

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P39059	COFA1_HUMAN	Homo sapiens	Collagen alpha-1(XV) chain	22.903221	T210	NaN	38253038	MAPRRNNGQCWCLLMLLSVSTPLP AVTQTRGATETASQGHLDLTQLIGVP LPSSVSFVTGYGGFPAYSGPGANV GRPARTLIPSTFFRDFAISSVVKPSST RGGVLFITDAFQKVIYGLRLSGVE DGHQRILYYTEPGSHVSQEAFAFSV PVMTHRWNRFAMIVQGEVTLN CEEHSRIPFORSSQALAFESAGIFM GNAGATGLERFTGSLQQLTVHPDPR TPEELCDPESSASGETSGLQEADG VAEILEAVTYTQASPEAKVEPINTPP TPSSPFEDMELSGEPVPEGTLETTN MSIIQHSSPKQGSGEILNDTLEGVHS VDGDPITDSGSGAGAFDIAEEKNLA ATAAGLAEVPISTAGEAEASSVPTGG PTLSMSTENPEEGVTPGPDNEERLA ATAAGEAEALASMPGEVEASGVAPG ELDLSMSAQLGEEATVGPSSDSL TTAAAAEVSLSSTFEDEEASGVPTDG LAPLTATMAPERAVTSGPGDEEDLA AATTEEPLITAGGEESGSPPPDGPPPL PLPTVAPERWITPAQREHVGKMGQA GPKGEGDAGEELPGPPEPSGVPVGP TAGAEAEAGSLGWGSDVGSQSGDL VGSEQLLRGPPGPPGPPGLPGIPGKP GTDVFMGPPGSPGEDPAGEPGPPG PEGQPGVDGATGLPGMKGEKARG PNGSVGEKDPGNRGLPGPPGKKG QAGPPGVMGPPGPPGPPGPGCT MGLGFEDTEGSGSTQLLNPKLSRP TAAIGLKGEKDRGPKGERGMDGAS IVGPPGPRGPPGHKVLNSLNITHG FMNFSDIPELVGPPGPDGLPGLPGF PGPRGPKGDTGLPGFPLKGEQGEK GEPGAILTEDIPLERLMGKKGEPGM HGAPGPMGPKGPPGHKGEFGLPGR PGRPGLNGLKGTGDPGVIMQGP LPGPPGPPGPPGAVINIKGAIFPIVVR PHCKMPVDTAHPGSELPITFHGVKG EKGSWGLPGSKGEKDGGAQGP PPLDLAYLRHFLNLLKGENGDKGFK GEKGEKGDINGSFLMSGPPGLP GAGQKGETVVGPPGPPGAPGLPGP PGFGRPGDPGPPGPPGPPAILGA AVALP FSNMDDMLQKAHLVIEGTFIYLRDS TEFFIRVRDGWKKLQGLGELIPADS PPPPALSSNPHQLLPPNPISANYE KPALHLAALNMPFSGDIRADFQCFK QARAAGLLSTYRAFLSSHLQDLSTIV RKAERYSLPIVNLKQVLFNNWDSI FSGHGGQFNMHIPYISFDGRDIMTD PSWPQKVIWHGSSPHGVRLVDNYC EAWRTADTAVTGLASPLSTGKILDQK AYSCANRLIVLCIENSFMTDARK	False	True	2.01	1.798	2.751	4.285	0.735	3.333	4.467