

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
E9PYG6	E9PYG6_MOUSE	Mus musculus	Ras GTPase-activating protein 1	29.125745	NaN	NaN	39927985	MMAAEAGSEEGGPATAGPGGGAAT GSSAYSAAACRVKLPAAALPVAAAPCPG LADADLVAALGGGAASGSGFLGTGP VAGVLGGAALTGSAAAGVAGAAAAG PAGDVAFKGTLSLPAETLPGGGGFP PLPPPPQLPPLGSGLGTVDEGDSLGD PEYEEEEVAIPLTAPPTNQWYHGKL DRTIAEERLRQAGKSGSYLRESDRR PGSFVLSFLSQTNVNVNHFRIAMCG DYYIGGRRFSSLSDLIGYYSHVSCLL KGEKLLYPVAPPEPVEDRRRVRAILP YTKVPDTDEISFLKGD MFIVHNELE DGWMWVTNLR TDEOGLIVEDLVEE VGREEDPHEGKIWFHGKISKQEA YLLMTVGVQVCSFLVRPSDNTPGDYS L YFR TNENIQRFKICPTPNNQFMMGG RYNSIGDIIDHYRKEQIVEGYLKEP VPMQDQGOVLNDTVDGKEIYNTIRR KTKDAFYKNIVKKG YLLKKGKGRW KNLYFILEGS DAQLIYFESEKRATK KGLIDL SVCSVYVHDSLFRPNCF QIVVQHFSEEHYFYFAGETPEQAED WMKGLQAFCSLRKSSPGTSSNKRLR QVSSLVLHIEAHKLPVKHFTNYPY NIYLSVQVAKTHAREGQNPVWSEE FVFDLPPDINRFEITLSNKT KSKD PDILFMRCQLSRLQKGHATDEWFLL SSHIPLKGIPEGSLRVRARYSMEKIM PEEYSEFKELILQKELHVYALSHV CGQDR TLLASILLKIFLHEKLESLLC TLNDREISMEDEATLFRATTLASTL MEQYMKATATQFVH HALKDSILKIM ESKQSCELSPSKLEKNEDVNTNLAH LLSILSELVEKIFMASEILPPTLRYYG CLQKSVQHKWPTNNTMRTRVVS GF VFLRLICPAILNPRMFNII SDSPS PIAARTLTLVAKSVQNLANLVEFGAKEP YMEGVNPFIKSNKHRMIMFLDELG NVPELPDTTEHSRTDLSRD LAALHEI CVAHSDELRTL SNERGVQQHVLKKL LAITELLQKQNYTKTNDIR	True	True	4.272	4.225	0.469	False	False	5.0	False