

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|----------------|---|--------------|---------------|-----------------------|----------|--|
| P23787 | TERA_XENLA | Xenopus laevis | Transitional endoplasmic reticulum ATPase | 11.140708 | NaN | S3 | 18617508 | MASGSDTKSDDLSTAILKQKSRPNR LIVDESINEDNSMVLSQAKMDELQ LFRGDTVLLKGKKRREAVCIVLSDDT CSDEKIRMNRVVRNLRVRLGDVIS IQPCPDVKYGKRVHVLPIDDTVEGIT GNLFEVYLKPYFLEAYRPIRKGDIFLV RGGMRAVEFKVVETDPSPYCIVAPD TVIHCEGEPIKREDEEESLNEVGydd IGGCRKQLAQIKEMVELPLRHPALF KAIGVKPPRGILLYGPPGTGKTLIARA VANETGAFFFLINGPEIMSKLAGESE SNLRKAFEEAEKNAPAIIFIDELDAIA PKREKTHGEVERRIVSOLLTMDGL KQRAHVIVMAATNRPN SIDPALRRF GRFDREVDIGIPDSTGRLEILQIHTK NMKLSDDVDLEQVANETHGHVGD LAALCSEAALQAIRKKMDLIDLEDET IDAEVMNSLAVTMDDFRWALSQSN PSALRETVVEVPQVTWEDIGGLEDV KRELQELVQYPVEHPDKFLKFGMTP SKGVLFYGPFGCKTLLAKAIANECQ ANFISIKGPELLTMWFGSEANVREI FDKARQAAPCVLFFDELDSIAKARG GNIGDGGGAADRVINQILTEM DGMS IKKNVFIIGATNRPD IIDPAILRPGRLD QLIYIPLPDEKSRMAILKANLRKSPV AKDVDVDFLAKMTNGFSGADLTEIC QRACKLAIRESIENEIRREDRQTNP SAMEVEEDDPVPEIRRDHFEEAMRF ARRSVSDN DIRKYEMFAQTLQQSRG FGSFRFPAGGQSGAGPSPGAGGSGG GGHFTEEDDDLYG |

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