	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparat
0	Q80TZ9	RERE_MOUSE	Mus musculus	Arginine-glutamic acid dipeptide repeats protein	41.206431	S1223;T1230	S53;S56;T120;S142;S304;T593;S594;S600;S613;S642;S656;S675;S679;S1098;S1105;S1107;T1111;T1251;S1258	36852467;22517741;34678516;40021952	MTADKDKDKDKEKDRDRDRDRERDKRDKARESENARPRRSCTLEGGAKNYAESDHSEDEDNDNNSATTEESNKKSRKKPPKKKSRYERTDTGEITSYITEDDVVYRPGDCVYIESRRPNTPYFICSIQDFKLVHSSQACCRSPAPAFCDPPACSLPVAPQPPQHLSEAGRGPGGSKRDHLLMNVKWYYRQSEVPDSVYQHLVQDRHNENDSGRELVITDPVIKNRELFISDYVDTYHAAALRGKCNISHFSDIFAAREFKARVDSFFYILGYNPETRRLNSTQGEIRVGPSHQAKLPDLQPFPSPDGDTVTQHEELVWMPGVSDCDLLMYLRAARSMAAFAGMCDGGSTEDGCVAASRDDTTLNALNTLHESSYDAGKALQRLVKKPVPKLIEKCWTEDEVKRFVKGLRQYGKNFFRIRKELLPSKETGELITFYYYWKKTPEAASSRAHRRHRRRQAVFRRIKTRTASTPVNTPSRPPSSEFLDLSSASEDDFDSEDSEQELKGYACRHCFTTTSKDWHHGGRENILLCTDCRIHFKKYGELPPIEKPVDPPPFMFKPVKEEDDGLSGKHSMRTRRSRGSMSTLRSGRKKQPTSPDGRASPINEDIRSSGRNSPSAASTSSNDSKAETVKKSAKKVKEEAASPLKSTKRQREKVASDTEDTDRITSKKTKTQEISRPNSPSEGEGESSDSRSVNDEGSSDPKDIDQDNRSTSPSIPSPQDNESDSDSSAQQQMLQAQPPALQAPSGAASAPSTAPPGTPQLPTQGPTPSATAVPPQGSPATSQPPNQTQSTVAPAAHTHIQQAPTLHPPRLPSPHPPLQPMTAPPSQSSAQPHPQPSLHSQGPPGPHSLQTGPLLQHPGPPQPFGLPSQPSQGQGPLGPSPAAAHPHSTIQLPASQSALQPQQPPREQPLPPAPLAMPHIKPPPTTPIPQLPAPQAHKHPPHLSGPSPFSLNANLPPPPALKPLSSLSTHHPPSAHPPPLQLMPQSQPLPSSPAQPPGLTQSQSLPPPAASHPTTGLHQVPSQSPFPQHPFVPGGPPPITPPSCPPTSTPPAGPSSSSQPPCSAAVSSGGSVPGAPSCPLPAVQIKEEALDEAEEPESPPPPPRSPSPEPTVVDTPSHASQSARFYKHLDRGYNSCARTDLYFMPLAGSKLAKKREEAIEKAKREAEQKAREEREKEKEKERERERERERREAERAAKASSSAHEGRLSDPQLSGPGHMRPSFEPPPTTIAAVPPYIGPDTPALRTLSEYARPHVMSPTNRNHPFYMPLNPTDPLLAYHMPGLYNVDPTIRERELRREIREIREIRERELRERMKPGFEVKPPELDPLHPATNPMEHFARHSALTIPPAAGPHPFASFHPGLNPLERERLALAGPQLRPEMSYPDRLAAERIHAERMASLTSDPLARLQMFNVTPHHHQHSHIHSHLHLHQQDPLHQGSAGPVHPLVDPLTAGPHLARFPYPPGTLPNPLLGQPPHEHEMLRHPVFGTPYPRDLPGAIPPPMSAAHQLQAMHAQSAELQRLAMEQQWLHGHPHMHGGHLPSQEDYYSRLKKEGDKQL	True	False	1.5	5.0	False	False	False