

Recombinant Human TRAK1 293 Cell Lysate

Cat. No. TRAK1-813HCL **Lot. No.** (See product label)

SPECIFICATION

Product Overview

Species	Human
Source	HEK293
Description	Antigen standard for trafficking protein, kinesin binding 1 (TRAK1), transcript variant 2 is a lysate prepared from HEK293T cells transiently transfected with a TrueORF gene-carrying pCMV plasmid and then lysed in RIPA Buffer. Protein concentration was determined using a colorimetric assay. The antigen control carries a C-terminal Myc/DDK tag for detection.
Components	This product includes 3 vials: 1 vial of gene-specific cell lysate, 1 vial of control vector cell lysate, and 1 vial of loading buffer. Each lysate vial contains 0.1 mg lysate in 0.1 ml (1 mg/ml) of RIPA Buffer (50 mM Tris-HCl pH7.5, 250 mM NaCl, 5 mM EDTA, 50 mM NaF, 1% NP40). The loading buffer vial contains 0.5 ml 2X SDS Loading Buffer (125 mM Tris-Cl, pH6.8, 10% glycerol, 4% SDS, 0.002% Bromophenol blue, 5% beta-mercaptoethanol).
Size	0.1 mg
Storage Instruction	Store at -80°C. Minimize freeze-thaw cycles. After addition of 2X SDS Loading Buffer, the lysates can be stored at -20°C. Product is guaranteed 6 months from the date of shipment.

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Applications

ELISA, WB, IP. WB: Mix equal volume of lysates with 2X SDS Loading Buffer. Boil the mixture for 10 min before loading (for membrane protein lysates, incubate the mixture at room temperature for 30 min). Load 5 ug lysate per lane.

GENE INFORMATION
Gene Name

TRAK1 trafficking protein, kinesin binding 1 [Homo sapiens]

Official Symbol

TRAK1

Synonyms

TRAK1; trafficking protein, kinesin binding 1; trafficking kinesin-binding protein 1; KIAA1042; MILT1; milton homolog 1 (Drosophila); O linked N acetylglucosamine transferase interacting protein 106; OGT(O Glc NAc transferase) interacting protein 106 KDa; OIP106; milton homolog 1; 106 kDa O-GlcNAc transferase-interacting protein; OGT(O-Glc-NAc transferase)-interacting protein 106 KDa; O-linked N-acetylglucosamine transferase interacting protein 106;

Gene ID

22906

mRNA Refseq

NM_001042646

Protein Refseq

NP_001036111

MIM

608112

UniProt ID

Q9UPV9

**Chromosome
Location**

3p22.1

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