

# Recombinant Human Tripartite motif-containing protein 72(TRIM72)

Catalog Number: CSB-MP744394HU



<b>Product Name:</b>	Recombinant Human Tripartite motif-containing protein 72(TRIM72)
<b>Alternative names:</b>	Mitsugumin-53
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<b>Relevance :</b>	Muscle-specific protein that plays a central role in cell membrane repair by nucleating the assembly of the repair machinery at injury sites. Specifically binds phosphatidylserine. Acts as a sensor of oxidation: upon membrane damage, entry of extracellular oxidative environment results in disulfide bond formation and homooligomerization at the injury site. This oligomerization acts as a nucleation site for recruitment of TRIM72-containing vesicles to the injury site, leading to membrane patch formation. Probably acts upstream of the $Ca^{2+}$ -dependent membrane resealing process. Required for transport of DYSF to sites of cell injury during repair patch formation. Regulates membrane budding and exocytosis. May be involved in the regulation of the mobility of KCNB1-containing endocytic vesicles.
<b>Mol. Weight:</b>	31kd
<b>Tag</b>	His-tagged
<b>Source:</b>	Mammalian cell derived
<b>Image:</b>	
<b>Purity:</b>	>90%(SDS-PAGE)
<b>Storage Buffer:</b>	20mM Tris-Hcl, 0.5M Nacl,pH 8.0, 50% glycerol
<b>Storage :</b>	Store at -20℃, for extended storage, conserve at -20℃ or -80℃.
<b>Notes :</b>	Repeated freezing and thawing is not recommended. Store working aliquots at 4℃ for up to one week.
<b>AA sequence:</b>	MSAAPGLLHQELSCPLCLQLFDAPVTAECGHSFCRACLGRVAGEPAADGTVLCPCCQAPT RPQALSTNLQLARLVEGLAQVPQGHCEEHLDP LSIYCEQDRALVCGVCASLGSHRGHRL PAAEAHARLKTQLPQQKLQLQEACMRKEKSVAVLEHQLVEVEETVRQFRGAVGEQLGKMR VFLAALEGLSLDREAERV RGEAGVALRRELGSLNSYLEQLRQMEKVLEEVADKPQTFLMK YCLVTSRLQKILAESP PPARLDIQLPIIS
<b>References:</b>	"Complete sequencing and characterization of 21,243 full-length human cDNAs." Ota T., Suzuki Y., Nishikawa T., Otsuki T., Sugiyama T., Irie R., Wakamatsu A., Hayashi K., Sato H., Nagai K., Kimura K., Makita H., Sekine M., Obayashi M., Nishi T., Shibahara T., Tanaka T., Ishii S. Sugano S. Nat. Genet. 36:40-45(2004)