

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P55072	TERA_HUMAN	Homo sapiens	Transitional endoplasmic reticulum ATPase	24.591228	S3	S3;S7;S13;S37;T436;S462;S702;S770;S775;S787;T805	23301498;27566657;30397120;35083852;26853435;33465208;35008409;33214551;34725712;30444036;32119511;22967762;35138101;34019948;26136220	MASGADSKGDDLSTAILKQKNRPNR LIVDEAINEDNSVVLSQPKMDELQ LFRGDTVLLKGGKRREAVCIVLSDDT CSDEKIRMNRVVRNLRVRLGDVIS IQPCPDVKYGKRIHVLPIDDTVEGITG NLFEVYLKPYFLEAYRPIRKGDIFLVR GGMRAVEFKVETDPSYCVAPDT VIHCEGEPIKREDEEESLNEVGYDDI GGCRKQLAQIKEMVELPLRHPALFK AIGVKPPRGILLYGPPGTGKTLIARAV ANETGAFFFLINGPEIMSKLAGESES NLRKAFEEAEKNAPAIIFIDELDAIAP KREKTHGEVERRIVSLLLTLMDGLK QRAHVIVMAATNRPN SIDPALRRFG RFDREVDIGIPDATGRLEILQIHTKN MKLADDVDLEQVANETHGHV GADL AALCSEALQAIRKKMDLIDLEDETI DAEVMNSLAVTMDDFRWALSQSNP SALRETVVEVPQVTWEDIGGLEDVK RELQELVQYPVEHPDKFLKFGMTPS KGVLFYGPPEGCKTLLAKAIANECQ ANFISIKGPELLTMWFGSEANVREI FDKARQAAPCVLFFDELDSIAKARG GNIGDGGGAADRVINQILTEMDGMS TKKNVFIIGATNRPDIIDPAILRPGRL DQLIYIPLPDEKSRVAILKANLRKSPV AKDVDLEFLAKMTNGFSGADLTEIC QRACKLAIRESIESEIRRERERQTNP SAMEVEEDDPVPEIRRDHFEEAMRF ARRSVSDNDIRKYEMFAQTLQQSRG FGSFRFPSGNQGGAGPSQSGGGGT GGSVYTEDNDDDLYG

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