

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	extracellular region
Q9ULB1	NRX1A_HUMAN	Homo sapiens	Neurexin-1	16.682301	S1024	NaN	30379171;37217939	MGTALLQRGGCFLLCLSLLLGCGWA ELGSGLEFPGAEGQWTRFPKWNAC CESEMSFQLKTRSARGLVLYFDDEG FCDFLELILTRGGRLQLSFSIFCAEPA TLLADTPVNDGAWHSVRIRRRQFRNT TLFIDQVEAKWVEVKSKRRDMTVFS GLFVGGPPELRAAALKLTLASVRER EPPFGWIRDVRVNSSQVLPVDSGEV KLDDPEPPNSGGGSPCEAGEEGEGGV CLNGGVCSVDDQAVDCSRTGFR GKDCSQEDNNVEGLAHLMMGDQG KSKGKEEYIATFKGSEYFCYDLSQNP IQSSSEITLSFKTLQRNGLMLHTGK SADYVNLALKNGAVSLVINLGSAGF EALVEPVNGKFNDAWHDVKVTRN LRQHSIGHAMVTISVDGILTTTGYT QEDYTMLGSDFFYVGGSPSTADLP GSPVSNFMGCLKEVYKNNNDVRL ELSLAKQGDPKMKIHGVVAFKCN VATLDPITFETPESFISLPKWNAKKT GSISFDFRITTEPNGLILFSHGKPRHQ KDAKHPQMIKVDFFAIEMLDGHLYL LLDMGSGTIKALKKKVNDGEWYH VDFQRDRSGTISVNTLRTPYTAPGE SEILDDELYLGGLPENKAGLVFPT EVWTALLNYGVGCIRDLDIDGQSK DIRQMAEVQSTAGVKPSCSKETAKP CLSNPCKNNGMCRDGNWRYVDCD SGTYLGRSCEREATVLSYDGSMMFM KIQLPVMHTEAEDVSLRFRSQRAY GILMATTSRDSADTLRLELDAGRVL TVNLDCIRINCNSKGPETLAFAGYNL NDNEWHTVRVRRGKSLKLTVDQDQ QAMTGOMAGDHTRLFHNIEGTIIT ERRYLSSVPSNFIGHLQSLTFNGMA YIDLCKNGDIDYCELNARFGFRNIIA DPVTFKTKSSYVALATLQAYTSMHLF FQFKTTSLDGLLYNSGDGDNDFIVVE LVKGYLHYVFDLGNGANLIKSSNK PLNDNQWHNVMSRDTSNLHTVKI DTKITTOITAGARNLDLKSPLYIGGV AKETYKSLPKLVHAKGFPQGCLASV DLNGRLPDLISDALFCNGQIERGCE GPSTTCQEDSCSNQGVCLQQWDGF SCDCSMTSFSGPLCNDPGTYIFSK GGGQITYKWPPNDRPSTRADRLAIG FSTVQKEAVLVRVDSGLGDYLEL HIHQKIGVKFNVGTDDIAIEESNAII NDGKYHVVRFRTRSGGNATLQVDSW PVIERYPAGRQLTIFNSQATIIIGGKE QGQPFQGLSGLYNGKLVNMAA ENDANIAIVGNVRLVGEVSSMTTE STATAMQSEMSTSIMETTTTLATSTA RRGKPTKEPISQTTDDILVASAEC SDEEDIDPCEPSSGGLANPTRAGGR EPYPGSAEVIRESSSTTGMVVGIVAA AALCILILLYAMYKYRNRDEGSYHVD ESRNYISNSAQSNQAVVKEKQPSSA KSSNKNKKNKDKKEYV	True	True	1.894	4.442	2.241	3.941	1.872	4.785	2.795