

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	
0	Q62889	NLGN3_RAT	Rattus norvegicus	Neurologin-3	23.439218	NaN	S745;T792	38843836	MWLQLGLPSLSLSPPTVGRSLCLIL WFLSLVLRASSTQAPAPTNTNTHFGKL RGARVPLPSEILGPVDQYLGVPYAAP PIGEKRFLPPEPPPSWSGIRNATHFP PVCPONIHTAVPEVMLPVWFTANLD IVATYIQEPNEDCLYLNIVYVPTEDVK RISKECARCPNKKICRKGSGAKKQ GEDLADNDGDEDEDIRDSGAKPVM VYIHGGSYMEGTGNMIDGSVLASYG NVIVITLNYRVGVLGFLSTGDQAAKG NYGLLDQIQALRWVSENIAFFGGDP RRITVFGSGIGASCVSLTSLSHHSEG LFQRAIIQSGSALSWSAVNYQPVKYT SLLADKVGCVLDTVDMVDCLRQK SAKELVEQDIQPARYHVAFGPVIDGD VIPDDPEILMEQGEFLNYDIMLGVN QGEGLKFVEGVVDPEDGVSGTDFDY SVSNFVDNLYGYPEGKDTLRETIKF MYTDWADRDNPEPTRRKTLLVALFTD HQWVPEPSVVTADLHARYGSPTYFYA FYHHCQSLMKPAWSDAAHGDEVY VFGVPMVGPTDLFPCNFSKNDVML SAVVMTYWTNFAKTGDPNKPVPQD TKFIHTKANRFEVAVSKYNPRDQL YLHIGLKPRVRDHYRATKVAFWKHL VPHLYNLHDMFHYTSTTTKVPPDPT THSSHITRRPNGKTWSTKRPAISPAY SNENAPGSWNGDQDAGPLLVENPR DYSTELSVTIAVGASLLFLNVLFAAA LYYRKDKRRQEPLRQPSFQRTGAP ELGTAPEEELAAALQLGPTHHECEAG PPHDTLRLTALPDYTLTLRRSPDDIP LMTPNITITMIPNSLVGLQTLHPYNTF AAGFNSTGLPNSHSTTRV	False	True	1.581	1.791	1.46	1.48	1.128	5.0	2.955