	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellu region
0	Q14005	IL16_HUMAN	Homo sapiens	Pro-interleukin-16	29.956363	S1001;S1275;T1277	S922	27655845;29351928	MESHSRAGKSRKSAKFRSISRSLML CNAKTSDDGSSPDEKYPDPFEISLAQ GKEGIFHSSVQLADTSEAGPSSVPDL ALASEAAQLQAAGNDRGKTCRRIFF MKESSTASSREKPGKLEAQSSNFLF PKACHQRARSNSTSVNPYCTREIDF PMTKKSAAPTDRQPYSLCSNRKSLS QQLDCPAGKAAGTSRPTRSLSTAQL VQPSGGLQASVISNIVLMKGQAKGL GFSIVGGKDSIYGPIGIYVKTIFAGGA AAADGRLQEGDEILELNGESMAGLT HQDALQKFKQAKKGLLTLTVRTRLT APPSLCSHLSPPLCRSLSSSTCITKDS SSFALESPSAPISTAKPNYRIMVEVSL QKEAGVGLGIGLCSVPYFQCISGIFV HTLSPGSVAHLDGRLRCGDEIVEISD SPVHCLTLNEVYTILSHCDPGPVPIIV SRHPDPQVSEQQLKEAVAQAVENTK FGKERHQWSLEGVKRLESSWHGRP TLEKEREKNSAPPHRRAQKVMIRSS SDSSYMSGSPGGSPGSGSAEKPSSD VDISTHSPSLPLAREPVVLSIASSRLP QESPPLPESRDSHPPLRLKKSFEILV RKPMSSKPKPPPRKYFKSDSDPQKS LEERENSSCSSGHTPPTCGQEAREL LPLLLPQEDTAGRSPSASAGCPGPGI GPQTKSSTEGEPGWRRASPVTQTSPI KHPLLKRQARMDYSFDTTAEDPWV RISDCIKNLFSPIMSENHGHMPLQP NASLNEEEGTQGHPDGTPPKLDTAN GTPKVYKSADSSTVKKGPPVAPKPA WFRQSLKGLRNRASDPRGLPDPALS TQPAPASREHLGSHIRASSSSSSIRQ RISSFETFGSSQLPDKGAQRLSLQPS SGEAAKPLGKHEEGRFSGLLGRGAA PTLVPQQPEQVLSSGSPAASEARDP GVSESPPPGRQPNQKTLPPGPDPLL RLLSTQAEESQGPVLKMPSQRARSF PLTRSQSCETKLLDEKTSKLYSISSQ VSSAVMKSLLCLPSSISCAQTPCIPK EGASPTSSSNEDSAANGSAETSALD TGFSLNLSELREYTEGLTEAKEDDD GDHSSLQSGQSVISLLSSEELKKLIE EVKVLDEATLKQLDGIHVTILHKEEG AGLGFSLAGGADLENKVITVHRVFP NGLASQEGTIQKGNEVLSINGKSLK GTTHHDALAILRQAREPRQAVIVTRK LTPEAMPDLNSSTDSAASASAASDV SVESTAEATVCTVTLEKMSAGLGFSL EGGKGSLHGDKPLTINRIFKGAASE QSETVQPGDEILQLGGTAMQGLTRF EAWNIIKALPDGPVTIVIRRKSLQSKE TTAAGDS	None	None	None	None	None	None	None	None	None