGT FHVNTQTDDD YAGFIFGYQD SSSFYVVMWK QTEQTYWQAT PFRAVAEPI QLKAVKSKTG PGEHLRNSLW HTGDTSDQVR LLWKDSRVNG WKDKVSYRWF LQHRPQVGYI RVRFYEGSEL VADSGVTIDT TMRGGRLGVF CFSQENIIWS NLKYRCNDTI PEDFQEFQQTQ SFDRLDNHHH HHH