

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membr
Q8CFE4	SCYL2_MOUSE	Mus musculus	SCY1-like protein 2	52.506565	S622;T722;S728;S741	S658;S677;T708	22517741;22645316;40885482;36852467;30059200;40997131	MESMLNKLKSTVTKVTADVTSAVM GNPVTREFDVGRHIASGGNGLAWKI FNGTKKSTKQEVAVFVFDKKLIDKY QKFEKDQIIDSLKRGVQLTRLRHP RLLTVQHPLEESRDCLAFCTEPVFAS LANVLGNWENLPSSISPDIKDYKLYD VETKYGLLQVSEGLSFLHSSVKMVH GNVTPENVILNKSGAWKIMGFDFCV SSSNPSEQEPKFPCKEWDPNLPSLC LPNPEYLAPEYILSVSCETASDMYSL GAVMYAVFNQGRPIFEVKNQDIYKS FSRQLDQLSRLGSSSLTSIPEEVREH VKLLLNVTPTVRPDADQMTKIPFFD DVGAVTLQYFDTLFQRDNLQKSQFF KGLPKVLPKLPKRIVIVORILPCLTSEF VNPDMVFPVLPNVLLIAEECTKEYEI KLILPELGPVFKQEPPIQILLIFLQKM DLLLTKTPPDEIKNSVLPVMVYRALEA PSIQIQELCLNIPTFANLIDYPSMKN ALIPRIKNAQLTSSLAVRVNSLVCL GKILEYLDKWFVLDLIPFLQIQIPSK EPAVLMGILGIYKCTFTHKKLGITKE QLAGKVLPHLIPLSIENNLNLNQFSS FIAVIKEMLSRLESEHRTKLEQLHV MQEQQRSLDIGNQMSTSEETKVAH SGSQQDKVFNNGADLLSGSESEN REDGMQKQKRGSLTLEEKQKLAK EQEQAQKLSQQPLKPOVHTPIAPIK QTKDLTDTLMENMSSLTSLSVSTPK ISASSTFTVPVSTGLGMMFSTPIDNT KRNLTNGLNANMGFQTSGFSPVNV PNQNFFSGTGTAGVTTMSLGAPPTM SNFSPLTIPPASVKQPQRPTDMSAL NNLFGPOKPKVSMNQLSQQKPNQ WLNQFAPPQGSVPMGSAAMGTQG NVMGQAAFGMQGNPFFNPQNFQAQ PPPTTMTSSSSASNDLKDLFG	False	False	1.748	1.431	0.469	False	3.398	0.657