

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
014964	HGS_HUMAN	Homo sapiens	Hepatocyte growth factor-regulated tyrosine kinase substrate	13.766969	S297;S299;S300;S305;S306;S310;S315	T216;T308;T329;T334	34019948;34725712;32119511;22661428;28510447;35008409;35254053;35138101;33214551;37340703;29485866;30620550;35132862;29351928;30379171	MGRGSGTFERLLDKATSQLLLETDW ESILQICDLIRQGDQAKYAVNSIKKK VNDKNPHVALYALEVMESVVKNCG QTVHDEVANKQTMELKDLLKRVQ EVNVRNKILYLIQAWAHAFRNEPKY KVVQDTYQIMKVEGHVVFPEFKESDA MFAAERAPDWVDAEECHRCRVQFG VMTRKHHCRACGQIFCGKCSSKYST IPKFGIEKEVRVCEPCYEQLNRKAEG KATSTTELPEYLTSPLSQSQSOLPPK RDETALQEEEEELQALALSQSEAE KERLRQKSTYTSYPKAEPMPASSAP PASSLYSSPVNNSAPLAEDIDPELAR YLNRYWEKKQEEARKSPTPSAPVP LTEPAAQPGEGHAAPTNNVENPLPE TDSQPIPPSGGPFSEPFHNGESEES HEQFLKALQNAVTTFVNRMKSNHM RGRSITNDSAVLSLFQSINGMHPQL LELLNQLDERRLYEGLQDKLAQIR DARGALSALREEHREKLRRAAEEAE RQRQIQLAQKLEIMRQKKQEYLEVQ RQLAIQRLQEKEKERQMRLEQQKQ TVQMRAQMPAFPLPYAQLQAMPAA GGVLYQPSGPASFPSTFSPAGSVEGS PMHGVYMSQPAPAAGPYPSMPSTA ADPSMVSAYMYPAGATGAQAAPQA QAGPTASPAYSSYQPTPTAGYQNVAS QAPQSLPAISQPPQSSTMGYMGQS VSMGYQPYNMQLMTTLPSQDASL PPQQPYIAGQQPMYQQMAPSGGPPQ QQPPVAQQPQAQGPPAQGSEAQLIS FD