

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi appara
Q9UKJ3	GPTC8_HUMAN	Homo sapiens	G patch domain-containing protein 8	9.350978	S908;S995	S491;S653;S738;S740;S758;S911;S914;S981;S1009;S1014;S1033;S1035;S1081;S1107;S1175	30379171;33214551;37217939;34019948	MADRFSRFNEDRDFQGNHFDQYEE GHLEIEQASLDKPIESDNIGHRLLOK HGWLKGGQLGKSLQGRTPPIVVK YDVMGMGRMEMELDYAEDATERR RVLEVEKEDTEELRQKYKDYVDKEK AIAKALEDLRANFYCELCDKQYQKH QEFDNHNSYDHAHKQRLKDLKQR EFARNVSSRSRKDEKKQEKALRRLH ELAEQRKQAECAPGSGPMFKPTTVA VDEEGGEDDKDESATNSGTGATASC GLGSEFSTDKGGPFTAVQITNTTGLA QAPGLASQGISFGIKNNLGTPLQKLG VSFSFAKKAPVKLESIASVFKDHAE GTSEDTGTPDEKSSDQGLQKVGDS GSSNLDGKKDEDPQDGGSLASTLS KLRMKREEGAGATEPEYYHYIPPA HCKVKPNFPFLFMRASEQMDGDN TTHPKNAPESKKGSSPKPKSCIKAAA SQGAECTVSEVSEQPKETSMTEPSE PGSKAEAKKALGGDVSDQSLESHSQ KVSETQMCESNSSKETSLATPAGKE SQEGPKHPTGPFPPVLSKDESTALQ WPELLIFTKAEPSISYSCNPLYDFDK LSRNDARTKGTEKPKDIGSSSKDH LQGLDPGEPNKSKEVGGKIVRSSG GRMDAPASGSACSLNKPQEPGGSH GSETEDTGRSLPSKKERSGKSHRHK KKKKKKSSKHKRKHKADTEEKSSK AESGEKSKRKRKRKKNKSSAPAD SERGPKPEPPGSGSPAPRRRRRAQ DDSQRRLPAEEGSSGKDEGGGGG SSQDHGGRKHKGELPPSSCRRRAG TKRSSRSHRSQPSGDESDDASS HRLHQKSPSYSEEEEEEDSGSEHS RSRSRSGRRHSSHRSSRRSYSSSD ASSDQSCYSRQRSYSDSDSYSDR SRRHSKRSHSDSDSYASSKHSKR HKYSSDDDYSLSCSQRSSRSRSH RERSRSRGRSSSSCSRSRSKRRSR STTAHSWQSRYSRDRSRSTRSPS QRSGSRKRSWGHEPESPEERHSGRRD FIRSKIYRSQSPHYFRSGRGEGPGKK DDGRGDDSKATGPPSQNSNIGTGRG SEGDCSPEDKNVTAKLLEKIQSRK VERKPSVSEEVQATPNKAGPKLKDP PQGYFGPKLPPSLGNKPVLPPLIGKLP ATRPNKCKEESGLERGEEQESET EEGPPGSSDALFGHOFSEETTGPL LDPPEESKSGEATADHPVAPLGTGA HSDCYPGDPPTISHNYLPDPDGDTL ESLDSSSQPGPVESLLPIAPDLEHF PSYAPPSGDPSEIESTDGAEDASLAPL ESQPIFTPEEMEKYSKLQAAAQOHI QQQLLAKQVKAFAASAALAPAL QPIHIQQPATASATSITTVQHAILQH HAAAAAIGIHPHPHPQPLAQVHHI PQPHLTPISLSHLTHSIIPGHATFLA SHPIHIIPASAIHPGPFTHFPVPHAAL YPTLLAPRAAAAATLHLHLLHPI FSGQDLQHPPSHGT	None	None	None	None	None	None	