

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9EQZ6	RPGF4_MOUSE	Mus musculus	Rap guanine nucleotide exchange factor 4	24.474581	NaN	NaN	33300544	MVAAHAAHSQSSAEWIACLDKRPLE RSSEEDVDIIFTRLKGVKAFKFFHPNL LRQICLCGYENLEKGITLFRQGDIG TNWYAVLAGSLDVKVSETSSHQDAV TICTLGIGTAFGESILDNTPRHATIVT RESSELLRIEQEDFKALWEKYROYM AGLLAPPYGVMETGSNNDRIPDKEN TPLIEPHVPLRPAHTITKVPSEKILRA GKILRIAILSRAPHMIRDRKYHLKTYR QCCVGTSLVDWMIQOTSCVHSRTQ AVGMWQVLEDDGVLNHVDQERHF QDKYLFYRFLDDEREDAPLPTEEK KECDEELQDTMLLSQMGPDAMMR MILRKPPGQRTVDDLEIHYDELLHIKA LSHLSITVKRELAVLIFESHAKGGT VLFNQGEETSWYIILKGSVNVVIYG KGVVCTLHEGDDFGKLALVNDAPRA ASIVLREDNCHFRLVDKEDFNRLR DVEANTVRLKEHDQDVLVLEKVPAG NRAANQGN SQPQKYTVMSGTPEK ILEHFLETIRLEPSLNEATDVLNDF VMMHCVFMPNTQLCPALVAHYHA QPSOCTEQERMDYALNNKRRVIRLV LQWAAMYGDLLQEDDVAMAFLEEF YVSVSDDARMMAAFKEQLPELEKIV KQISEDAPKQKHKVLLQFNTGD ERAQKRQPIRGSEVLFKVICIDHTY TTIRVPVAASVKEVISAVADKLGSGE GLIIVKMNSGGKVVLSKNDVSVFTT LTINGRLFACPREQFDSLTPLEQEG PTTGTVGTFELMSSKDLAYQMTTYD WELFNCVHELELIYHTFGRHNFKKT TANLDFLRRFNEIQFWVTEVCLC SQLSKRVQLLKKFIKIAAHCKEYKNL NSFFAIVMGLSNVAVSRLALTWEKL PSKFKKFYAEFESLMDPSRNHRAYR LTAACKLEPPLIPFMPLLIKDMTFHE GNKTFIDNLVNF EKMRMIANTARTV RYYRSQPFNPDAQAANKNHQDVRS YVRQLNVIDNQR TLSQMSHRLPRR P	True	True	5.0	2.716	2.355	1.827	0.87	4.544	1.749