

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	e r
Q8N3D4	EH1L1_HUMAN	Homo sapiens	EH domain-binding protein 1-like protein 1	16.682301	S884;S1511	S170;S173;S191;S285;S310;S367;S430;T454;S895;S964;S1168;S1257;S1273	30379171;37217939	MTSVWKRLQRVGKRAAKFQFVACY HELVLLECTKKWQPKLVVVVWTRRN RRICSKAHSWQPGIQNPYRGTVVW MVPENVDISVTLYRDPHVDQYEAKE WTFIIEESKQQRKVLATAEVDLAR HAGPVPVQVPVRLRLKPKSVKVVQA ELSLTSLGVLREGRATDDDDMQSLA SLMSVKPSDVGNLDDFAESDEDEA HGPGAPEARARVQPPDPSRELKTLC EEEEEGQGRPQQAVASPSNAEDTSP APVSAPAPPARTSRGQGSERANEAG GQVGPAPRPPETSPEMRSRQPAQ DTAFTPAPRLRKGSDALRPPVPQGE DEVKASGAPPAGLGSARETQAQAC PQEGTEAHGARLGPSIEDKSGSDPF GRQRLKAEEMDTEDRPEASGVDTPEP RSGGREANTKRSRVGAGEAEISSAV CQVDAEQRSKVRHVDTKGPEATGV MPEARCRGTPEAPRGSQGRGLGVRT RDEAPSGLSLPPAEPAGHSGQLGDL EGARAAAGQEREGAEVRRGGAPGIEG TGLEQGPSVGAISTRPOVSSWQCAL LSTAQGAISRGLGGWEAEAGSGDL ETETEVVGLVLTQEKVEVGGSGFP ETRTLEIEILGALEKEAARSRLVESEV AGTAQCEGLETQETEVGVIETPGTET EVLGTQKTEAGSGVLQTRTTIAETE VLVTQEISGDLGPKIEDTIQSEMLG TQETEVEASRVPESEAEAGTEAKILGT QETIARDSGVREIEAEIAESDILVAQE IEVGLLGVLGIEGAAEGAILGTQIEIA SRDSGVPGLEADTTGIQVKEVGGSE VPEIATGTAETEILGTQEIASRSSGVP GLESEVAGAQETEVEGGSGISGPEAG MAEARVLMTRKTEIIVPEAEKEEAQT SGVQEAETRVGSALKYEALRAPVTO PRVLGSQEAKEISGVQGSSETQVLRV QEAEAGVWGMSEKSGAWGAQEA EMKVLESPEPKSGTFKAQEAEGVLE GNEKGEAEGLSTEASLPEAQVAVSG AGAGAPRASSPEKAEEDRRLPGSQA PPALVSSQSLEWCOEVTITGYRGV RITNFTTSWRNGLAFCAILHRFYDPK IDYASLDPLNIKQNNKQAFDGFAL GVSRLLEPADMVLLSVPDKLIVMTYL CQIRAFCTGQELQLVQLEGGGGAGT YRVGSAQSPDDLDAGGLAQRRLRG HGAEGPOEPKAAADRADGAAPGVAS RNAVAGRASKDGGAEAPRESRPAEV PAEGLVNGAGAPGGGGVRLRRPSVN GEPGSVPPRAHGSFHVADADLLK KRRSRLRNSSFSMDDPDAGAMGA AAAEQGAPDPSAPGPPTAADSQQP PGGSSPSEPPSPGEEAGLQRFQD TSQYVCAELQALEQEQRQIDGRAAE VEMQLRSLMESGANKLQEEVLIQE WFTLVNKNALIRRDQLQLLMEE QDLERRFELLSRELRAMLAIEDWQK TSAQQHREQLLLEELVSLVNQRDEL VRDLDHKERIALEEDERLERGLEQR RRKLSRQLSRRERCVL	False	False	3.699	2.701	1.826	1.22	1.203	2.49	