

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
Q91V14	S12A5_MOUSE	Mus musculus	Solute carrier family 12 member 5	33.64034	NaN	T57;S1044;S1047;S1048	33300544;22645316	MSRRFTVTSLPAAASAADPESRR HSVADPRRLPREYDKGDNPKFESSP FINSTDTEKGREYDGRNMFEEEM DTSPMVSSLLSGLANYTNLPQGSRE HEEAENNEGKPKVQAPRMGTFFM GVYLPCLQONIFGVILFLRLTWVVGIA GIMESFCMVFICCSCTMLTAISMSAI ATNGVVPAGGSYYMISRLGPEFGG AVGLCFYLGTTFAGAMYLGTEILLA YLPFAMAIFKAEDASGEAAAMLNN MRVYGTCLVTCMATVVFVGVKYVVK FALVFLGCVILSILAIYAGVIKSAFDP NFPICLLGNRTLNRHGFDVCAKLA EGNETVTRLWGLFCSSRLLNATCD EYFTRNNVTEIQGIPGAASGLIKENL WSSYLTKGVIVERRGMPVSVGLADGT PVDMDHPYVFSDMTSYFTLLVGIYF PSVTGIMAGSNRSGDLRDAQSIPT GTILAIATTSAVYISSVVLFGACIEGVV LRDKFGEAVNGNLVVGTLAWPSPW VIVIGSFFSTCGAGLQSLTGAPRLLQ AISRDGIVPFLQVFGHGKANGPTW ALLLTACICEIGILIASLDEVAPILSMF FLMCYMFVNLACAVQTLRLRTPNWR PRFRYYHWTLNFLGMSLCLALMFIC SWYYALVAMLIAGLIYKYIEYRGAEK EWGDGIRGLSLSAARYALLRLEEGPP HTKNWRPQLLVLRVDQDQNVVHP QLLSLTSQKAGKGLTIVGSVLEGT LDNHPQAQRAEESIRRLMEAEKVKG FCQVVISSNLRDGVSHLIQSGGLGGL QHNTVLVGWPRNWRQKEDHQTWR NFIELVRETTAGHLALLVTKNVSMF PGNPERFSEGSIDVWVHDGGML MLLPFLRRHHKVWRCKMRIFTVA QMDDNSIQMKKDLTTFYHLRITAE VEVVMHESDISAYTYEKTLMVEQR SQILKQMHLLTKNEREREIQSITDESR GSIRRNPNPRLRLNVPEETACDN EEKPEEEVQLIHDQSAPSCPSSSPSP GEEPEGERETDPEVHLTWTKDKSVA EKNKGPSVSSSEGIKDFFSMKPEWE NLNQSNVRRMHTAVRLNEVIVNKS RDAKLVLNMPGPPRRNRNGDENYM EFLEVLTEQLDRVMLVRGGREVITI YS	False	True	1.936	2.155	1.749	1.872	1.457	5.0	2.381