	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ext reg
0	Q8SY33	GAWKY_DROME	Drosophila melanogaster	Protein Gawky	28.06426	T436;T437;S892;S1077;T1081;T1084	NaN	40245499	MREALFSQDGWGCQHVNQDTNWE VPSSPEPANKDAPGPPMWKPSINNG TDLWESNLRNGGQPAAQQVPKPSW GHTPSSNLGGTWGEDDDGADSSSV WTGGAVSNAGSGAAVGVNQAGVNV GPGGVVSSGGPQWGQGVVGVGLGS TGGNGSSNITGSSGVATGSSGNSSN AGNGWGDPREIRPLGVGGSMDIRN VEHRGGNGSGATSSDPRDIRMIDPR DPIRGDPRGISGRLNGTSEMWGHH PQMSHNQLQGINKMVGQSVATAST SVGTSGSGIGPGGPGPSTVSGNIPTQ WGPAQPVSVGVSGPKDMSKQISGW EEPSPPPQRRSIPNYDDGTSLWGQQ TRVPAASGHWKDMTDSIGRSSHLM RGQSQTGGIGIAGVGNSNVPVGANP SNPISSVVGPQARIPSVGGVQHKPDG GAMWVHSGNVGGRNNVAAVTTWG DDTHSVNVGAPSSGSVSSNNWVDD KSNSTLAQNSWSDPAPVGVSWGNK QSKPPSNSASSGWSTAAGVVDGVDL GSEWNTHGGIIGKSQQQQKLAGLNV GMVNVINAEIIKQSKQYRILVENGFK KEDVERALVIANMNIEEAADMLRAN SSLSMDGWRRHDESLGSYADHNSS TSSGGFAGRYPVNSGQPSMSFPHN NLMNNMGGTAVTGGNNNTNMTAL QVQKYLNQGQHGVAVGPQAVGNSS AVSVGFGQNTSNAAVAGAASVNIAA NTNNQPSGQQIRMLGQQIQLAIHSG FISSQILTQPLTQTTLNLLNQLLSNIK HLQAAQQSLTRGGNVNPMAVNVAI SKYKQQIQNLQNQINAQQAVYVKQQ NMQPTSQQQQPQQQQLPSVHLSNS GNDYLRGHDAINNLQSNFSELNINK PSGYQGASNQQSRLNQWKLPVLDK EINSDSTEFSRAPGATKQNLTANTS NINSLGLQNDSTWSTGRSIGDGWPD PSSDNENKDWSVAQPTSAATAYTDL VQEFEPGKPWKGSQIKSIEDDPSITP GSVARSPLSINSTPKDADIFANTGKN SPTDLPPLSLSSSTWSFNPNQNYPS HSWSDNSQQCTATSELWTSPLNKS SSRGPPPGLTANSNKSANSNASTPT TITGGANGWLQPRSGGVQTTNTNW TGGNTTWGSSWLLLKNLTAQIDGPT LRTLCMQHGPLVSFHPYLNQGIALC KYTTREEANKAQMALNNCVLANTTI FAESPSENEVQSIMQHLPQTPSSTS SSGTSGGNVGGVGTSANNANSGSA ACLSGNNSGNGNGSASGAGSGNNG NSSCNNSAAGGGSSSNNTITTVANS NLVGSSGSVSNSSGVTANSSTVSVV SCTASGNSINGAGTANSSGSKSSAN NLASGQSSASNLTNSTNSTWRQTS QNQALQSQSRPSGREADFDYISLVY SIVDD	True	False	2.244	4.099	False	0.912	False	0.506	0.6