

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
P55072	TERA_HUMAN	Homo sapiens	Transitional endoplasmic reticulum ATPase	21.162562	S3	S3;S7;S13;S37;T436;S462;S702;S770;S775;S787;T805	26136220;33214551;32119511;33465208;30444036;34019948;26853435;27566657;22967762;23301498;30397120	MASGADSKGDDLSTAILKQKNRPNR LIVDEAINEDNSVVLSQPKMDELQ LFRGDTVLLKGGKRREAVCIVLSDDT CSDEKIRMNRVVRNLRVRLGDVIS IQPCPDVKYGKRIHVLPIDDTVEGITG NLFEVYLKPYFLEAYRPIRKGDIPLVR GGMRAVEFKVETDPSYCVAPDT VIHCEGEPIKREDEEESLNEVGYDDI GGCRKQLAQIKEMVELPLRHPALFK AIGVKPPRGILLYGPPGTGKTLIARAV ANETGAFFFLINGPEIMSKLAGESES NLRKAFEEAEKNAPAIIFIDELDAIAP KREKTHGEVERRIVSLLLTLMDGLK QRAHVIVMAATNRPN SIDPALRRFG RFDREVDIGIPDATGRLEILQIHTKN MKLADDVDLEQVANETHGHVGADL AALCSEALQAIRKKMDLIDLEDETI DAEVMNSLAVTMDDFRWALSQSNP SALRETVVEVPQVTWEDIGGLEDVK RELQELVQYPVEHPDKFLKFGMTPS KGVLFYGPPEGCKTLLAKAIANECQ ANFISIKGPELLTMWFGGESEANVREI FDKARQAAPCVLFFDELDSIAKARG GNIGDGGGAADRVINQILTEMDGMS TKKNVFIIGATNRPDIIDPAILRPGRL DQLIYIPLPDEKSRVAILKANLRKSPV AKDVDLEFLAKMTNGFSGADLTEIC QRACKLAIRESIESEIRRERERQTNP SAMEVEEDDPVPEIRRDHFEEAMRF ARRSVSDNDIRKYEMFAQTLQQSRG FGSFRFPSGNQGGAGPSQGGGGT GGSVYTEDNDDDLYG

0