

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q64514	TPP2_MOUSE	Mus musculus	Tripeptidyl-peptidase 2	29.424336	S937;S938	S915	22645316;37507081;25153642;28528544;34418053	MATAATEEPFFHGLLPKKTGASS FLCRYPEYDGRGVLI AVLDTGVDPGA PGMQVTTD GKPKIIDIT TSGSDVN TATEVEPKDGEIIGLSGRVLKIPANW TNPLGKYHIGIKNGYDFYPKALKERI QKERKEKIWDPIHRVALAEACRKQE EFDIANNSSQANKLIKEELQSQVE LLNSFEKKYSDPGPVYDCLVWH DGE TWRACVDSNENGDLSKCAVLRNYK EAQEYSSFGTAEMLNYSVNIYDDGN LLSIVTSGGAHGTHVASIAAGHFPEE PERNGVAPGAQILSIKIGDTRLSTME TGTGLIRAMIEVINHKCDLVNYSYGE ATHWPNNGRICEVINEAVWKHNTIY VSSAGNNGPCLSTVGC PGGTTSSVI GVGAYVSPDMMVAEYSLREKLPAN QYTWSSRGPSADGALGVSISAPGGAI ASVPNWTLRGTQLMNGTSMSSPNA CGGIALVLSGLKANNVDYTVHSVRR ALENTAIKADNIEVFAQGHGIIQV DK AYDYLIQNTSFANRLGFTVTVGN NR GIYLRDPVQVAAPSDHGVGIEPVFPE NTENSEKISFQLHLALTSNSSWVQC PSHLELMNQCRHINIRVDP RGLREG LHYTEVCGYDIASP NAGPLFRVPITA VIAAKVNESSHYDLAFTDVHFKPGQI RRHFVEVPEGATWAEVTVCSCSSEV SAKFVLHAVQLVKQRAYSHEFYKF CSLPEKGTLIEAFPVLGGKAIEFCIAR WWASLSDVNIDYTISFHGIVCTAPQL NIHASEGINRFDVQSSLKYEDLAPCI TLKSWVQTLRPVNAKTRPLGSRDVL PNNRQLYEMVLTYSFHQPKSGEVTP SCPLLCELLYESEFDSQLWIIFDQNK RQMGSGDAYPHQYSLKLEKGDYTIR LQIRHEQISDLDR LKDLPFIVSHRLS NTLSLDIHENHSLALLGKKKSSSLTL PPKYNQPFVTSLPDDKIPKGAGPGC YLAGSLTSLKTELGKKAGQSAKRQ GKFKKDVIPVHYLLIPPPTKIKNGSK DKEKDSEKEKDLKEEFTEALRDLKI QWMTKLDSTDIYNELKETYPAYLPLY VARLHQLDAEKERMKRLNEIVDAAN AVISHIDQTALAVYIAMKTDP RPDA TIKNDMDKQKSTLIDALCRKGCALA DHLLHTQPHDGAAGDAEAKEEEG ESTMESLSETYWETTKWTDLFDTKV LIFAYKHALVNKMYGRGLKFATKLV EEKPTKENWKNCIQLMKLLGWTHC

