	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P23787	TERA_XENLA	Xenopus laevis	Transitional endoplasmic reticulum ATPase	26.754248	NaN	S3	18617508	MASGSDTKSDDLSTAILKQKSRPNR LIVDESINEDNSMVSLSQAKMDELQ LFRGDTVLLKGKKRREAVCIVLSDDT CSDEKIRMNRVVRNNLRVRLGDVIS IQPCPDVKYGKRVHVLPIDDTVEGIT GNLFEVYLKPYFLEAYRPIRKGDIFLV RGGMRAVEFKVVETDPSPYCIVAPD TVIHCEGEPIKREDEEESLNEVGYDD IGGCRKQLAQIKEMVELPLRHPALF KAIGVKPPRGILLYGPPGTGKTLIARA VANETGAFFFLINGPEIMSKLAGESE SNLRKAFEEAEKNAPAIIFIDELDAIA PKREKTHGEVERRIVSQLLTLMDGL KQRAHVIVMAATNRPNSIDPALRRF GRFDREVDIGIPDSTGRLEILQIHTK NMKLSDDVDLEQVANETHGHVGAD LAALCSEAALQAIRKKMDLIDLEDET IDAEVMNSLAVTMDDFRWALSQSN PSALRETVVEVPQVTWEDIGGLEDV KRELQELVQYPVEHPDKFLKFGMTP SKGVLFYGPPGCGKTLLAKAIANECQ ANFISIKGPELLTMWFGESEANVREI FDKARQAAPCVLFFDELDSIAKARG GNIGDGGGAADRVINQILTEMDGMS IKKNVFIIGATNRPDIIDPAILRPGRLD QLIYIPLPDEKSRMAILKANLRKSPV AKDVDVDFLAKMTNGFSGADLTEIC QRACKLAIRESIENEIRRERDRQTNP SAMEVEEDDPVPEIRRDHFEEAMRF ARRSVSDNDIRKYEMFAQTLQQSRG FGSFRFPAGGQSGAGPSPGAGGGSG GGHFTEEDDDLYG	None	None	None	None	None	None	None	None	None