

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
094898	LRIG2_HUMAN	Homo sapiens	Leucine-rich repeats and immunoglobulin-like domains protein 2	18.379595	NaN	T906	30379171	MAPAPLGVPEEQLLGCRSRVLSRLL FIAQTALLLLPAAGAGLCPAPCSCRIP LLDCSRRKLPAPSWRALSGLLPPDT AILDFSHNRLSNWNISLESQTLQEV KMNYNELTEIPYFGPEPTSNITLLSLV HNIIPEINAQALQFYPALESLLDSSNI ISEIKTSSFPRMQLKYLNLSNNRITT LEAGCFDNLSSLLVVKLNRRNRMS MIPPKIFKLPHLQFLELKRNRKIVEG LTFQGLDSLRLKMQRNGISKLDKG AFFGLNNMEELEHNNLTRVNKG WLYGLRMLQQLYVSQNAIERISPD WFCQRLSELDLSYNQLTRLDESFAF VGLSLLERLNLGDNRVTHIADGVFR FLSNLQTLDLRNNESWAIEDASEAF AGLTSCLKLILQGNQIKSITKKAFIGL ESLEHLDLNNNAIMSIQENAFSQTH LKEILINTSLLCDCHLKWLLQWL DNNFQHSVNVSCAHPWLAGQSIL NVDLKDVFCDDFLKPQIRTHPETIA LRGMNVTLTCTAVSSDSPMSTVW RKDSEILYDVTENFVRYWQAGEA LEYTSILHLFNVNFTDEGKYQCIVTN HFGSNYSQKAKLTVNEMPSFLKTP MDLTIRTGAMARLECAAEGHPAPQI SWQKDGDTDFPAARERRMHVMPE DDVFFIANVKIEDMGIYSCMAQNTA GGLSANASLTVLETPSFIRPLEDKTV TRGETAVLQCIAGGSPAPRLNWTKD DGPLLVTERRHFAAANQLLIIVDAGL EDAGKYTCIMSNLTLGTERGHIYLNVI SSPNCDSQSSIGHEDDGWTTVGVIV IVVCCVVGTSLIWVVIYHMRRKNE DYSITNTEELNLPADIPSYLSSQGTLS EPQEGYSNSEAGSHQQLMPPANGYI HKGTGGTGTRVICSDCYDNANIYS RTREYCPYTYIAEDVLDQTLSSLMV QMPKETYLVHPPQDTTALLESIPSAN REPSAFPTNHERISEKLPSTQMSG ETLQRPVWNINRELGLPHPPFSQPP VHESPQLHQNEGLAGREPDCSASS MSCHRLQDHAFDFSRTRNIQDGSE GT	False	False	1.113	2.429	0.684	0.673	1.441	4.348	3.516