	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q8CGB6	TENS2_MOUSE	Mus musculus	Tensin-2	25.775171	NaN	T91;S118;S120;S455;T456;S466;T474;S481;T483;S820;S825;S830;S832;S835;T909;S931;S941;S972;T977;S991;S1003;S1087;T1173;S1238	34678516	MKSSGPVERLLRALGRRDSSRATSR PRKAEPHSFREKVFRKKTPVCAVCK VTIDGTGVSCRVCKVATHRKCEAKV TSSCQALPPAELRRSTAPVRRIEHLG STKSLNHSKQRSTLPRSFSLDPLME RRWDLDLTYVTERILAAAFPARPDE QRHRGHLRELAHVLQSKHRDKYLL FNLSEKRHDLTRLNPKVQDFGWPE LHAPPLDKLCSICKAMETWLSADPQ HVVVLYCKGSKGKLGVIVSAYMHYS KISAGADQALATLTMRKFCEDKVAT ELQPSQRRYVSYFSGLLSGSIRMNSS PLFLHYVFVPVLPAFEPNTGFQPFLK IYQSMQLVYTSGVYRIAGPGPQQLCI SLEPALLLKGDVMVTCYHKGGQGTD RTLVFRVQFHTCTIHGSRLTFPKDQL DEAWADERFPFQASVEFVFSSSPEK VKGNTPRNDPSVSVDYNTTEPAVRW DSYENFNQHHEDSVDGALAHTRGP LDGSPYAQVQRVPRQTPPAPSPELPP PPMLSVSSDSGHSSTLTTEHTAESP GRPPPTAAERQELDRLLGGCGVASA GRGAGRETAILDDEEQPSVGGGLHL GMYSGHRPGLSRRCSCRQGFREPC GVPNGSYYRPEGTLERRRPPYGGYE GHPQGYAEASVEKRRLCRSLSEGPY PYAPELGKPANGDFGYRPAGYREVVI LEDPGVPALCSCPACEEKLALPTAAL YGLRLEREAAEGWSSEVGKPLLHPV RPGHPLPLLVPACGHHHAPMPDYG CLKPPKVGEEGHEGCSYAVCSEGRY GHSGYPALVTYGYGGAVPSYCPAYG RAPHSCGSPSEGRGYPSPGAHSPRA GSVSPGSPPYLQPRKLGYEISAEDGR DKYPLSGHLASTGPLASTESPEPSW RDGSSGHSTLPRSPRDPQCSASSEL SGPSTPLHTSSPVQGKESNRRQDTT RSPSLAPTQRLSPGEALPSVVQGVAE KTPELLTSSRPEQLDPSPFSQTSAPG SPNGWPQERSPGGHTNSASPRSPVP TTLPGLRHAPWQGPRGTSDSPDGSP LTPVPTQMPWLVGSPEPPQSSPTPA FPLATSYDANGPIQPPLPEKRHLPGS GQQPSPPARSTNQHVTFASPLPDVT QPPEHPLQENQSNVKFVQDTSKFW YKPHLSRDQAIALLKDKDPGAFLIRD SHSFQGAYGLALKVATPPPSAQPWK GDPSEQLVRHFLIETGPKGVKIKGCP TEPYFGSLSALVSQHSISPISLPCCLR IPSKDPLEETPEAPVPTNMSTAADLL RQGAACSVLYLTSVETESLTGPQAVA KASSAALSCSPVPVPAIVHFKVSAQG ITLTDNQRKLFFRRHYPVNSITFSST DPQDRRWTNPDGATSKIFGFVAKKP GSPWENVCHLFAELDPDQPASAIVT FITKVLLGQRK	None	None	None	None	None	None	None	None	None