

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
Q8VD04	GRAP1_MOUSE	Mus musculus	GRIP1-associated protein 1	36.83566	S700	S620;S631;S633;S634;S653;S655;S656;S657;S795	40885482	MAQALSEEEFQRMQTQLELRNNTNN YQLSDELRRKNGVELSSLRQKVAYLD KEFSKAQKALSCKKAQEVVLLSE NEMLQAKLHSQEEDFRLQNSTLMA EFSKLCQSLEQLELENRQLKEGVPG AAGAHVDGELLRLQAENTALQKNM AALQERYGKEAVRPSAVGEGQGDPP GDVLPPTLAPMPLAEVELKWEMERE EKLLWEQLQGLESSKQAEYSRLQE ELAKLSEKLSKQESFCRLQTEKETL FNDSRNKIEELQQRKEADLKAQLAR TQKLQEQLEAANQSLAELRDQRQGE RLEHAAALRALQDQIQTAKTQELNM LREQTSELASELQHRQAEYEELMGQ KDDLNSQLQESLRANSRLLEQLQEI GQEKEQLTQDLQEARSAEKRVKVM LDELAMETLQEKSQHKEELGAVRLR HEKELLGVRARYERELRELHEDKKR QEEELRGQIREEKARTRELENLQHT VEELQAQVHSMGAKGWFERRLKE AEESLQQQQEQEETLKLCREEHAA ELKGDDEELQNVREQLQQAQEERD GHVKTISNLKQEVKDTVDGQRILEK KGSAYLKDLKRQLHLERKRADKLQE RLQEILTNSKSRGTGLEELVLEMNSP SRTQTGDSSSVSFSYREILKEKES AIPARLSLSSPQAQPPRPAELSDEEV AELFQLAETQEKWMLLEKVKHL EVSSASMAEDLCRKSIIETVVMDSR IDVSVAAAGHTDRSGLGSLRDLVKP GDENLREMNNKLQNMLEEQLTKN MHLHKDMEVLSQEIVRLSKECVGSP DPDLEPGEAN	False	False	3.346	3.585	0.908	0.846	2.067	1.373	0.756