

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8CHT3	INT5_MOUSE	Mus musculus	Integrator complex subunit 5	25.685855	NaN	S278	39627609	MSALCDPPGAPGPPGPAPATHGPAP LSAQELSQEIKAFITGVDPILGHQLS AREHARCGLLLRSLPPARAALVDH LRGVFDESVAHLAALEESPVAGPP HLRPPPPSHVPTGGPGLDVVHEVQ QVLCFIRANPKAWAPVISAWSIDL MGQLSSTYSGQHQRVPHATGSLNE LLQLWMGCRATRLMDIYVQCSAL IGSCPACVDALLDTSVQHSPhFDW VVAHIGSSFPGTIISRVLSCGLKDFCV HSGAGGGASACGNSSQPPSTDPFPG SPAIPGEKRVPKIASVVGILGHLASR HGDSIRRELLRMFHDSLGGSGGR NGEPLQATVPFLLQLAVMSPALLG TVSGELVDCLKPPAVLSQLQQLQG FPREELDNMLNLAHLVLSQASGTGA YRLQLVDTAMPASVITTOGLAVPD TVREACDRLIQLLLHLQKLVHHRG GSPGEGVLGPPPPRPVFLDALRN HVGELCGETLRLEKRFLWQHQLL GLLSVYTRPSCGPEALGHLLSRARSP EELSLATQLYAGLVVSLSGLLPLAFR SCLARVHAGTLQPPFTARFLRNALL VGWEQQGGEGPSALGARFGESASA HLADLAPLLHPPEEVAAAAALLAI CPFPSEALSPSOLLGLVRAGVHHFF SSLRLHGPPGVASASQLLTRLSTP AGLKAVLQLLVEGALHRGNTELFGG EMDGDNETLSIVSTPLASALLDINR RHTAAVPGGGIWSVFHAGVIGRGL KPPKIVQSRNHQEVYINTQSLVSLLV HCCSASGNSEREGCWGAPTLSPEAA KAVAVTLVESVCPDAAGAELAWPPE DHARATVERDLRIGRRFREQLLFEF LKLVAAPPALCYCVLLRGLLAALL SHWEASRHPDTTHSPWHLEASCILV AVMAEGSLLPPALGNMHEVFSQLAP FEVRLLLSVWGFLEHGPLPKQKFI QSERGRFIRDFAREGGAEGGPHLSV LHSVLRNIDRLGLFSGRFQAPPST LLRQGT	True	False	2.0	4.257	0.528	False	False	2.906	False