

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9BZQ6	EDEM3_HUMAN	Homo sapiens	ER degradation-enhancing alpha-mannosidase-like protein 3	13.899183	NaN	NaN	37217939;38665916	MSEAGGRGCGSPVPQRRARWLVAATAAFCLVSATSVVWTAGAEPMSREEKQKLGNOVLEMFDHAYGNYMEHAYPADELMPLTCRGRVVRGQEPSRGDVEDALGKFSLTLDLTLVVLNKTKEFEDAVRKVLRDVENLDNDVVVSVFETNIRVLGGLLGGHSLAIMLKEKGEYMQWYNDELLOMAKQLGYKLLPAFNNTSGLPYPRINLKFIRKPEARTGTETDTCTACAGTLLEFAALSRFTGATIFEYYARKALDFLWEKRQRSSNLVGVITINIHTGDWVRKDSGVGAGIDSYYEYLLKAYVLLGDDSFLERFNTHYDAIMRYSISQPPLLDVHIHKPMLNARTWMDALLAFFPGLQVLKGDIPAIETHEMLYQVIKKNHFLPEAFTTDFRVHWAQHPLRPEFAESTYFLYKATGDPYYLEVGKTLIENLNKYARVPCGFAAMKDVRTGSHEDRMDSFLLAEMFKYLKLLFADKEDIIFDIEDYIFTTEAHLPLWLSTNQSISKKNTTSEYTELDDSNFDWTCPN TQILFPNDPLYAQSIRESPLKNVVDKSCPRGIIRVEESFRSGAKPPLRARDFMATNPEHLEILKMGVSLIHLKDGRLVQVHAIQAASSIDAEDGLRFMQEMIELSSQQQKEQQLPPRAVQIVSHFFFGRVLTAGPAQFGLDLSKHKETRGFVASSKPSNGCSELNPEAVMGKIALIQRGQCMFAEKARNIQNAGAIGGIVIDDNEGSSDTAPLFQMAGDGKDTDDIKPMLFLFSKEGSIIIDAIREYEEVEVLLSDKAKDRDPEMENEOPSSENDSQNSGEGQISSSQEVLDLVDQESSEENSLNSHPESLSLADMDNAASISPSEQTSNPTENHETTNLNGECTDLDNQLQEQSETTEEDSNPNVSWGKVKVPIDLSILADWNEIEAFEMMEKDEL	False	True	2.519	1.308	3.033	4.498	1.574	1.212	1.412