

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9QX74	SHAN2_RAT	Rattus norvegicus	SH3 and multiple ankyrin repeat domains protein 2	12.481722	NaN	S457;T486;S586;S724;T903;S1334;S1338;S373;S373;S162;S162;S162	18683930	MKSLLNFTKKEVPPFREAPAYSNRR RRPPNTLAAPRVLLRSNSDNNLNAG APEWAVCSAATSHRSLSPQLLQOTP SKPDGATKSLGSYAPGPRSRSPSLNR LGGAGEDGKRPPPHWHVVGSPFTP GANKDSLSTFEYPGPRRKLYSAPVG RLFVAIKPYQPQVDGEIPLHRGDRVK VLSIGEGGFWECSARGHIGWFPAEC VEEVQCKPRDSQAETRADRSKKLFR HYTVGSYDSFDAASDCIIEDKTVVLO KKDNEGFVLRGAKADTPIEEFTPT PAFPALQYLESVDEGGVAWQAGLRT GDFLIEVNNENVVKVGHQVVMNI RQGGNHLVLKVVTVTRNLDPPDDTAR KKAPPPPKRAPTTALTLRSKSMATAEL EELGLSLVDKASVRKKKDKPEEIVPA SKPSRTAENVAIESRVATIKQRPTSR CFPAASDVNSVYERQGIAMTPTVP GSPKGPFLGLPRGTMRRQKSIDSRI LSGITEEERQFLAPPMLKFTRSLSM DTSEDIPPPQSVPPSPPPSPTTYNC PRSPTRVYGTIKPAFNQNPVAKVPP ATRSDTVATMMREKGMFYRRELD FSLDSEDVYSRSPAPQAAFRTKRGQ MPENPYSEVGKIASKAVYVPAKPARR KGMLVKQSNVEDSPEKTCSEIPIIIV KEPSTSSSGKSSQGSMEIDPQATEP GQLRPDDSLTVSSPFAAAIAGAVRDR EKRLAARRNSPAFLSTDLDGEDVGL GPPAPRMQPSKFPEEGGFGEDETE QPLLPTPGAAPRELENHFLGGGEAG AQGEAGGPLSSTSKAKGPESGPA LKSSSPASPENYVHPLTGRLLDPSS LALALSARDRAMQESQQGHKGEAP KADLNKPLYIDTKMRPSVESGFPPVT RQNTRGPLRRQETENKYETDLSKDR RADDKKNMLINIVDTAQKKSAGLLM VHTVDIPVAGPPLEEEEDREDGDTK PDHSPSTVPEGVPKTEGALQISAAPE PAAAPGRTIVAAGSVEEAVILPFRIPP PPLASVDLDEDLFTTEPLPPLEFAN SFDIPDDRAASVPALADLVKQKSDT PQPPSLNSSQPANSTDSKKPAGISN CLPSSFLPPESFDAVTDGIEEVDS RSSSDHHLETTSTISTVSSISTLSSEG GESMDTCTVYADGQAFVVDKPPVPP KPKMKPIVHKSALYQDTLPEEDTD GFVIPPAPPPPPGSAQAGVAKVIQP RTSKLWGDVTEVKSPILSGPKANVIS

