

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
Q812A2	SRGP3_MOUSE	Mus musculus	SLIT-ROBO Rho GTPase-activating protein 3	24.474581	NaN	S817;S820;S821;S837;S858;S954	33300544	MSSQTKFKKDKKEIIAEYEAQIKEIRT QLVEQFKCLEQQSESRLQLLQDLQE FFRRKAEIELEYSRSLKLAERFSSKI RSSREHQFKKQYLLSPVNCWYLVL HQTRRESRDHATLNDIFMNNVIVRL SQISEDVIRLFKKSKEIGLQMHEELL KVTNELYTMKTYHMYHAESISAES KLKEAEKQEEKQFNKSGELSMNLL RHEDRPQRRSSVKIEKMKEKRQAK YSENLKCTKARNDYLLNLAATNAA ISKYYIHDVSDLIDCCDLGFHASLAR TFRTYLSAEYNLETSRHEGLDVIENA VDNLDSSRDKHTVMDMCSQVFCPP LKFEFQPHMGDEVQVSAQQPVQT ELLMRYHQLQSRLATLKIENEVVRK TLDATMQLQDMLTVEDFDVSDAF QHSRSTESIKSASETYMSKINIAKR RANQQEEMFYFTKFKEYVNGSNLI TKLQAKHDLKQTLGGERAECGTT RPPCLPPKQKMRPRPLSVYSHKL FNGSMEAFIKDSGOAIPVAESCIRF INLYGLQQGIFRVPGSQVEVNDIKN SFERGEDPLVDDQNERDINSVAGVL KLYFRGLENPLFPKERFQDLISTIKL ENPADRVHPIQQILITLPRVIVVMR YLF AFLNLSQYSDENMMDPYNLAI CFGPTLMHIPDGDQPVSCQAHVNEV IKTIIHHHEAIFPSPRELEGPVYEKCM AGGEEYCDSPHSEPGTIDEVDHDNG TEPHTSDEEVEQIEAIAKFDYVGRSP RELSFKKGASLLLYHRASEDWWEG RHNGVDGLIPHQYIVQDMDDAFSD SLSQKADSEASSGPLLDDKASSKND LQSPTEHISDYGGVMGRVRLRSD GAAIPRRRSGDTHSPPRGLGPSIDT PPRAAACPSPHKIPLSRGRIESPEK RRMATFGSAGSINYPDKKALTEGLS MRSTCGSTRHSSLGDHKSLEAEALA EDIEKTMSTALHELRELERQNTVKQ APDVVLDTLEPLKNPPGPISSEPA LHTIVIRDPDAAMRRSSSSSTEMMT TFKPALSARLAGAQLRPPPMRPV VVQHRSSSSSSGSPAVTPTTEKM FPNSSSDKSGTM	None	None	None	None	None	None	None	None	

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