

| UniprotKB ID | Entry name | organism | full name | oglnacscore | oglnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|---|-------------|--------------|--|-------------------------------------|---|
| Q01853 | TERA_MOUSE | Mus musculus | Transitional endoplasmic reticulum ATPase | 14.863586 | NaN | S3;S7;S13;S37;T436;S462;S702;S770;S775;S787;T805 | 33300544;37453647;36288343;37507081 | MASGADSKGDDLSTAILKQKNRPNR LIVDEAINEDNSVVSLSQPKMDELQ LFRGDTVLLKGKKRREAVCIVLSDDT CSDEKIRMNRVVRNLRVRLGDVIS IQPCPDVKYGKRIHVLPIDDTVEGITG NLFEVYLKPYFLEAYRPIRKGDIPLVR GGMRAVEFKVETDPSYCVAPDT VIHCEGEPIKREDEEESLNEVGYDDI GGCRKQLAQIKEMVELPLRHPALFK AIGVKPPRGILLYGPPGTGKTLIARAV ANETGAFFFLINGPEIMSKLAGESES NLRKAFEEAEKNAPAIIFIDELDAIAP KREKTHGEVERRIVSLLLTLMDGLK QRAHVIVMAATNRPN SIDPALRRFG RFDREVDIGIPDATGRLEILQIHTKN MKLADDVDLEQVANETHGHVGADL AALCSEALQAIRKKMDLIDLEDETI DAEVMNSLAVTMDDFRWALSQSNP SALRETVVEVPQVTWEDIGGEDVK RELQELVQYPVEHPDKFLKFGMTPS KGVLFYGPPEGCKTLLAKAIANECQ ANFISIKGPELLTMWFGESEANVREI FDKARQAAPCVLFFDELDSIAKARG GNIGDGGGAADRVINQILTEMDGMS TKKNVFIIGATNRPDIIDPAILRPGRL DQLIYIPLPDEKSRVAILKANLRKSPV AKDVDLEFLAKMTNGFSGADLTEIC QRACKLAIRESIESEIRRERERQTNP SAMEVEEDDPVPEIRRDHFEEAMRF ARRSVSDNDIRKYEMFAQTLQQSRG FGSFRFPSGNQGGAGPSQSGGGGT GGSVYTEDNDDDLYG |

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