

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O75962	TRIO_HUMAN	Homo sapiens	Triple functional domain protein	14.211856	S2291	S1627;S1632;S1633;T1824;S2282;S2455;S2459;S2631	38253038;37217939	MSGSSGGAAAPAASSGPAAAAASAAG SGCGGGAGEGAEAAKDLADIAAFF RSGFRKNDEMKAAMDVLPILKEKVAY LSGGRDKRGGPILTFPARSNHDRIR QEDLRRLISYLACIPSEEVCCKRGFTVI VDMRGSKWDSIKPLLKILQESFPCCI HVALIIPDNFWQKQRTNFGSSKFE FETNMVSLEGLTKVVDPSQLTPEFD GCLEYNHEEWIEIRVAFEDYISNATH MLSRLEELQDILAKKELPQDLEGAR NMIEEHSQKKKVIKAPIEDLDLEGQ KLLQRIQSSESFPPKNSGSGNADLQ NLLPKVSTMLDRLHSTRQHLHQMWW HVRKCLKDQCFQLRLFQDAEKMF DWITHNKGLFLNSYTEIGTSHPHAM ELQTQHNHFAMNCMNVYVNINRIM SVANRLVESGHYASQQIRQIASQLEQ EWKAFAAALDERSTLLDMSSIFHQK AEKYMSNVDSWCKACGEVDLPSEL QDLEDAIHQQGIYEHITLAYSEVSQ DGKSLDKLQRPLTPGSSDSLTAASA NYSKAVHHVLDVIHEVLHHQRQLE NIWQHRKVRLHQRLQLCVFQQDVQ QVLDWIENHGEAFLSKHTGVGKSL HRARALQKRHEDFEEVAQNTYTNA DKLLEAAEQLAQTGECDPPEIYQAA HQLEDRIQDFVRRVEQRKILLDMSV SFHTHVKELWTWLEELQKELDDV YAESVEAVQDLIKRFGQQQQTTLQV TVNVIKEGEDLIQQLRDSAISSNKTP HNSSINHIETVLOQLDEAQSQMEEL FQERKIKLELFLQLRIFERDAIDIISD LESWNDELSQQMNDFTDTELTIAE QRLQHHADKALTMNNLTFDVIHQG QDLLQYVNEVQASGVLLCDRDVD MATRVQDLLEFLHEKQQELDLAAE QHRKHLEQCQVLRHLQAEVKQVLG WIRNGESMLNAGLITASSLQAEQQL QREHEQFQHAIEKTHQSALQVQQK AEAMLQANHYDMDMIRDCAEKVAS HWQQLMLKMEDRLKLVNASVAFYK TSEQVCSVLESLEQEYKREEDWCGG ADKLGPNSETDHVTPMISKHLEQKE AFLKACTLARRNADVFLKYLHRNSV NMPGMVTHIKAPEQQVKNILNELFQ RENRLHYWTMRKRRLDQCQYVV FERSAKQALEWIHDNGEFYLSHTS TGSSIQHTQELLKEHEEFQITAKQTK ERVKLLIQLADGFCEKGAHAHAAEIKK

CVTAVDKRYRDFSLRMEKYRTSLEK  
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STQSNSESSSSSNISTMLVTHDYT  
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