

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9UPX8	SHAN2_HUMAN	Homo sapiens	SH3 and multiple ankyrin repeat domains protein 2	25.208493	NaN	S831;T860;S960;S1099;T1278;S1709;S1713	30620550;37340703;35132862;28657654	MPRSPTSSSEDEMAQSFSDYVSGSES DSSKEETIYDTIRATAEKPGGARTEE SQGNTLVIRVVIHDLQQTCKIRFNP ATVWVAKQRILCTLTQSLKDVLNYG LFQPASNGRDGKFLDEERLLREYPQ PVGEGVPSLEFRYKRVYKQASLDE KQLAKLHTKTNLKKCMDHIQHRLV EKITKMLDRGLDPNFHDPETGETPL TLAAQLDDESVEVIKALKNNGGAHLDF RAKDGMTALHKAARARNQVALKTLL ELGASPDYKDSYGLTPLYHTAIVGGD PYCCELLLHEHATVCKDENGWHEI HQACRYGHVQHLEHLLFYGADMSA QNASGNTALHICALYNQDSCARVLL FRGGNKELKNYNSQTPFQVAIIAGN FELAEYIKNHKETDIVPFREAPAYSN RRRRPPNTLAAPRVLLRSNSDNNLN ASAPDWAVCSTATSHRSLSPQLLQQ MPSKPEGAAKTIGSYVPGPRSRSPSL NRLGGAGEDGKRPQPLWHVGSPPFA LGANKDSLSAFEYPGPKRKLYSAPV GRLFVAVKPYQPQVDGEIPLHRGDR VKVLSIGEGGFWECSARGHIGWFPA ECVEEVQCKPRDSQAETRADRSKLL FRHYTVGSYDSFDTSSDCIIEEKTIV LQKKNNEGFGFVLRGAKADTPIEEF TPTPAFPALQYLESVDEGGVAWQAG LRTGDFLIEVNNENVVKVGHQRQVNV MIRQGGNHLVVKVTVTRNLDPDDT ARKKAPPPKRAPTTALTLRSKSMTS ELEELVDKASVRKKKDKPEEIVPASK PSRAAENMAVEPRVATIKQRPSSRC FPAGSDMNSVYERQGIAMTPTVPG SPKAPFLGIPRGTMRQKSIDSRIFL SGITEEERQFLAPPMLKFTRLSMP DTSEDIPPPQSVPPSPPPSPTTYNC PKSPTPRVYGTIKPAFNQNSAAKVSP ATRSDTVATMMREKGMFRRELDLDR YSLDSEDLYSRNAGPQANFRNKRQ MPENPYSEVGKIASKAVYVPAKPARR KGMLVKQSNVEDSPEKTCIPIPTIIV KEPSTSSSGKSSQGSSMEIDQAPEP PSQLRPDESLTVSSPFAAAIAGAVRD REKRLEARRNSPAFLSTDLGDEDVG LGPPAPRTRPSMFPEEGDFAEDSA EQLSSPMPSATPREPENHFVGGAEA SAPGEAGRPLNSTSKAQGPSSPAV PSASSGTAGPGNYVHPLTGRLLDPS SPLALALSARDRAMKESQOGPKGEA PKADLNKPLYIDTKMRPSLDAGFPTV TRQNTRGPLRRQETENKYETDLGRD RKGDDKKNMLIDIMDTSQQKSAGLL MVHTVDATKLDNALQEEDEKAEVE MKPDSSPSEVPEGVSETEGALQISAA

