	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	ex re
0	Q6ZS30	NBEL1_HUMAN	Homo sapiens	Neurobeachin-like protein 1	20.761126	T268;T1871	NaN	30379171;35289036;38253038	MASRERLFELWMLYCTKKDPDYLKL WLDTFVSSYEQFLDVDFEKLPTRVD DMPPGISLLPDNILQVLRIQLLQCVQ KMADGLEEQQQALSILLVKFFIILCR NLSNVEEIGTCSYINYVITMTTLYIQQ LKSKKKEKEMADQTCIEEFVIHALAF CESLYDPYRNWRHRISGRILSTVEKS RQKYKPASLTVEFVPFFYQCFQESE HLKESLKCCLLHLFGAIVAGGQRNA LQAISPATMEVLMRVLADCDSWED GDPEEVGRKAELTLKCLTEVVHILLS SNSDQRQVETSTILENYFKLLNSDH SALPNQRRSRQWENRFIALQIKMLN TITAMLDCTDRPVLQAIFLNSNCFE HLIRLLQNCKVFQGQLDCLAISTIQA LTAVMNKSPAAKEVFKERIGYTHML EVLKSLGQPPLELLKELMNMAVEGD HTSVGILGISNVQPLLLLIQWLPELQ SHDLQIFISDWLKRICCINRQSRTTC VNANMGIRIIETLDLHSSLHQTCAE NLIAIHGSLGSQSVSSEEIRRLLRLLR VDESESVHPYVTPVTRAILTMARKLS LESALQYFNLSHSMAGISVPPIQKW PGSAFSFSAWFCLDQDQLTLGIANK GGKRKQLYSFFTGSGMGFEAFITHS GMLVVAVCTKREYATVMLPDHSFC DSLWHNITVVHMPGKRPFGQSFVYI YDNGQQKVSAPLRFPAMNEPFTSCC IGSAGQRTTTPPPSQIPDPPFSSPITP HRTSFGGILSSASWGGTIEKSKLITK LISAGTQDSEWGCPTSLEGQLGSVII FYEPLQPPQVKALYLAGPNCLSPWK CQESDMADLPGNILLYYTAKACKNSI CLDLSTNCLHGRLTGNKVVNWDIK DIINCIGGLNVLFPLLEQISHFSEGQI PEEKNESTVPESVTPVEGDWLVWTS TKASESRLERNLVATFILIVKHFIQR HPINQGNLIHSHGVATLGALLQKVP STLMDVNVLMAVQLLIEQVSLEKN MQLLQQMYQYLLFDFRIWNRGDFP FRIGHIQYLSTIIKDSRRVFRKKYGVQ FLLDTLRIYYGNGCKYNELSLDDIRTI RTSLYGLIKYFLCKGGSHEEIQSIMG YIAATNEEEQLFGILDVLFSLLRTSPT RGQLFLLLFEPGNADILYALLLNQKY SDRLREIIFKIMEQMLKCTNVYERSK QHIRLREVGYSGLGLLLNEALVNTSL IKNLTHQIINTDPVINFKDLLSVVYIS HRAHINVRVAICRKVLQILQFQPDAA HQISQQVGWQDTLVRLFLKAKFENG NTLHKHSRAVLMKDNDKNMSTEDT KKNSDEKTDEEKITSFASANVSSDQ WSLEDRHSLDSNTPLFPEDSSVGEL SFKSENQEEFWHSNPSHLSLDLSGI DSCEMSDSGSQVPDSLPSTPSPVES TKSFSVHSDRESSITNDMGFSDDFS LLESQERCEEELLQLLTHILNYVMC KGLEKSDDDTWIERGQVFSALSKPGI SSELLRPSDEIKLTLLQKMLEWAISE NREAKTNPVTAENAFRLVLIIQDFLQ SEGLVNSNMWTEKLLEDMMLLFDC LSVCYSESPVWVKLSQIQIQLLLGFI GRGNLQVCAMASAKLNTLLQTKVIE NQDEACYILGKLEHVLSQSIKEQTEI YSFLIPLVRTLVSKIYELLFMNLHLPS LPFTNGSSSFFEDFQEYCNSNEWQV YIEKYIVPYMKQYEAHTFYDGHENM ALYWKDCYEALMVNMHKRDREGG ESKLKFQELFVEPFNRKARQENLRY NNMLKQLSSQQLATLRRWKAIQLYL TCERGPWAKRKQNPIHWKLANVEN YSRMRLKLVPNYNFKTHEEASALRD NLGIQHSQPSSDTLLLEVVKQVKVS DMVEDKLDLPEEDITARVNVDEKEE QDQKEKLVLMEDCELITIIDVIPGRL EITTQHIYFYDGSIEKEDGVGFDFKW PHSQIREIHLRRYNLRRSALEIFHVD QSNYFLNFKKEVRNKIYSRLLSLHSP NSYYGSRSPQELFKASGLTQKWVNR EISNFDYLIQINTMAGRTYNDLAQYP	None	None	None	None	None	None	None	None	No

									VFPWILQDYTSEELDLNNPAVFRDL SKPIGVVNEKNAKAMREKYENFEDP MGTIDKFHYGTHYSNSAGVMHYLIR VEPFTTLHIQLQSGRFDCADRQFHSI PATWQALMDNPYDVKELIPEFFYFP EFLENQNQFNLGRLQISKELVNDVI LPKWAKSAEDFIYKHRKALESEYVSA HLHEWIDLIFGYKQRGPAAVEALNV FYYCSYEGAVDLDALTDEKERKALE GMINNFGQTPCQLLKEPHPPRLSAE EAVQKPTKIDTSTLNLFQHLPELKSF FIEGISDGIPLLKATIPKNQYRSFMSQ GSPELLITISMNYVIGTHGWLPYDRN ISNYFTFIKDQTVTNPKTQRSINGSF APGLEITSKLFVVSHDAKLLFSAGYW DNSIQVMSLTKGKIISHIIRHMDIVT CLATDYCGIHLISGSRDTTCMIWQIT QQGGVPVGLASKPFQILYGHTNEVL SVGISTELDMAVSGSRDGTVIIHTIQ KGQYMRTLRPPCESSLFLTIPNLAIS WEGHIVVYSSTEEKTTLKDKNALHL FSINGKYLGSQILKEQVSDICIIGEHI VTGSIQGFLSIRDLHSLNLSINPLAM RLPIHCVCVTKEYSHILVGLEDGKLI VVGVGKPAEMRSGQLSRKFWGSSK RLSQISAGETEYNTQDSK								