

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q4ACU6	SHAN3_MOUSE	Mus musculus	SH3 and multiple ankyrin repeat domains protein 3	18.754664	NaN	T122;S373;S375;S387;S394;S482;T555;S694;S781;S790;S801;S890;S897;T912;T930;S995;T1130;S1134;S1159;S1163;S1166;T1234;S1253;S1420;S1510;S1521;S1529;S1539;S1634;S1636;S1638	22645316;33300544	MDGPGASAVVVRVGIPDLQQTCKLR LDPTAPVWAAKQRVLCALNHSLQDA LNYGLFQPPSRGRAGKFLDEERLLQ DYPPNLDTPLPYLEFRYKRRVYAQNL IDDKQFAKLHTKANLKKFMDPVQVHL STDKVARLLDKGLDPNFHDPDSGEC PLSLAAQLDNATDLLKVLNRNGGAHL DFRTRDGLTAVHCATRQRNAGALTT LLDLGASPDYKDSRGLTPLYHSALG GGDALCCCELLLHDHAQLGTTDENG WQEIHQACRFGHVQHLEHLLFYGA NMGAQNASGNTALHICALYNQESC ARVLLFRGANKDVRNYSQTAFQVA IIAGNFELAEVIKTHKSDVVPFRET PSYAKRRRLAGPSGLASPRPLQRSAS DINLKGDPAAASPGPTLRSLPHQLLL QRLQEEKDRDRDGELENDISGPSAG RGGHNKISPSGPGGSGPAPGPGPAS PAPPAPPPRGPKRKLYSAVPGRKFIA VKAHSPQGEGEIPLHRGEAVKVLISG EGGFWEGETVKGRTGWFPADCVEEV QMRQYDTRHETREDRTKRLFRHYT VGSYDSLTSYSDYVIDDKVAILQKRD HEGFGFVLRGAKAETPIEEFTPTPAF PALQYLESVDVEGVAWRAGLRTGDF LIEVNGVNVVKVGHKQVVGLIRQGG NRLVMKVVSVTRKPEEDGARRRAPP PPKRAPSTTLTLRSKSMTAELEELAS IRRRKGEKLDEILAVAAEPTLRPDIAD ADSRAATVKQRPTSRRITPAEISSLFE RQGLPGPEKLPGLRKGIPRTKSVGE DEKLASLLEGRFPRSTSMQDTVREG RGIPPPPQTAPPPPPAPYYFDSGPPPT FSPPPPPGRAYDTRVSSFKPGLEARL GAGAAGLYDPSTPLGPLYPERQKRA RSMILQDSAPEVGDVPRPAPAATPP ERPARRPRPSGPDSPYANLGAFSASL FAPSKPQRRKSPLVKQLQVEDAQER AALAVGSPGPVGGSFAREPSPTHRC PRPGSLDYSSGEGLGLTFGGPSPGPV KERRLEERRRSTVFLSVGAIEGSPPS ADLPSLQPSRSIDERLLGTGATTGRD LLPSPVSALKPLVGGPSLGPSTFI HPLTGKPLDPSSPLALALAARERALA SQTSPRSPTPVHSPDADRPGPLFVD VQTRDSERGPLASPAFSPRSPAWIPV PARREAEEKPPREERKSPEDKKSMLS VLDTSLQRPAGLIVVHATSNGQEPSR LGAEERPGTPELAPAPMQAAVAE

PMPSPRAQPPGSIPADPGPGQGSSE
EPELVFAVNLPPAQLSSSDEETREE
LARIGLVPPPPEEFANGILLTTPPGPG
PLPTTVPSASGKPSSELPPAPESAA
DSGVVEADTRSSSDPHLETTSTISTV
SSMSTLSSSEGELTDHTSFADGHT
FLEKPPVPPKPKLSPLGKGPVTFR
DPLLQSSDSELMAQQHHAASTGLA
SAAGPARPRYLFQRRSKLWGPVES
RGLPGPEDDKPTVISELSSRLQQLNK
DTRSLGEEPVGGLGSLLDPAKKSPIA
AARLFSSLGELSTISAQRSPGGPGGG
ASYSVRPSGRYPVARRAPSPVKPASL
ERVEGLGAGVGGAGRPFGLTPPTILK
SSLSIPHEPKVRFVRSVSARSRS
PSPSPLPSPSPSGGPSAGPRRPFQOK
PLQLWSKFDVGDWLESIHLGEHRD
RFEDHEIEGAHLPALTKEDFVELGV
TRVGHRMNIERALRQLDGS