

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q0KL02	TRIO_MOUSE	Mus musculus	Triple functional domain protein	18.754664	NaN	S1627;S1632;S1633;T1824;S2282;S2458;S2462;S2634	22645316;33300544	MSGSSGGATAPAASSGPAAAAAAG SGCGGGAGEGAEAAKDLADIAAFF RSGFRKNDEMAMDVLPILKEKVAY LSGGRDKRGGPILTFPARSNHDRIR QEDLRRLISYLACIPSEEVCKRGFTVI VDMRGSKWDSIKPLLKILQESFPCCI HIALIIPDNFWQKQRTNFGSSKFEEF ETNMVSLEGLTKVVDPSQLTPEFDG CLEYNHEEWIEIRVAFEEYISNAAH MLSRLEELQDVLAKKELPDLEGAR NMIDEHSQKKKVIKAPIEDLDLEG QKLLQRIQSSDSFPKNSGSGNADL QNLLPKVSTMLDRLHSTRQHLHQM WHVRKCLKDQCFQLRLEFQDAEKM FDWITHNKGLFLNSYTEIGTSHPHA MELQTQHNHFAMNCMNVYVNNINRI MSVANRLVESGHYASQKIQIANQL EQEWKAFAAALDERSTLLDMSSIFH QKAEKYMSNVDSWCKACGEVDLPS ELQDLEDAIHQQGIYEHITLAYSEV SQDGKSLLDKLRPLTPGSSDSLTA ANYSKAVHHVLDVIHEVLHHQRQLE NIWQHRKVRLHQRLQLCVFQQDVQ QVLDWIENHGEAFLSKHTGVGKSL HRARALQKRHEDFEEVAQNITYTNA DKLLEAAEQLAQTGECDPPEIYQAA HQLEDRIQDFVRRVEQRKILLDMSV SFHTHVKELWTWLEELQKELLDDV YAESVEAVQDLIKRFGQQQQTTLQV TVNVIKEGEDLIQQLRDSAISNKTP HNSSINHIETVLQQLDEAQSQMEEL FQERKIKLELFLQLRIFERDAIDIISD LESWNDELSQQMNDFTDTELTIAE QRLQHHADKALTMNNLTFDVIHQG QDLLQYVNEVQASGVELLCDRDVD MATRVQDLLEFLHEKQQELDLAAE QHRKHLEQCVQLRHLQAEVKQVLG WIRNGESMLNAGLITASSLQAEQL QREHEQFQHAIEKTHQSALQVQQK AEAMLQANHYDMDMIRDCAEKVAS HWQQLMLKMEDRLKLVNASVAFYK TSEQVCSVLESLEQYKREEDWCGG ADKLGPNSETDHVTPMISKHLEQKE AFLKACTLARRNADVFLKYLHRNSV SMPGMVTHIKAPEQQVKNILNELFQ RENRLHYWTMRKRRLDQCQYVV FERSAKQALEWIHDNGEFYLSHTS TGSSIQHTQELLKEHEEFQITAKQTK ERVKLLIQLADGFCEKGHAHAAEIKK

CVTAVDKRYRDFSLRMEKYRTSLEK  
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QSRLPRV