

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
P55067	NCAN_RAT	Rattus norvegicus	Neurocan core protein	23.439218	NaN	NaN	38843836	MGAESVWASGLLVLWLLLLVSGDQ DTQDTTTEKGLHMLKSGSGPIQAA LAELVALPCFFTLQPRQSPLGDIPRIK WTKVQTASGQRQDLPLILVAKDNVVR VAKGWQGRVSLPAYPRHRANATLLL GPLRASDSGLYRCQVVKGIEDEQDL VTLEVTGVVVFHYRAARDRYALTFAEA QEACHLSSATIAAPRHLQAAAFEDGF DNC DAGWLSDRTVRYPIQSRPGCY GDRSSLPGVRSYGRRDPQELYDVYC FARELGGEVFVYGPARRLLTAGARA LCORQGAALASVGOHLAWHEGLD QCDPGWLADGSVRYPIQTPRRRCGG SAPGVRTVYRFANRTGFPAPGARFD AYCFRAHHHTPQRGDSEIPSSGDEG EIVSAEGPPAPELKPRLGEQEVITPD FQEPLVSSGEDEPLDLTRTQASQETL ASTPGGPTLASWLLTGVTSSGVPSP SSLGVDMEETTPSGTQVAPTPTMRR GRFKGLNGRHFQQQGPEQDQLEAA EASAPPTLEVTADHMGPSAATEAL ESDQSHSPWAILTNEVDVPGAGSLG SRSLESRKWSPLISPSTVPSTDST PGLKPGADEAPGVKSAIHHPPWLP EPVAVSSIPSEALSAVSLQASPGDGS PDFPIVAMLRAPKLWLLPHSTLVPN VSPILPASPLPSSVPEEQAVRPVVF GAEDPETPFQTTMAAPGEASHGSPE ADSIEIEGISSMQATKHPISGPWASL DSSNVTNVPVPSDAGILGTESGVLDL PGSPTS DGQATVDMVLATWLPPLPGH GLDTGSQSTPMEAHGVTMSVEPTVA LEGGATKDPMEATMDVVPSTVDATS GSEPKSSISSTHVVVTAAGDQGTPTL TPTSSEGQVVAQESLGTLSLPSHP WSSLASSMDEVASVSSGEPTRLWDI PSTLIPVSLGLDESDLKVVAESPLE GFWEVAVSGQEDPTDPCENNPC LH GGTCRTNGTMYGCSCDQGYAGENC EIDIDDCLCSPCENGGTCIDEVNGFI CLCLPSYGGNLCEKDTGCDRGWH KFQGH CYRYFAHRRAWEDAERDCR RRAGHLTSVHSP EEFKINSFGHEN SWIGLNDRTVERDFQWTDNTGLQY ENWREKQPDNFFAGGEDCVMVA HENGRWNDVPCNYNLPYCKKGTV LCGPPPAVENASLVGVRKVKYNVHA TVRYQCDEGFSQHHVATIRCRSNGK WDRPQIVCTKPRRSHRMRHHHHHP HRHHKPRKEHRKHKRHPAEDWEK DEGDFC	False	True	0.607	0.764	0.639	False	False	4.208	5.0