

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	extra regio
0 Q9NQ88	TIGAR_HUMAN	Homo sapiens	Fructose-2,6-bisphosphatase TIGAR	20.761126	NaN	NaN	34229054;34019948;23301498	MARFALTVVRHGETRFNKEKIIQGQ GVDEPLSETGFKQAAAAGIFLNNVK FTHAFSSDLMRKQTMHGILERSKF CKDMTVKYDSRLRERKYGVVEGKAL SELRAMAKAAREECPVFTPPGGETL DQVKMRGIDFFEFLLCQLILKEADQK EQFSQGPSNCLLETSLAEIFPLGKN HSSKVNDSGIPGLAASVIVVSHGA YMRSFLDYFLTDLKCSPATLSRSEL MSVTPNTGMSLFIINFEEGREVKPT VQCICMNLQDHLNGLTETR	True	False	4.712	5.0	5.0	2.254	0.793	1.968	1.929