

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
P51692	STA5B_HUMAN	Homo sapiens	Signal transducer and activator of transcription 5B	21.366884	NaN	T90;S128;S193;T682;T699	28510447	MAVWIIQAQQLQGEALHQMQUALYGO HFPIEVRHYLSQWIESQAWDSVDLD NPQENIKATQLLEGLVQELQKKAEH QVGEDGFLLKIKLGHYATQLQNTYD RCPMELVRCIRHILYNEQRLVREAN NGSSPAGSLADAMSQKHLQINQTFE ELRLVTQDTENELKKLQQTQOYFIIQ YQESLRQAQFGPLAQLSPQERLSRE TALQQKQVSLEAWLQREAO TLQQYR VELAEKHQKTLQLLRKQQTILDDDEL IQWKRQQLAGNGGPPGSLDVLQS WCEKLAEIWQNRQQIRRAEHLCCQ LPIPGPVEEMLAEVNATITDIISALVT STFIIKQPPQVLKTQTKFAATVRLLV GGKLNVMNPPQVKATIISEQQAAS LLKNENTRNDYSGEILNCCVMEY HQATGTLSAHFRNMSLKRKRSDRR GAESVTEEKFTILFESQFSVGGNELV FOVKTLSLPVVIVHGSQDNNATAT VLWDNAFAEPGRVVFVAVPKVLWP QLCEALNMKFKAQEVQSNRGLTKEN LVFLAQKLFNNSSSHLEDYSGLSVS WSQFNRENLPGRNYTFWQWFDGV MEVLKKHLKPHWNDGAILGFVNKO QAHDLLINKPDGTFLLRFSDEIGGI TIAWKFDQSERMFWNLMPTTRDF SIRSLADRLGDLNYLIYVFPDRPKDE VYSKYTPVPCESATAKAVDGYVKPQ IKQVVPEFVNASADAGGGSATYMDQ APSPAVCPQAHYNYMYPQNPDSVLDT DGDFDLEDTMDVARRVEELLGRPM DSQWIPHAQS	True	False	4.685	5.0	2.689	2.36	1.927	2.916	3.238