	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	extracellular region
0	Q06704	IMH1_YEAST	Saccharomyces cerevisiae	Golgin IMH1	24.475439	NaN	S308;S660;S827;T830	33229814	MFKQLSQIGKNLTDELAKGLADDM SPTPSEQQIEDDKSGLPKEIQAKLRK FEKYEQKYPLLLSAYKNEKLKSEKLE AVEKILAENTPISNIDDAVDTLPAFF QDLNNKNNLLNDEIKRLTKQNSEIP ESASSETLKDKEEEFLKKEQNYKND IDDLKKKMEALNIELDTVQKEKNDT VSGLREKIVALENILKEEREAKKQKE EVSISELKEELAIKNHSLEDSRMKIT ELEQNLSSKSTIMEEKSSELAELNIT LKEKERKLSELEKKMKELPKAISHQ NVGNNNRRKKNRNKGKKNKGGITT GDISEEETVDNSINTEEYDKLKENLQ ELQEKYKDCEDWKQKYEDIEAELKD AKELENSQLEKSAKELETLNTELIDT KKSLKEKNSELEEVRDMLRTVGNEL VDAKDEIKESSSKQNEEVKTVKLEL DDLRHKNATMIEAYEAKNTELRSKI ELLSKKVEHLKNLCTEKEKEQTTSQ NKVAKLNEEISQLTYEKSNITKELTS LRTSYKQKEKTVSYLEEQVKQFSEQ KDVAEKSTEQLRKDHAKISNRLDLL KKENETLHNDIAKNSNSYEEYLKEN GKLSERLNILQEKYNTLQNVKSNSN EHIDSIKRQCEELNVKLKESTKKILS LEDELNEYANIVQDKTREANTLRRL VSDSQTDDSSKQKELENKLAYLTDE KNKLEAELDLQTSRKATELQEWKHT VTELKSEIHALKLREEGLKSEVDALK HVNNDIKRKTQATSDDSDQLEQITS NLKLSLSKADEKNFELQSANEKLLN LNNELNKKFDRLLKNYRSLSSQLNA LKERQYSDKSGRVSRSGSIGTLANA NIDSSPANNSNPTKLEKIRSSSSLEL DSEKNEKIAYIKNVLLGFLEHKEQR NQLLPVISMLLQLDSTDEKRLVMSL K	None	None	None	None	None	None	None	None	None