

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 Q91X34	BAAT_MOUSE	Mus musculus	Bile acid-CoA:amino acid N-acyltransferase	23.627151	NaN	S125;S418	37453647	MAKLTAVPLSALVDEPVHIQVTGLAP FQVVCLQASLKDEKGNLFSSQAFYR ASEVGEVDLEHDPSTLGGDYMGVHP MGLFWSLKPEKLLGRLIKRDVMNSP YQIHKACHPYFPLQDIVVSPPLDSL LERWYVAPGVKRIQVKESRIRGALFL PPGEGPFPGVIDLFGGAGGLMEFRA SLLASRGFATLALAYWNYDDLPSRL EKVDLEYFEEGVEFLLRHPKVLGPG VGILSVCIGAEIGLSMAINLKQIRATV LINGPNFVSQSPHVYHGQVYPPVPS NEEFVVTNALGLVEFYRTFQETADK DSKYCFPIEKAHGHFLFVVGEDDKN LNSKVHANQAIQALMKNKGNWTL LSYPGAGHLIEPPYTPLCQASRMPILI PSLSWGGEVIPHAAAQEHSWKEIQK FLKQHLLPDLSSQL	True	False	5.0	2.239	2.152	2.025	0.605	1.638	1.6