

Product datasheet for TP501113

OriGene Technologies, Inc.

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Mal (NM 010762) Mouse Recombinant Protein

Product data:

Product Type: Recombinant Proteins

Description: Purified recombinant protein of Mouse myelin and lymphocyte protein, T cell differentiation

protein (Mal), with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug

Species: Mouse

Expression Host: HEK293T

Expression cDNA Clone >MR201113 protein sequence or AA Sequence: Red=Cloning site Green=Tags(s)

MAPAAASGGSTLPSGFSVFTTFPDLLFVCEFVFGGLVWILIASSLVPLPLAQGWVMFVSVFCFVATTSLM ILYIIGTHGGETSWITLDAAYHCVAALFYLSASVLEALATISMFDGFTYKHYHENIAAVVFAYVVTLIYV

VHAVFSLIRWKSS

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-MYC/DDK

Predicted MW: 16.6 kDa

Concentration: >0.05 μg/μL as determined by microplate BCA method

Purity: > 80% as determined by SDS-PAGE and Coomassie blue staining

Buffer: 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol

Note: For testing in cell culture applications, please filter before use. Note that you may experience

some loss of protein during the filtration process.

Storage: Store at -80°C after receiving vials.

Stability: Stable for 12 months from the date of receipt of the product under proper storage and

handling conditions. Avoid repeated freeze-thaw cycles.

RefSeq: NP 034892

 Locus ID:
 17153

 UniProt ID:
 009198

 RefSeq Size:
 2792

 Cytogenetics:
 2 F1





Mal (NM_010762) Mouse Recombinant Protein - TP501113

RefSeq ORF: 462

Synonyms: Mpv17; VI; Vip17

Summary: This gene encodes a highly hydrophobic integral membrane protein belonging to the MAL

family of proteolipids. The encoded protein has been localized to the endoplasmic reticulum of T-cells and is a candidate linker protein in T-cell signal transduction. In addition, this proteolipid is localized in compact myelin of cells in the nervous system and has been implicated in myelin biogenesis and/or function. The protein plays a role in the formation, stabilization and maintenance of glycosphingolipid-enriched membrane microdomains. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2010]