

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	
0	O35316	SC6A6_MOUSE	Mus musculus	Sodium- and chloride-dependent taurine transporter	25.385059	NaN	S322	28528544	MATKEKIQCLKDFHKDILKPSPGKS PGTRPEDEADGKPPQREKWSSKIDF VLSVAGGFVGLGNVWRFPYLCYKNG GGAFLLPYFIFLFGSGLPVFFLEVIIG QYTSEGGITCWEKICPLFSGIGYASIV IVSLLNVYYIVILAWATYYLFHSFQKD LPWAHCNHSWNTQPQCMEIDLRRN ESHWVSLSTANFTSPVIEFWERNVL SLSSGIDNPGSLKWDLALCLLLVWL VCFFCIWKGVRSTGKVYVFTATPFFA MLLVLLVRGLTLPGAGEGKIFLYPD ISRLGDPQVWIDAGTQIFFSYAICLGA MTSLGSYNKYKNSYRDCMLLGCL NSGTSFVSGFAIFSILGFMAQEQV DIADVAESGPGLAFAIYKAVTMMPL PTFWLILFIMLLLLGLDSQFVEVEG QITSLVDLYPSFLRKGYRREIFAILCS ISYLLGLTMVTEGGMYVFLQFDYYA ASGVCLLWVAFFECFVIAWIYGGDN LYDGEDMIGYRPGPVMKYSWAVIT PALCVGCFVFLVKYVPLTYNKVYRY PDWAIGLGWGLALSSMLCIPLVIVIL LCRTEGPLRVRIKYLITPREPNRWAV EREGATPFHSRVTLMNGALMKPSH VIVETMM	False	True	1.268	1.484	1.567	1.17	0.602	4.725	1.433