

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
Q92834	RPGR_HUMAN	Homo sapiens	X-linked retinitis pigmentosa GTPase regulator	18.379595	NaN	S418;S518	30379171	MREPEELMPDSGAVFTFGKSKFAE NNPGKFWFKNDVPVHLSGDEHSA VVTGNNKLYMFGSNNWGQLGLGSK SAISKPTCVKALKPEKVKLAACGRNH TLVSTEGGNVYATGGNNEGQLGLG DTEERNTFHVISFFTSEHKIKQLSAG SNTSAAALTEDGRLFMWGDNSEGQI GLKNVSNVCVPPQVVTIGKPVSWISC GYYHSAFVTTDGGELYVFGPEPENGKL GLPNQLLGNHRTPLVSEIPEKVIQV ACGGEHTVVLTEHAVYTFGLGQFGQ LGLGTFLEFETSEPKVIENIRDQTISIYI SCGENHTALITDIGLMYTFGDGRHG KLGLENFNTNHFIPITLCSNFLRFIV KLVACGGCHMVVFAAPHRGVAKEIE FDEINDTCLSVATFLPYSSLTSGNVL QRTLSARMRRRERERSPDSFSMRRT LPPIEGTLGLSACFLPNSVFPRCSE NLQESVLSQDLMOPEEPDYLLDE MTKEAEIDNSSTVESLGETTDILNM THIMSLNSNEKSLKLSPVQKQKKQQ TIGELTQDTALTENDDSDYEEMSE MKEGKACKQHVSQGIFMTQPATTIE AFSDEEVGNDTGQVGPQADTDGEG LQKEVYRHENNNNGVDQLDAKEIEKE SDGGHSQKESEAEIDSEKETKLAEI AGMKDLREREKSTKKMSPFFGNLP DRGMNTESEENKDFVKKRESCKQD VIFDSERESVEKPDYSMEGASESQQ GIADGFQQPEAIEFSSGEKEDDEVET DQNIRYGRKLIEQGNEKETKPIISKS MAKYDFKCDRLSEIPEEKEGAEDSK NGIEEQEVEANEENVKVHGGRKE KTEILSDDLTDKAEDHEFSKTEELKL EDVDEEINAENVESKKKTVDDES PTGYHKTGAERTNDSSAETIEK KEKANLEERAICEYENPKGYMLDD ADSSSLEILENSETTPSKDMKTKKI FLFKRVPSINQKIVKNNNEPLPEIKSI GDQIILKSDNKDADQNHMSQNHQN IPPTNTERRSKSCSTIL	None	None	None	None	None	None	None	None	

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