

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P14735	IDE_HUMAN	Homo sapiens	Insulin-degrading enzyme	20.660394	NaN	NaN	30397120;34725712;38665916	MRYRLAWLLHPALPSTFRSVLGARL PPPERLCGFQKKTYSKMNNPAKRIG NHITKSPEDKREYRGLELANGIKVLL ISDPTTDKSSAALDVHIGSLSDPPNIA GLSHFCEHMLFLGTKKYPKENEYSQ FLSEHAGSSNAFTSGEHTNYYFDVS HEHLEGALDRFAQFFLCPLFDESCK DREVNADVSEHEKNVMNDAWRFL QLEKATGNPKHPFSKFGTGKNKYTLE TRPNQEGIDVRQELLKFHSAYYSSN LMAVCVLGRESLDDLTNLVVKLFSE VENKNVPLPEFPEHPFQEEHLKQLY KIVPIKDIRNLYVTFPIPDLOKYYKSN PGHYLGHLIGHEGPGSLLSELKSKG WVNTLVGGQKEGARGFMFFIINVDL TEEGLLHVEDIILHMFQYIQKLAEG PQEWVVFQECKDLNAVAFRFKDKER PRGYTSKIAGILHYYPLEEVLTAEYLL EEFRPDLIEMVLDKLRPENVRVAIVS KSFEGKTDRTTEEWYGTQYKQEAIPD EVIKKWQNADLNGKFKLPTKNEFIP TNFEILPLEKEATPYPALIKDTAMSK LWFKQDDKFFLPKACLNFEFFSPFA YVDPLHCNMAYLYLELLKDSLNEYA YAAELAGLSYDLQNTIYGMYSVKGY NDKQPILLKKIIEKMATFEIDEKRFEI IKEAYMRSLNNFRAEQPHQHAMY LRLLMTEVAWTKDELKEALDDVTLF RLKAFIPQLLSRLHIEALLHGNITKQ AALGIMQMVEDTLIEHAHTKPLPS QLVRYREVQLPDRGWFVYQQRNEV HNKCGIEIYYQTDMQSTSENMFLEL FCQIIESEPCFNTLRTKEQLGYIVFSGP RRANGIQGLRFHQSEKPPHYLESRV EAFLLTMEKSIEDMTEAFQKHQAL AIRRLDKPKKLSAECACYWGEIISQ YNFDRDNTEVAYLKLTKEDIHKFYK EMLAVDAPRRHKVSVHLAREMDS CPVVGEPFCQNDINLSQAPALPQPE VIQNMTEFKRGLPLFPLVKPHINFM AAKL	True	True	5.0	5.0	5.0	2.113	2.045	4.558	5.0