

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							
Q9R0X5	RPGR_MOUSE	Mus musculus	X-linked retinitis pigmentosa GTPase regulator	34.74265	S822	S518	34418053;22645316	MAESES LVPDTGAVFTFGTKKFAEN IPSKFWFKNDIPICLSCGDEHTAIVT GNNKLYMFGSNNWGQLGLGSKAAI IKPTCIKALKPEKVKLAACGRNHTLV STDTGGVYAAGGNNEGOLGLGDTD DRDTFHQIVFFTPADTIKQLSAGANT SAALTEGKLFMWGDNSEGQIGLE DKSNVCIPHEVTVGKPISWISCGYYH SAFVTMDGELYTFGEPENKGLGLPN ELLMNHRSPQRVLGIPERVIQVACC GGHTVVLTKEVVYAFGLGQFGQLGL GTFLFETSEPKIIRIKDQKICHISCG ENHTALMTELGLLYTFGDGRHGKL GLGMENFTNQFFPTLCSNFLRFAVQ LIACGGCHMLVFATPRLGTIDEPKFE DVYEPYISTGFSINDLSPRSSLNRSL SARLRRRERERPPCSASVMGTLPLPL EGTSASTSAYFYPPSSPPFHLSVNNYP EKSPSESMEPLDSDFEDKMNKDT ETENSSAVDSENFGETNDILNMTH MMTTSSNEKLLDFSPIQKQONQDTF EKVMESTPCTENEDSYEYEEMSKIK EVTVYKQYLAKGIYMIRPAEILEAFSD EEVGNGLDQVEEPRVFTDGKGLQSK QVGKESDEEIVSEKTEVMEVADVK KIRESEENSKSDSLFDDLPDKTMNS ESEDNKDIAEERRSSEQNMTFDSET ELVEEPDSYMECERHSEQDSAELE QPKLVEYSSEKDEKDEKDDDEVET ENLWYDRNCTEQETENVFRATRFFP KFDLKHDLHLSGIPEEQEGPEDSEGN VVVEQVVQAQKENLEFEFGDRKEAKA EAPSDVITEKEAPQLSETVKPEEGEM DEEISILNVEDTVEERKEGEKEIVE EGSIPETEGSETIDITDEKLDVLEKEE DSASLLQRALREYNENPKGHMYDR VKSSSSEILGGNDPTSKDIKAKKISF FNRMSLTGQKLMQNTNDPLPEIKPI GDQIALQSDKGDANQNHMGQNLQ DSTTPNMEGKSKSCTIL	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None