

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
Q0GNC1	INF2_MOUSE	Mus musculus	Inverted formin-2	22.102647	T100	S351;S1172;S1174;T1203;S1216;S1218;T1223;T1230	22645316;33300544;24788674;30016717	MSVKEGAQRKWAALKEKLGPDSD PTEANLESAEPELCIRLLQMPSPVNY SGLRKRLESSDGGWMVQFLEQSGL DLLLEALARLSGRGVARISDALLQLT CISCVRAVMNSQQGIEYILSNQGYVR QLSQALDTSNVMVKQVFELLAALC IYSPEGHALTLDALDHYKMVCSQOY RFSVIMSELSDSNVYVVTLLSVIN AAILGPEDLRSRAQLRSEFIGLQLLDI LTRLRDLEDADLLIQLEAFEEAKAED EEELQRISDGINMNSHQEVFASLFH KVSCSPASAQLLSVLQGLMHLEPAG RSGQLLWEALENLVNRVLLASDAQ ACTLEEVVERLLSIKGRPRPSPLDKA HKSVQTNVQNGSSSQNTTPTTK VEGQQPVVASPCQHVGSIQSSVDIA PQPVALEQCITALPLTPPLSSSTPVL PPTPPPLPGGATSPLPPPPPLPPPL PGSGTTSPPPPPPPPPLPPPLPGSGT ISPPPPPPPPPLPGTGAVSPPPPPPPL SLPDSHKTQPPPPPPPLPGMCPVPP PPPLPRAGQIPPPPPPLPGFSVPSMMG GVEEIIVAQVDHSLGSAWVPSHRRV NPPTLRMKKLNWQKLPSNVARERN SMWATLGSPCTAAVEPDFSSIEQLF SFPTAKPKPEPSAAPARKEPKEVTFLD SKKSLNINIFLKQFKCSNEEVTSMI QAGDTSKFDVEVLKQLLKLPEKHEI ENLRAFTEERAKLSNADQFYVLLLDI PCYPLRVECMMLCEGTAIVLDMVRP KAQLVLTACESLLTSQRLPVFCQLIL KIGNFLNYGSHTGDADGFKISTLLKL TETKSQSRVTLHHVLEEEVEKSHP DLLQLSRDLEPPSQAAGINVEIIHSE ASANLKKLLEAERKVSASIPEVQKQY AERLQASIEASQELDKVFDAIEQKLL ELADYLCEDPQQLSLEDTFSTMKTF RDLFTRALKENKDRKEQMAKAERR KQQLAEEREARRPRDEDGKPIRKGPG KQEEVCVIDALLADIRKGFQLRKTAR GRGDTEASGRVAPTDPKATEPATA SNPTQGTNHPASEPLDTTAADEPQG WDLVDAVTSPQPSKEEDGPPALER RSSWYVDAIDFLDPEDTPDAQPSEG VWPVTLGDQALNPLEFSSNKPPGV KSSHQDATDPEALWGVHQTEADST SEGPEDEAQRGQSTHLPRTPGGEDE DGEDTAPESALDTSLDRSFSEDAVT DSSGSGTLPRVQGRVSKGTSKRRKK RPSRNQEEFVPSDDIKAKRLCVIQ