

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6T4R5	NHS_HUMAN	Homo sapiens	Actin remodeling regulator NHS	8.666114	S1479	T332;T401;S415;S533;S739;S1176;T1262;S1329;S1499	35254053	MPFAKRIVEPQWLRCRQRRPAPGPAV DASGGSAEPPPPLQPPGRRDLDEVE APGPEEPARAVPAPSGLPPPPPPLPA PADQTQPPHGEASVAGEESTAGIPEA APAAGEASSAAAAAAILMLDLCAV SNAALARVLRQLSDVARHACSLFQE LESDIQLTHRRVWALQGKLGGVQRV LSTLDPKQEAVPVSNLDIKLSVYY RAPWHQQRNIFLPATRPPCVEELHR HARQSLQALRREHRSRSDRREQRA AAPLSIAAPPLPAYPPAHSQRRREFK DRHFLTFNSTRSPSPTECCHMTPWS RKSHPPEDEDTDVMLGQRPKNPIH NIPSTLDKQTNWSKALPLPTPEEKM KQDAQVISSCIIPINVTGVGFDREASI RCSLVHSQSVLQRRRKLRRRKTISGI PRRVQQEIDSDESPVARERNVIVHT NPDPSNTVNRISGTRDSECQTEDILI AAPSRRRIRAQRGQSIASLSHSAGN ISALADKGMTMFTPAVSSRTRSRSLP REGNRGGDAEPKVGAKPSAYEEGES FVGDHERTPNDFSEAPSSPSAQDHQ PTLGLACSQHLHSPQHLKLSERGRSR LSRMAADSGSCDISSNSDTFGSPIH CISTAGVLLSSHMDQKDDHQSSSGN WSGSSSTCPSQTSETIPPAASPLTG SSHCDSELSLNTAPHANEDASVFT EQYNDHLDKVRGHRANSFTSTVAD LLDDPNNSNTSDSEWNYLHHHHDA SCRQDFSPERPKADSLGCPSTMA TYDSFLEKSPDKADTSSHFSVDTE GYTSMHFDCGLKGNKSYVCHYAAL GPENGQVGASPLPDCAWQDYLD HKRQGRPSISFRKPKAKPTPPKRSS LRKSDGNADISEKKEPKISSGQHLP HSSREMKLPLDFANTPSRMENANL PTKQEPSWINQSEQGIKEPQLDASDI PPFKDEVAESTHYADLWLLNDLKTN DPYRSLSNSSTATGTTVIECIKSPSS ESQTSQSESRAATPSLPSVDNEFKLA SPEKLAGLASPSSGYSSQSETPTSSF PTAFFSGPLSPGGSKRKPVPKPKSS LQQPSTKDGITISLKDLELPIPPHTL DLSALHNVLNKPFFHRRHPLHVFTH NKQNTVGETLRSNPPPSLAITPTILK SVNLRKINKSEEVKQKEENNTDLPY LEESTLTTAALSPSKIRPHTANKSVS RQYSTEDTILSFLDSSAVEMGPKL HLEKNSTFDVKNRCDPETTTSAGSS

												LLDSNVTKDQVRTETEPENTPTKN CAFPTTEGFQRVSAARPNDLDGKIIQY GPGPDEETLEQVQKAPSAGLEEVAQP ESVDVITSQSDSPTRATDVSNOFKH QFVMSRHHDKVPGTISYESEITSVN SFPEKCSKQENIASGISAKSASDNSK AETQGNVDEASLKESSPDDSIISP LSEDSQAEAEGVVFVSPNKPRTTEDL FAVIHRSKRKVLGRKDSGDMSVRSK SRAPLSSSSSSASSITSPSSNVTPNS QRSPGLIYRNAKKSNTSNEEFKLLLL KKGSRSDSSYRMSATEILKSPILPKP PGELTAESPQSTDDAHQGSQGAEAL SPLSPCSPRVNAEGFSSKSFATSASA RVGRSRAPPAASSRYSVRCRLYNTP MQAISEGETENSDGSPHDDRSSQSS T
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