

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
O94822	LTN1_HUMAN	Homo sapiens	E3 ubiquitin-protein ligase listerin	23.656127	T706;T1336;S1337	NaN	28510447;29351928	MGGKKNKQRTKGNLRPSNSGRAAEL LAKEQGTVPGFIFGFTSQSDLGYVPA IQGAEEIDSLVDSDFRMVLRKLSKDD VTTKLLKAMQEFMGTCERDTEITVKG VLPYWPFRIFCKISLDHDDRVRREATQQ AFEKLLIKVKKQLAPYLKSLMGYWL MAQCDTYTPAAFAAKDAFAEAFPPS KQPEAIAFCCKDEITSVLQDHLIKETP DTLSDPQTVPEEEREAKFYRVVTCSL LALKRLLCLLPDNELEKFKSLL SQNKFWYKHSVQIRSAFELVS ALCQRIPQLMKEEASKVSPSVLLSID DSDPIVCPALWEAVLYTLTTIEDCWL HVNAAKSVFPKLSTVIREGGGRGLATV IYPYLLPFISKLPQSITNPKLDFKKNF LTSLVAGLSTERTKTSSLESSAVISAF FECLRFIMQQNLGEEIEIQLMVDNQ LIPFIDAVLKDPGLQHGQLFNHIAET LSSWEAKADTEKDEKTAHNLENVLI HFWERLSEICVAKISEPEADVESVLG VSNLLQVLQPKPKSLLKSSKKNKGV RFADEILESNNKENEKCVSEGEKIEG WELTTEPSLTHNSSGLLSPLRKKPL EDLVCKLADISINYNERKSEQHLRF LSTLLDSFSSSRVFKMLLGDEKQSI QAKPLEIAKLVQKNPAVQFLYQKLLG WLNEDQRKDFGLVDILYSALRCCD NDMERKVVLDLTKVDLKWNSLLKI IEKACPSSDKHALVTPWLKGDILGEK LVNLADCLCNEDLESRVSESHFSE RWTLLSLVLSQHVKNLYLIGDVYVE RIIVRLHETLFTKTKLSEAESSDSSVS FICDVAYNYFSSAKGCLLMPSSDGLL LTLFQLCAQSKEKTHLPDFLICKLKN TWLSGVNLLVHQTDSYKSTFLHL SALWLKNQVQASSLDINSLQVLLSA VDDLNTLLESDSYLMGVYIGSVM PNDSEWEKMRQSLPMQWLHRPPL EGRLSLNYECFKTDFKEQDIKTLPSH LCTALLSKMVLIALRKETVLENNEL EKIAELLYSLQWCEELDNPPIFLIGF CEILOKMNITYDNLRLVGNLNTSGLLO LLFNRSREHGTLSLIIAKLILRSIS SDEVKPHYKRKESFFPLTEGNLHTI QSLCPFLSKEEKKEFSAQCIPALLGW TKKDLCTSTNGGFGHLAIFNSCLQTK SIDDGELLHGILKIIISWKEHEIDFL FSCNLSSEASPEVLGVNIEIRFSLFL KYCSSPLAESEWDFIMCSMLAWLET TSENQALYSIPLVQLFACVSCDLACD LSAFFDSTLDTIGNLPVNLISEWKE FFSQGIHSLLLPILVTVTGENKDVSE TSFQNALMKPMCETLTYISKEQLLS HKLPARLVADQKTNLPEYLQTLNLT LAPLLFRARPVQIAVYHMLYKLMPE LPQYDQDNLSYGDEEEEPALSPPA ALMSLLSIQEDLLENVLGCIPVQIV TIKPLSEDFCYVLYLLTWKLLITFFK AASSQLRALYSMYLRKTKSLNKLly HLFRLMPENPTYAETAVEVFNKDPK TFFTEELQLSIRETTMLPYHIPHLAC SVYHMTLKDLPAMVRLWVNSSEKR VFNIVDRFTSKYVSSVLSFQEISSVQ TSTQLFNGMTVKARATTREVMATYT IEDIVIELHQLPSNYPLGSIHVESGKR VGVAVQQWRNWMQLSTYLTHQN GSIMEGLALWKNVNDKRFEGVEDC MICFSVIHGFNYSPLPKKACRTCKKFF HSACLYKWFTSSNKSTCPLCRETFF	None	None	None	None	None	None	None		