	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q63633	S12A5_RAT	Rattus norvegicus	Solute carrier family 12 member 5	23.439218	NaN	T57;T929;T1030;S1045;S1048;S1049	38843836	MSRRFTVTSLPPAASAASADPESRR HSVADPRRLPREDVKGDGNPKESSP FINSTDTEKGREYDGRNMALFEEEM DTSPMVSSLLSGLANYTNLPQGSKE HEEAENNEGGKKKPVQAPRMGTFM GVYLPCLQNIFGVILFLRLTWVVGIA GIMESFCMVFICCSCTMLTAISMSAI ATNGVVPAGGSYYMISRSLGPEFGG AVGLCFYLGTTFAGAMYILGTIEILLA YLFPAMAIFKAEDASGEAAAMLNN MRVYGTCVLTCMATVVFVGVKYVNK FALVFLGCVILSILAIYAGVIKSAFDPP NFPICLLGNRTLSRHGFDVCAKLAW EGNETVTTRLWGLFCSSRLLNATCD EYFTRNNVTEIQGIPGAASGLIKENL WSSYLTKGVIVERRGMPSVGLADGT PVDMDHPYVFSDMTSYFTLLVGIYF PSVTGIMAGSNRSGDLRDAQKSIPT GTILAIATTSAVYISSVVLFGACIEGVV LRDKFGEAVNGNLVVGTLAWPSPW VIVIGSFFSTCGAGLQSLTGAPRLLQ AISRDGIVPFLQVFGHGKANGEPTW ALLLTACICEIGILIASLDEVAPILSMF FLMCYMFVNLACAVQTLLRTPNWR PRFRYYHWTLSFLGMSLCLALMFIC SWYYALVAMLIAGLIYKYIEYRGAEK EWGDGIRGLSLSAARYALLRLEEGPP HTKNWRPQLLVLVRVDQDQNVVHP QLLSLTSQLKAGKGLTIVGSVLEGTF LDNHPQAQRAEESIRRLMEAEKVKG FCQVVISSNLRDGVSHLIQSGGLGGL QHNTVLVGWPRNWRQKEDHQTWR NFIELVRETTAGHLALLVTKNVSMF PGNPERFSEGSIDVWWIVHDGGML MLLPFLLRHHKVWRKCKMRIFTVA QMDDNSIQMKKDLTTFLYHLRITAE VEVVEMHESDISAYTYEKTLVMEQR SQILKQMHLTKNERE REIQSITDESR GSIRRKNPANTRLRLNVPEETACDN EEKPEEEVQLIHDQSAPSCPSSSPSP GEEPEGEGETDPEKVHLTWTKDKSA AQKNKGPSPVSSEGIKDFFSMKPEW ENLNQSNVRRMHTAVRLNEVIVNK SRDAKLVLLNMPGPPRNRNGDENY MEFLEVLTEQLDRVMLVRGGGREVI TIYS	False	True	1.964	2.193	1.703	1.541	1.07	5.0	2.48