

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	ext reg
Q3T906	GNPTA_HUMAN	Homo sapiens	N-acetylglucosamine-1-phosphotransferase subunits alpha/beta	29.286762	S384;S385;S390	NaN	28411811;37340703	MLFKLLQRQTYTCLSHRYGLYVCFLLGVVVTIVSAFQFGVEVLEWSRDQYHVLFDYSYRDNIAGKSFQNRCLPMPIDVVYTWVNGTDLELLKELQVREQMEEEQKAMREILGKNTTEPTKKSEKQLECLLTHCIKVPMLVLDPALPANITLKDPLSLYPSFHSASDIFNVAKPKNPSTNVSVVVFDSKDVEDAHSGLLKGNSRQTVWRGYLITDKEVPLVLMQDLAFLSGFPPTFKETNQLKTKLPENLSSKVLLQLYSEASVALLKLNPKDFQELNKQTKKNMTIDGKELTSPAYLLWDLAISQSKQDEDISASRFEDNEELRYSLSIERHAPWVRNIFIVTNGQIPSWLNLNPRVITIVTHQDVFRNLSHLPTFSSPAIESHIHRIEGLSQKFIYLNDDVMFGKDVVDDFYSHSKGQKQVLTWVPVNC AEGCPGSWIKDGYCDKACNNSACDWDGGDCSGNSGGSRVYAGGGGTGSIGVGPWQFGGINSVSYCNQGCANSWLADKFCQACNVLSCGFDAGDCGQDHFHELYKVILLPNQTHYIIPKGECLPYFSAEVAKRVEGAYSDNPIRHASIANKWKTIHLIMHSGMNATTIHFNLTFQNTNDEEFKMQITVEVDTRREGPKLNSTAQKGYENLVSPITLLPEAEILFEDIPKEKRFKFRHDDVSTRRAQEEVKIPLVNISLLPKDAQLSLNTLDLQLEHGDITLKGYNLSKASLLRSFLMNSQHAKIKNQAIITDETNDSLVAPEKQVHKSILPNSLGVSERLQRLTFPAVSVKVNGHDQGGQNPPLDLETTARFRVETHQKTIGGNVTKEKPPSLIVPLESQMTKEKKTIGKEKENSREMEENAENHIGVTEVLLGRKLQHYTDSYLGFLPWEKKKYFQDLLDEEESLKTQLAYFTDSKNTGRQLKDTFADSLRYVNKILNSKFGFTSRKVPAMMPHMI DRIVMQELQDMFPEEFDKTSFHKVRHSEDMQFAFSYFYLLMSAVQPLNISQVFDEVDTDQSGVLSDREIRTLATR IHELPLSLQDLTGLEHMLINCSKMLPADITQLNIPPTQESYYPNLPVTKSLVTNCKPVTDKIKHAYKDKNKYRFEIMGEEIEAFKMIRTNVSHVVGQLDDIRKNPRKVFCLNDNIDHNHKAQTVAVLRDVFYESMFPSPQFELPREYRNRFLHMHELQEWRAIRDKLFWTHCVLATLIMFTIFSFFAEQLIALKRKIFPRRRIHKEASPNRIRV	False	True	1.338	1.736	1.443	1.495	5.0	3.106	1.8