

β site



agrin_triad CQNGAQC SLGVQGYLCNCPQY YTGENC SQAI.....EVFGPSVAFDGNVYTLRNNISA
agrin_trica CLNNGLCVPELDQFICKCPSNTKGKYCE.....ISQEETP.AIKFNGATFLQFRN....
agrin_apime CHNGGVCLPLLNSYLCKCATGYNGLHCEFFMGYDVSTELTERPVRFKGDNLFQFRH....
agrin_cioin CDNGGVCHPRGAEYMCVCLPY YTGDNCEQEHSTDLLQDEQATAIYLDGTTKIMYRNAVKA
agrin_disom CQNGGVCS PRLREYDCMCQRGFSGPQCEKALEEKSASGSESVA..FNGRTFIEYHNTVTR
agrin_chick CQNGGTCS PRLREYECACQRGFSGAHCEKVII EKAAGDAEAIA..FDGRTYMEYHNAVTK
agrin_rat CLNGGSCVPREATYECLCPGGFSGLHCEKGLVEKSVG DLETLA..FDGRTYIEYLNVAIE
agrin_human CLNGASCVPREAAAYVCLCPGGFSGPHCEKGLVEKSAGDVDTLA..FDGRTFV EYLNVAITE

agrin_triadDEKEFNFN TIELOLKT FN LHGLILWSGPRNYSQNORRPFIALAVTNGNIQL
agrin_tricaRGYRRRKPERGNRYEIKLRTFASDALLLWRSK...SRSIREDYFSIAV VNGYPEI
agrin_apimeRNGRRRKQ QSNKFELRLRTHPDGLI AWIGR...G...KVEHLILSLHGGQVLL
agrin_cioin I.....SRARTHNNYETVFR TTA RHGLLLMVGK...ARE.GVDYIALA IHDGRLHL
agrin_disom SHLANEIPDEKAVQVNYFEMS IKT EATKGLILWSGK...IAE.KSDYIALAVVDGFVQM
agrin_chick SHLSNEIPA EKALQSNHFELSLRTEATQGLV LWIGK...AAE.RADYMALA IVDGHLQL
agrin_rat SELTNEIPA EKALQSNHFELSLRTEATQGLV LWIGK...AAE.RADYMALA IVDGHLQL
agrin_human SELTNEIPV EKALQSNHFELSLRTEATQGLV LWSGK...ATE.RADYVALA IVDGHLQL