

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
0 Q9QXD6	F16P1_MOUSE	Mus musculus	Fructose-1,6-bisphosphatase 1	23.627151	NaN	T216;T245;T265	37453647	MANHAPFETDISTLTRFVMEQGRKA QGTGELTQLLNSLCTAIKAISSAVRQ AGIAQLYGIAGSTNVTGDQVKKLDIL SNDLVINMLKSSYATCVLVSEENTN AIIIEPKRGKYVVCFDPLDGSSNIDC LVSIGTIFGIYRKKSTDEPSEKDALQP GRDLVAAGYALYGSATMLVLAMDCG VNCFMLDPSIGEFIMVDRDVKMKKK GNIYSLNEGYAKDFDPAINAYLQKKK FPPDGSAPYGARYVGSMAVADHRTL VYGGIFLYPANKKSPSGKLRLLYECN PIAYVMEKAGGLATTGDKDILDIVPT EIHQKAPVVMGSSSEDVQEFLEIYRK HKAK	True	False	4.696	3.814	2.233	1.537	0.788	2.296	4.37