

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q99MR3	S12A9_MOUSE	Mus musculus	Solute carrier family 12 member 9	25.810548	NaN	S6	29187734	MASESSPLLAYRLLGEEGAFFPPNG AGVSGVPSSRKLSTFLGVVPTVLS MFSIVVFLRIGFVVGHAGLLQALAM LLVAYIILALTVLSVCAIATNGAVRGG GAYFMISRTLGPVEVGGSIGLMFYLA NVCCGAVSLLGLVESILDVFGADAT GSSGIQVLPQGYGWNLLYGSLLLGL VGGVCTLGAGLYARASFLTFLVSGS LASVLVSFVAVGPRNIPLAPRPGTNA SSVPHRHGHFTGFNGSTLRDNLGA GYAEDYTTGAMMTFASVFAVLFNGC TGIMAGANMSGELKDPSRAIPLGTII AVAYTFFIYILLFLLSSFTCDRALLQE DYGFFRDISLWPPVLVIGIYATALSAS MSSLIGASRILHALAQDDLFGVILAP AKVVSGGGNPWGAVLYSWGVLVQLV LLAGKLNLAAVVTVFYLVAYAADVLDL SCLSLEWASAPNFRPTFSLFSWHTC LLGVASCLLMMFLISPGAAGGSLLL MGLLSALLTARGGPSSWGYVSQALL FHQVRKYLRLDVRKEHVKFWRPQL LLLVGNPRGALPLRLANQLKKGGL YVLGHVTLGDLDLSPDPVQPQYGA WLSLVDLAQVKAFVDLTLSPSVRQG AQHLLRISGLGGMKPNTLVLFYDD APPQDHFLTDPAFSEPAEAGTREGGS PALSTLFPFRAPGSPRALSPQDYVA TVADALKMNKNVVLARACGALPPER LSRGSSSAQLHHVDVWPLNLLRPR GGPGYVDVCGFLLOMATILSMVPA WHSARLRIFLCGPREAPGAAEGRL RALLSQLRIRAEVQEVVWGEAETG EPEEEEGDFVNGGRGDEEAEALACS ANALVRAQQGRGTVGGPGGPEGRD GEEGPTTALTFLYLPRPADPARYPR YLALLETLSRDLGPTLLIHGVTPTC TDL	False	True	1.259	1.318	1.434	0.73	0.817	4.461	1.575