

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q5T7N3	KANK4_HUMAN	Homo sapiens	KN motif and ankyrin repeat domain-containing protein 4	28.529092	S172	NaN	29485866;38253038	MEKTDAKDQSSQGDEEKDPPKSHPYSVETPYGFHLDLDFLKYVDDIEKGN TIKRIPIHRRAKQAKFSTLPRNFSLPD SGARPPAAPPLQNWSFVVPREASLG TQEQNQSPPLGNAPQASTSRSEVSY HRKALLAEATROLEAAEPEDAELTF GSGRPQLLRASSMPATLLHSRASEE PGLSLGPPAPPALPPLQEGEGSVCDGT FEPAEGLAGFHSSSPRASTRIPELVQ EGAEPPPEGVVKVPHLPLPGPPFSF QNVLVVLEDKEDAHNAREAEVLF GSPTSPPLPSPIPENELLEEIELN ISEIPPPVPEVDMRSIGIRVTEESLG LARVDPGSISSLKQVSALEGELSGR TEELAQVRTALQQEEEEIKAREQRIR ELEFTVAQLEGQFHQENAKDTQGG TDVMVNTDPVHGLLTRESCDKGIEV NLLGSMESESWGHRGEENGLWGP DGHKQGNQSPAERVLLPQLSLPQGP EQVLTSSVHSFLSTELRIEAGTEQE GGPQGGTRGAGGFLWGS DRKTPPA GREETSNNLPGKEHPGRPPSSPTDA TIGQYVKIQELLQEQWNCLEHGYP ELASAIKQPAASKLSSIQSLLSSNLL LSAYSAQAHPPKEPPASSSSPPVEISP STSLKSIMKKKDYGFRAAGNGTKKN LQFVGVNGGYETTSSEETSGEDSTP EDLSDSEAEKKCDGPDHKHKVDAH LTCEAGQGIPEGTCHAAQESGPGEE VPHSKAERYKPSSEFLNACRALSQH LPETGTTTQLLRQSLNTISQEWFR VSSRKSSPAVVASYLHEVQPHSPH FLKLLVNLADHNGNTALHYSVSHSN FSIVKLLLETGVCNVDHONKAGYTA VMITPLASAEETNEDMAVVWKLREG NVNIQATQGGQTALMLGVSHDRED MVQALLSCQADVNLQDHDGSSALM VACHHGNVDLVRLLLAHPACDSSLT DKAGRTALSIALKSPHMEIAGLLRA HAEQGRSLGL	None	None	None	None	None	None	None	None	

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