

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membr
Q5SSZ5	TENS3_MOUSE	Mus musculus	Tensin-3	31.528337	S745	T323;S332;S361;S370;S440;S516;S571;T632;S648;S687;S690;S769;T773;S804;S860;S894;S960;S1144;S1149;S1288;S1436	25153642;28528544;34418053	MEDSHELDLTYVTERIAVSPFASCS EESYLHSLQEVTRMLKCKKHGDNYLV LNLSEKRYDLTKLNPKIMDVGWPEL HAPPLDKMCTICKAQESWLNNDPQ HVVVHICRGGKGRIGVVISSYMHFT NVSASADQALDRFAMKKFYDDKISA LMEPSQKRYVQFSLGSLGAMKMN TSPFLHFVIMHGVPSFDTGGACRP FLKLYQAMQPVYTSGIYVNGSENPS RIRIAIEPAQLLKGDMVKCYHKKFR SATRDVIFRLQFHTGAVQGYLLFG KEELDACKDDRFDPYGKIELVFSAT PEKIQGSEHLYSDQGVTVDYNTADP LIRWDSYENMSADGEVLHTQGPD GSLYAKVRKKSASDTGIPSSPQGM TSSPDHGDHTLSVSSDSGHSTASAR TDKTEERLTPGARRGLSPQEAELD QLLSGFGLSDSASSHKDMTDMRSK YSGTRHVVPAQVHVNGDAALKDRET DILDDEMPHDLHSVDSLGLTSSSE GPOSTHLGPFTCLKSSQNSLLSDGF GNGVAEDHNGVLSPLDGLGVDTLY DRERMCGGREQKPLQPLLRKPSAPT PVQAYGQSNYSTQTWVRQQQMVAA HQYFASDGEARLGSRSTVDNTGLA QPPPHIPVTPNRGASSRVAVQGISN GPNPPDTQQLCPGKALQPRFODDRV TNGVHQEPNTGSSPGSFTLDIDQIE QLNRLILELDPTEFPIPTHLNALGISA VCPDGVGSLRCSGRLDSDVGGPRS PGRQDDPIGGRLRKLKLSIGQYDND ASQVTFKCGWKGAGVDPAPSLGFS SSPEDIKETVITAYPSDLNMDGRIPN SKESSMCLTPSFPVSPETPYVKTSPR YPPFSPPEQLSSPASLHKGREPRGC PEIISHTVGMSESPVGPKPTMLRAD MPATPNFQQVFASSCTVSSNGPGQR RESPPSAERQWVESPSTLTLGN SHPSESPLGTHEFCSSGKDSPLGPC FQSSQLQASFHSHSLSMSEPPQALP PAGSQTFLGFNTVTTATSVLPPGEDA GTLVNSHGTSPAPGTPLLTGAAD NGFLPHNFLTSPGASSHHSPGLQN QNVSLPGQPPLPEKKRASEGDRSLG SVSPSSSGFSSPHSGSTMSIPFPNVL PDFCKPSEVASPLPDSPNDKLVIVKF VQDTSKFWYKADISREQAIAMLKDK APGSFIVRDSHSFRGAYGLAMKVAT PPPSVLHLNKKAGDLSNELVRHFLI ECTPKGVRLKGCSENPYFGSLTALV CQHSITPLALPCKLLIPERDPLEEIAE NSPQTAANSAEELLKQGAACNVVYL NSVEMESLTGHQAVQKALSMTLVQ EPPPVSTVVHFKVSAQGITLTDNQRK LFFRRHYPVSSVIFCALDPQDRKWIK DGPSSKVFGFVARKQGSATDNVCHL FAEHDPEQPASAINFVSKVMIGSPK KI	False	False	2.712	3.356	1.416	1.11	0.705	1.765