

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	extracellul region							
Q9I7D3	CAPR1_DROME	Drosophila melanogaster	Caprin homolog	25.496522	NaN	S356;S758;T764;S765;S777;T778;S791;S798;S806	33925313	MPSAANTATVTTAAIATTVAASANTN NNSVSKEKRSSIIISLGGSVIPGTAVAA GAGVATTNGQSNSSAAATADGSSQTD LASSNNNGNAAKSKANAVSAAAVVP EPYNPLKQLLVITIEHKIRNLEKRKTK LESYRAIQSSGKELSGDQASAVAKYD AVLANLEFARELAKHIQQQSKEAEK EOKKQARKDNLAKTIAETAKIREVLII QNVLNCFNDDQVRSDFLNGENGAK KLENTELELLEKFFIETQTRRPETAD DVSFIATAQKSALFYSTINARPKSF GEVSFEKLRSLFQQIQDSGYLDKYLL VPLAENAVANSTDSGTSGDGGESGD LLNDGSGELDAELPSEPSARNLEE GLDKLHLGVQAEILDQHOSYQQQQE RPSHLLPEQHQRAPSLHQQNSMV QQVSTQVYQAAPQAGGTTTPVHVLY APAPPAQQQPQQPPPHQLLSPPV NLQALQTGTAAPPAQQQPHPHTHFA PNVRAVEQNYFKOPPOHPQPGIQA PQTPQQQFMQHSRPLAEVLGTGR FHFLQDSELDNPEALAPPPPTNQGL VFEQQPQPNNHVEEPTQAILTLTFT NQSFPSPQPQPQQGPPQQQLFS PVLIIHQRRQPPQQQPMLIMPRL PTQQPQPAPQIGMQAGDVTSGFS YENAVVSYEQIQQLKQQQQQQ QQQNQLQLDEPSSVNSSSVGAG GDVENNEWHARNNDTNAATAALT NVLKGETTQATPPAPPTKWSSEMNA MSSAASNGSSNTSHKQEWATPRDH STGGGNSGYADNNSNHNHWNNSQ GDGRRNSGNYQRRGGDRDRDRD QGGRGDERRNGDRGNYSRQYGN SSNTNCRNSGNSGVYFRNNEESGN GGGNNYYQNGGGGGTYNKESRYE SSGGGGSYRGQRGGNQQRNVNGS YGGGRPMGDRGRGGAGNQQGGGG GYINRQNSQRMPLGLENK	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None

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