

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
Q8BML9	SYQ_MOUSE	Mus musculus	Glutamine-tRNA ligase	25.685855	NaN	S70;S495	39627609	MATPDSLALFTGLGLENKARETLK NEALSTQLREAAATQAHQILGSTIDKA TGVLLYDLVSRRLDTRRRSFLVSYIA NKKIHTGLQLSAALEYVRSHPQDPID TKDFEQECGVGVVVTPEQIEEAVEST INKHQLQLLAERYRFNMGLLMGEA RAALRWADGKMIKNEVDMQVLHLL GPKMEADLVKKPKVAKARLEETDRK TAKDVVEKGEVAGQTLSLMEQLRGE ALKFHKPGENYKTPGYVITPYTMDLL KQHLEITGGQVTRFPPEPENGILHIG HAKAINFNFGYAKANNNGICFLRFDD TNPEKEEAKFFTAIYDMVTWLGYTP YKVTYASDYFDQLYAWAVELIHGGL AYVCHQRVEELKGHNPSPWRDR PKEESLLFEAMRKGKFAEGEATLR MKLVMEDGKMDPVAYRVKYTPHHR TGDKWCYPTYDYTHCLCDSIEHITH SLCTKEFQARRSSYFWLCNALKVYC PVQWEYGRLNLHYAVVSKRKILQLV AAGAVRDWDDPRLFTLTALRRRGFP PEAINNFCARVGVTVVAQTMEPHLL EACVRDVLNDAAPRAMAVLEPLQVV ITNFPAPKPLDIRVNFPADETKGFH QVPFASTVFIERSDFKEESEPGYKRL ASGQPVGLRHTGYVIELQNIVRGSSG CVERLEVTCRRADAGEKPKAFIHVV SQPLVCEIRLYECLFQHKNPEDPVEV PGGFSDLNFPASLQVVEGALVDCSV ALAKPFDKQFERLGYFSVDPDSHQ GQIVFNRTVTLKEDPGKI	True	False	4.323	2.443	1.737	1.022	False	0.925	0.974