

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9WV48	SHAN1_RAT	Rattus norvegicus	SH3 and multiple ankyrin repeat domains protein 1	13.680566	NaN	T186;S540;S671;S791;S898;S1291;S1442;S638;S641	22645316	MTHSPATSEDEERHSASECPEGGSE SDSSPDGPGRGPQGRGRGSGAPG NLASTRGLQGRSMSVPDDAHFSMM VFRIGIPDLHQTKCLRFNPDATIWTA KQQVLCALSESLQDVLNYGLFQPAT SGRDANFLEEERLLREYQSFKEGV PYLEFRYKTRVYKQTNLDEKQLAKL HTKTGLKKFLEYVQLGTSDKVARLL DKGLDPNYHSDSDSETPLTLAAQTE GSVEVIRTLC LGGAHIDFRARDGMT ALHKAACARHCLALTALLDLGGSPN YKDRRGLTPLFHTAMVGGDPRCCEL LLYNRAQLGIADENGWQEIHQACQR GHSQHLEHLLFYGAEPGAQNASGN TALHICALYNKETCARILLYRGANKD VKNNNGQTPFQVAVIAGNFELGELI RNHREQDVVPFQESPKYAARRRGPP GAGLTVPPALLRANSDTSMALPDW MVFSAPGASSSGTPGPTSGPQQSQ PSAPSTKLSSGTLRSASSPRGARARS PSRGRHPEDAKRQPRGRPSSSGTPR DGPAGGTGGSGGPGGSLGSRGRRR KLYSAVPGRSFMAVKSYQAQGEI SLSKGEKIKVLSIGEGGFWEQVKG RVGWFPSDCLEEVANRSQEGRQES RSDKAKRLFRHYTVGSYDSFDAPSLI DGIDSGSDYIIKEKTVLLQKKDSEGF GFVLRGAKAQTPIEFTPTPAFPALQ YLESVDEGGVAWRAGLRMGDFLIEV NGQNVVKVGHQRQVNMIRQGGNTL MVKVVMVTRHPDMDEAVHKKASQ QAKRLPPPAILRSKSMTSELEEMVS PWKKKIEYEQQPAAVPSMEKKRTVY QMALNKLDEILAAAQQTISASESPGP GGLASLGKHRPKGFFATESSFDPHH RSQPSYDRPSFLPPGPGLMLRQKSIG AAEDDRPYLAPPAMKFSRSLSVPGS EDIPPPPTTSPPEPPYSTPPAPSSSGR LTPSPRGGFNPSSGGPLPASSPSSF DGPSPDTRGGGREKSLYHSAALPP AHHHPPHHHHHHHAPPPQPHHHHA HPPHPPEMETGGSPDDPPRLALGP QPSLRGWRGGGPSPTSGAPSPSHHS SSGGSSGPTQAPALRYFQLPPRAASA AMYVPARSGRGRKGPLVKQTKVEGE PQKGSIPSASSPTSPALPRSEPPPAGP SEKNSIPIPTIIKAPSTSSSGRSSQGS STEAEPPTQPDGAGGGGSSPSPAPAT SPVPPSPVPTPASPSGPATLDFTS

QFGAALVGAARREGGWQNEARRRS
TLFLSTDAGDEDEGGDSGLGPGGPPG
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PQPGDPPPP GTDSGIEEVDSRSSSDH
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QNWPKPPLPPLPTGSGVSSSTAAP
GATSPSASSASASTRHLOGVEFEMR
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