

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5SSZ5	TENS3_MOUSE	Mus musculus	Tensin-3	16.895078	S745	T323;S332;S361;S370;S440;S516;S571;T632;S648;S687;S690;S769;T773;S804;S860;S894;S960;S1144;S1149;S1288;S1436	34418053;28528544;25153642	MEDSHELDTLYVTERIIAVSFPASCS EESYLHSLQEVTRMLKCKHGDNYLV LNLSEKRYDLTKLNPKIMDVGWPEL HAPPLDKMCTICKAQESWLNNDPQ HVVVIHCRGGKGRIGVVISSYMHFT NVSASADQALDRFAMKKFYDDKISA LMEPSQKRYVQFLSGLLSGAMKMN TSPLFLHFVIMHGVPSTGACRP FLKLYQAMQPVYTSGIYNVGSNPS RIRIAIEPAQLLKGDIMVKCYHKKFR SATRDVIFRLQFHTGAVQGYGLLFG KEELDSACKDDRFPDYGKIELVFSAT PEKIQGSEHLYSDQGVTVDYNTADP LIRWDSYENMSADGEVLHTQGPVD GSLYAKVRKKSASDTGIPSSPQGMPA TSSPDHGDHTLSVSSDSGHSTASAR TDKTEERLTPGARRGLSPQEKALD QLLSGFGLEDSASSHKDMTDMRSK YSGTRHVPAQVHVNGDAALKDRET DILDDEMPHHDLSVDSLGLTSSSE GPQSTHLGPFTCLKSSQNSLLSDGF GNGVAEDHNGVLSPLDGLGVDTLY DRERMCGGREQKPLQPLLRKPSAPT PVQAYGQSNYSTQTVWRQQQMVAA HQYSFASDGEARLGSRSTVDNTGLA QPPHIPVTPNRGASSRVAVQRGISN GPNPPDTQQLCPGKALQPRFQDDR TNGVHQEPNTGSSPGSPTLDIDQSIE QLNRLILELDPTFEPIPTHNLALGISA VCPDGVGSLRCSGRLDSVDGPGRS PGRQGDDPIGGRLRKL SIGQYDNDA ASQVTFSKCGWGKAGVDPAPSLGSF SSPEDIKETVITAYPSDLNMIDGRIPN SKESSMCLTPSFPVSPETPYVKTSR YPPFSPPEQLSSPASLHKGREPRGC PEIISHTVGMSESPVGPKPTMLRAD MPATPNFQQVFASSCTVSSNGPGQR RESPPSAERQWVESPSTLTLGN SHPSESPLGTHEFCSSGKDSPLPC FQSSELQASFHSHELSMSEPQALP PAGSQTFGLFNTVTTATSVLPPGEDA GTLVNSHGTSPAPGTPLLTTGAAD NGFLPHNFLTSPGASSHHSPGLQN QNVSLPGQPPLPEKKRASEGDRSLG SVSPSSSGFSSPHSGSTMSIPFPNVL PDFCKPSEVASPLPDSNDKLVIVKF VQDTSKFWYKADISREQAIAMLKDK APGSFIVRDSHSFRGAYGLAMKVAT PPPSVLHLNKKAGDLSNELVRHFLI ECTPKGVRLKGCNEPYFGSLTALV CQHSITPLALPCKLLIPERDPLEEIAE

NSPQTAANSAEELLKQGAACNVWYL
NSVEMESLTGHQAVQKALSMTLVQ
EPPPVSTVVHFKVSAQGITLTDNQRK
FFRRHYPVSSVIFCALDPQDRKWIK
DGPSSKVFVARKQGSATDNVCHL
FAEHDPEQPASAIVNFVSKVMIGSPK
KI