

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membran
Q06210	GFPT1_HUMAN	Homo sapiens	Glutamine-fructose-6-phosphate aminotransferase [isomerizing] 1	14.762029	NaN	S103;S261	34229054;23301498;35138101;34019948;30397120;34725712;38665916	MCGIFAYLNYHVPRTTRREILETLIKG LQRLEYRGYDSAGVGFDDGGNDKDW EANACKIQLIKKKGKVKALDEEVHK QQDMDLDIEFDVHLGIAHTRWATH GEPSPVNSHPQRSKNNFEVIVHNG IITNYKDLKKFLESKGYDFESETDTE TIAKLVKMYDNRRESQDTSFTTLVER VIQOLEGAFALVFKSVHFPQAQVGT RRGSPLLIGVRSEHKLSTDHIPILYRT ARTQIGSKFTRWGSQGERGKDKKGS CNLSRVSTTCLFPVEEKAVEYYFAS DASAVIEHTNRVIFLEDDVAAVVD GRLSIHRKRTAGDHPGRAVOTLQM ELQIMKGNFSSFMQKEIFEQPEV VNTMRGRVNFDDYTVNLGGLKDH KEIQRCRRLILIACGTSYHAGVATRO VLEELTELPVMVELASDFLDRNTPV FRDDVCFFLSQSGETADTLMGLRYC KERGALTVGITNTVGSISRETDCGV HINAGPEIGVASTKAYTSQFVSLVMF ALMMCDDRISMQERRKEIMLGLKR LPDLIKEVLSMDDEIQKLATELYHQK SVLIMGRGYHYATCLEGALKIKEITY MHSEGILAGELKHGPLALVDKLMVP IMIIMRDHTYAKCQNALQQVVARQG RPVVICDKEDTETIKNTKRTIKVPHS VDCLQGILSVIPLQLLAFHLAVLRGY DVDFFRNLAKSVTVE	True	True	4.469	3.103	2.48	2.122	1.755	2.091