	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	O18738	DAG1_BOVIN	Bos taurus	Dystroglycan 1	39.03173	NaN	T790;T892	41018975	MRMSVGSAVPLPLWGRTFLLLLSVA VTQSHWPSEPSEAVRDWENQLEAS MHSALSDLHETVPTVVGIPDGTAVV GRSFRVTIPTDLIASNGEVIKVSAAGK EALPSWLHWDPQSHTLEGLPLDTD KGVHYISVSAARLGANGSHVPQTSS VFSIEVYPEDHSEPQSLRAASPDPGE VVSLVCAADEPVTVLTVILDADLTKM TPKQRIDLLRRMRGFSEVEPHNMKL VPVVNNRLFDMSAFMAGPGNAKKV VENGALLSWKLGCCLNQNSVPDIRG VEVPAREGAMSAQLGYPVVGWHIAN KKPSLPKRIRRQIHATPTPVTAIGPPT TAIQEPPSRIVPTPTSPAIAPPTETMA PPVRDPVPGKPTVTIRTRGAIIQTPTL GPIQPTRVSEAGTTVPSQIRPTMTIPG YMEPSTVTTPPTTTTKKPRVSTPRPA TPSTDSSTTTTRRPTKKPRTSRPVPR VTTKAPITRLETASPATRMRTTTSGV PHGGEPNQRPELKNHIDRVDAWVG TYFEVKIPSDTFYDNEDTTTDKLKLT LKLREQQLVGEKSWVQFNSNSQLM YGLPDSSHVGKHEYFMHATDKGGL SAVDAFEIHVHRRPQGDKAPARFKA KLTGDPAAVTNDIHKKIALVKKLAFA FGDRNCSTITLQNITRGSIVVEWTN NTLPLEPCPKEQITALSRRIAEDDGK PRGAFVNALEPDFQAMSITVTGSGS CRHLQFVPVAPPMRVPSEAPATEVP DRDPEKSSEDDVYLHTVIPAVVVAAI LLIAGIIAMICYRKKRKGKLTLEDQAT FIKKGVPIIFADELDDSKPPPSSSMPL ILQEEKAPLPPPEYPNQSMPETTPLN QDTVGEYAPLRDEDPSAPPYQPPPPF TAPMEGKGSRPKNMTPYRSPPPYVP P	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None