

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	extracell region
Q8K1S4	UNC5A_MOUSE	Mus musculus	Netrin receptor UNC5A	47.04859	T421;S427	NaN	22645316;34678516;40885482	MAVVRPGLWPALLGIVLTAWLRGSGA QQSATVANPVPGANPDLLPHFLVEP EDVYIVKNKPVLLVCKAVPATQIFFK CNGEWVRQVDHVIERSTDGSSGLPT MEVRINVSRRQVQKVFGLLEEYWCQ CVAWSSSGTTKSKAYIRIAYLRKNF EQEPLAKEVSLSEQGIVLPCRPPGIP PAEVEWLRNEDLVDPSLDPNVYITR EHSLVVRQARLADTANYTCVAKNIV ARRRSASAAVIVYVNGGWSTWTEWS VCSASCGRGWQKRSRCTNPAPLN GGAFCEGQNVQKTACATLCPVDGS WSPWSKWSACGLDCTHWSRECS DPAPRNGGEECRGADLDRNCTSDL CLHTSSGPELVALYIGLVAVAVCLILL LLVLVLIYCRKKEGLSDVADSSILTS GFQPVSIKPSKADNPHLLTIQPDLS TTTTYQGSLCPRQDGPSPKFQLSNG HLLSPLGSGRHLHHSSPTSEAEF VSRLSTQNYFRSLPRGTSNMAYGTF NFLGGRLMIPNTGISLLIPPAIPRGK IYEIYLLHKPEDVRLPLAGCQTLSP IVSCGPPGVLLTRPVILAMDHCGEPS PDSWSLRLKKQSCGSGWEDVLHLG EESPSHLYYCQLEAGACYVFTEQLG RFALVGEALVAATKRLRLLLFAPVA CTSLEYNIRVYCLHDTHDALKEVVO LEKQLGGQLIQEPRVLHFKDSYHNL RLSIHDVPSLWKSLLVSYQEIPFY HIWNGTQQYLHCTFTLERVNASTD LACKVWWVQVEGDGQSFNINFNIT KDTRFAEMLALESEGGVPALVGPSA FKIPFLIROKIITSLDPPCSRGADWRT LAQKLHLDSHLSFFASKPSPPTAMIL NLWEARHFPPNGNLGQLAAAVAGLG QPDAGLFTVSEAE	True	True	4.173	1.437	0.856	0.784	False	4.571	2.168