	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	Q9BXP2	S12A9_HUMAN	Homo sapiens	Solute carrier family 12 member 9	38.068807	NaN	S6	41018975;34725712	MASESSPLLAYRLLGEEGVALPANG AGGPGGASARKLSTFLGVVVPTVLS MFSIVVFLRIGFVVGHAGLLQALAM LLVAYFILALTVLSVCAIATNGAVQG GGAYFMISRTLGPEVGGSIGLMFYLA NVCGCAVSLLGLVESVLDVFGADAT GPSGLRVLPQGYGWNLLYGSLLLGL VGGVCTLGAGLYARASFLTFLLVSGS LASVLISFVAVGPRDIRLTPRPGPNG SSLPPRFGHFTGFNSSTLKDNLGAG YAEDYTTGAVMNFASVFAVLFNGCT GIMAGANMSGELKDPSRAIPLGTIVA VAYTFFVYVLLFFLSSFTCDRTLLQE DYGFFRAISLWPPLVLIGIYATALSAS MSSLIGASRILHALARDDLFGVILAP AKVVSRGGNPWAAVLYSWGLVQLV LLAGKLNTLAAVVTVFYLVAYAAVDL SCLSLEWASAPNFRPTFSLFSWHTC LLGVASCLLMMFLISPGAAGGSLLL MGLLAALLTARGGPSSWGYVSQALL FHQVRKYLLRLDVRKDHVKFWRPQ LLLLVGNPRGALPLLRLANQLKKGG LYVLGHVTLGDLDSLPSDPVQPQYG AWLSLVDRAQVKAFVDLTLSPSVRQ GAQHLLRISGLGGMKPNTLVLGFYD DAPPQDHFLTDPAFSEPADSTREGS SPALSTLFPPPRAPGSPRALNPQDYV ATVADALKMNKNVVLARASGALPPE RLSRGSGGTSQLHHVDVWPLNLLR PRGGPGYVDVCGLFLLQMATILGMV PAWHSARLRIFLCLGPREAPGAAEG RLRALLSQLRIRAEVQEVVWGEGAG AGEPEAEEEGDFVNSGRGDAEAEAL ARSANALVRAQQGRGTGGGPGGPE GGDAEGPITALTFLYLPRPPADPARY PRYLALLETLTRDLGPTLLVHGVTPV TCTDL	False	True	0.63	0.889	0.791	0.509	0.818	4.403	4.181